vSMC - PARALLEL SEQUENTIAL MONTE CARLO IN C++

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VERSION DEVELOP

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ABSTRACT

Sequential Monte Carlo is a family of algorithms for sampling from a sequence of distributions. Some of these algorithms, such as particle filters, are widely used in the physics and signal processing researches. More recent developments have established their application in more general inference problems such as Bayesian modeling.

These algorithms have attracted considerable attentions in recent years as they admit natural and scalable parallelization. However, these algorithms are perceived to be difficult to implement. In addition, parallel programming is often unfamiliar to many researchers though conceptually appealing, especially for sequential Monte Carlo related fields. A C++ template library is presented for the purpose of implementing general sequential Monte Carlo algorithms on parallel hardware.

1 INTRODUCTION

Sequential Monte Carlo (SMC) methods are a class of sampling algorithms that combine importance sampling and resampling. They have been primarily used as "particle filters" to solve optimal filtering problems; see, for example, Cappé, Godsill, and Moulines (2007) and Doucet and Johansen (2011) for recent reviews. They are also used in a static setting where a target distribution is of interest, for example, for the purpose of Bayesian modeling. This was proposed by Del Moral, Doucet, and Jasra (2006b) and developed by Peters (2005) and Del Moral, Doucet, and Jasra (2006a). This framework involves the construction of a sequence of artificial distributions on spaces of increasing dimensions which admit the distributions of interest as particular marginals.

smc algorithms are perceived as being difficult to implement while general tools were not available until the development of Johansen (2009), which provided a general framework for implementing smc algorithms. smc algorithms admit natural and scalable parallelization. However, there are only parallel implementations of smc algorithms for many problem specific applications, usually associated with specific smc related researches. Lee et al. (2010) studied the parallelization of smc algorithms on gpus with some generality. There are few general tools to implement smc algorithms on parallel hardware though multicore CPUs are very common today and computing on specialized hardware such as GPUs are more and more popular.

The purpose of the current work is to provide a general framework for implementing SMC algorithms on both sequential and parallel hardware. There are two main goals of the presented framework. The first is reusability. It will be demonstrated that the same implementation source code can be used to build a serialized sampler, or using different programming models (for example, OpenMP and Intel Threading Building Blocks) to build parallelized samplers for multicore CPUs. The second is extensibility. It is possible to write a backend for vSMC to use new parallel programming models while reusing existing implementations. It is also possible to enhance the library to improve performance for specific applications. Almost all components of the library can be reimplemented by users and thus if the default implementation is not suitable for a specific application, they can be replaced while being integrated with other components seamlessly.

Section 2 introduces the SMC algorithm and notations used throughout this guide. Section 3 introduces the basic features of the library. It is followed by section 4, which details a few more advanced usage of the library. Section 5 to 9 introduces some useful additional features. It is not practical and of little interest to the users to include all the details of the library in this guide. A Doxygen¹ generated reference manual can be access online². Last, appendix A shows the complete source code of programs discussed in the following guide.

http://www.stack.nl/~dimitri/doxygen/

²http://zhouyan.github.io/vSMCDoc/develop/

2 SEQUENTIAL MONTE CARLO

2.1 SEQUENTIAL IMPORTANCE SAMPLING AND RESAMPLING

Importance sampling is a technique which allows the calculation of the expectation of a function φ with respect to a distribution π using samples from some other distribution η with respect to which π is absolutely continuous, based on the identity,

$$\mathbb{E}_{\pi}[\varphi(X)] = \int \varphi(x)\pi(x) \, \mathrm{d} x = \int \frac{\varphi(x)\pi(x)}{\eta(x)} \eta(x) \, \mathrm{d} x = \mathbb{E}_{\eta}\left[\frac{\varphi(X)\pi(X)}{\eta(X)}\right] \tag{1}$$

And thus, let $\{X^{(i)}\}_{i=1}^N$ be samples from η , then $\mathbb{E}_{\pi}[\varphi(X)]$ can be approximated by

$$\hat{\varphi}_1 = \frac{1}{N} \sum_{i=1}^{N} \frac{\varphi(X^{(i)})\pi(X^{(i)})}{\eta(X^{(i)})}$$
 (2)

In practice π and η are often only known up to some normalizing constants, which can be estimated using the same samples. Let $w^{(i)} = \pi(X^{(i)})/\eta(X^{(i)})$, then we have

$$\hat{\varphi}_2 = \frac{\sum_{i=1}^N w^{(i)} \varphi(X^{(i)})}{\sum_{i=1}^N w^{(i)}}$$
(3)

or

$$\hat{\varphi}_3 = \sum_{i=1}^N W^{(i)} \varphi(X^{(i)}) \tag{4}$$

where $W^{(i)} \propto w^{(i)}$ and are normalized such that $\sum_{i=1}^{N} W^{(i)} = 1$.

Sequential importance sampling (sis) generalizes the importance sampling technique for a sequence of distributions $\{\pi_t\}_{t\geq 0}$ defined on spaces $\{\prod_{k=0}^t E_k\}_{t\geq 0}$. At time t=0, sample $\{X_0^{(i)}\}_{i=1}^N$ from η_0 and compute the weights $W_0^{(i)} \propto \pi_0(X_0^{(i)})/\eta_0(X_0^{(i)})$. At time $t\geq 1$, each sample $X_{0:t-1}^{(i)}$, usually termed particles in the literature, is extended to $X_{0:t}^{(i)}$ by a proposal distribution $q_t(\cdot|X_{0:t-1}^{(i)})$. And the weights are recalculated by $W_t^{(i)} \propto \pi_t(X_{0:t}^{(i)})/\eta_t(X_{0:t}^{(i)})$ where

$$\eta_t(X_{0:t}^{(i)}) = \eta_{t-1}(X_{0:t-1}^{(i)})q_t(X_{0:t}^{(i)}|X_{0:t-1}^{(i)})$$
(5)

and thus

$$W_{t}^{(i)} \propto \frac{\pi_{t}(X_{0:t}^{(i)})}{\eta_{t}(X_{0:t}^{(i)})} = \frac{\pi_{t}(X_{0:t}^{(i)})\pi_{t-1}(X_{0:t-1}^{(i)})}{\eta_{t-1}(X_{0:t-1}^{(i)})q_{t}(X_{0:t}^{(i)}|X_{0:t-1}^{(i)})\pi_{t-1}(X_{0:t-1}^{(i)})}$$

$$= \frac{\pi_{t}(X_{0:t}^{(i)})}{q_{t}(X_{0:t}^{(i)}|X_{0:t-1}^{(i)})\pi_{t-1}(X_{0:t-1}^{(i)})}W_{t-1}^{(i)}$$
(6)

and importance sampling estimate of $\mathbb{E}_{\pi_t}[\varphi_t(X_{0:t})]$ can be obtained using $\{W_t^{(i)}, X_{0:t}^{(i)}\}_{i=1}^N$.

However this approach fails as t becomes large. The weights tend to become concentrated on a few particles as the discrepancy between η_t and π_t becomes larger. Resampling techniques are applied such that, a new particle system $\{\bar{W}_t^{(i)}, \bar{X}_{0:t}^{(i)}\}_{i=1}^M$ is obtained with the property,

$$\mathbb{E}\Big[\sum_{i=1}^{M} \bar{W}_{t}^{(i)} \varphi_{t}(\bar{X}_{0:t}^{(i)})\Big] = \mathbb{E}\Big[\sum_{i=1}^{N} W_{t}^{(i)} \varphi_{t}(X_{0:t}^{(i)})\Big]$$
(7)

In practice, the resampling algorithm is usually chosen such that M = N and $\bar{W}^{(i)} = 1/N$ for i = 1, ..., N. Resampling can be performed at each time t or adaptively based on some criteria of the discrepancy. One popular quantity used to monitor the discrepancy is *effective sample size* (ESS), introduced by Liu and Chen (1998), defined as

$$ESS_t = \frac{1}{\sum_{i=1}^{N} (W_t^{(i)})^2}$$
 (8)

where $\{W_t^{(i)}\}_{i=1}^N$ are the normalized weights. And resampling can be performed when ESS $\leq \alpha N$ where $\alpha \in [0,1]$.

The common practice of resampling is to replicate particles with large weights and discard those with small weights. In other words, instead of generating a random sample $\{\bar{X}_{0:t}^{(i)}\}_{i=1}^{N}$ directly, a random sample of integers $\{R^{(i)}\}_{i=1}^{N}$ is generated, such that $R^{(i)} \geq 0$ for $i=1,\ldots,N$ and $\sum_{i=1}^{N}R^{(i)}=N$. And each particle value $X_{0:t}^{(i)}$ is replicated for $R^{(i)}$ times in the new particle system. The distribution of $\{R^{(i)}\}_{i=1}^{N}$ shall fulfill the requirement of Equation (7). One such distribution is a multinomial distribution of size N and weights $(W_t^{(i)},\ldots,W_t^{(N)})$. See Douc, Cappé, and Moulines (2005) for some commonly used resampling algorithms.

2.2 SMC SAMPLERS

smc samplers allow us to obtain, iteratively, collections of weighted samples from a sequence of distributions $\{\pi_t\}_{t\geq 0}$ over essentially any random variables on some spaces $\{E_t\}_{t\geq 0}$, by constructing a sequence of auxiliary distributions $\{\tilde{\pi}_t\}_{t\geq 0}$ on spaces of increasing dimensions, $\tilde{\pi}_t(x_{0:t}) = \pi_t(x_t) \prod_{s=0}^{t-1} L_s(x_{s+1}, x_s)$, where the sequence of Markov kernels $\{L_s\}_{s=0}^{t-1}$, termed backward kernels, is formally arbitrary but critically influences the estimator variance. See Del Moral, Doucet, and Jasra (2006b) for further details and guidance on the selection of these kernels.

Standard sequential importance sampling and resampling algorithms can then be applied to the sequence of synthetic distributions, $\{\tilde{\pi}_t\}_{t\geq 0}$. At time t-1, assume that a set of weighted particles $\{W_{t-1}^{(i)}, X_{0:t-1}^{(i)}\}_{i=1}^N$ approximating $\tilde{\pi}_{t-1}$ is available, then at time t, the path of each particle is extended with a Markov kernel say, $K_t(x_{t-1}, x_t)$ and the set of particles $\{X_{0:t}^{(i)}\}_{i=1}^N$ reach the distribution $\eta_t(X_{0:t}^{(i)}) = \eta_0(X_0^{(i)}) \prod_{k=1}^t K_t(X_{t-1}^{(i)}, X_t^{(i)})$, where η_0 is the initial distribution of the particles. To correct the discrepancy between η_t and $\tilde{\pi}_t$, Equation (6)

is applied and in this case,

$$W_t^{(i)} \propto \frac{\tilde{\pi}_t(X_{0:t}^{(i)})}{\eta_t(X_{0:t}^{(i)})} = \frac{\pi_t(X_t^{(i)}) \prod_{s=0}^{t-1} L_s(X_{s+1}^{(i)}, X_s^{(i)})}{\eta_0(X_0^{(i)}) \prod_{k=1}^t K_t(X_{t-1}^{(i)}, X_t^{(i)})} \propto \tilde{w}_t(X_{t-1}^{(i)}, X_t^{(i)}) W_{t-1}^{(i)}$$

$$(9)$$

where \tilde{w}_t , termed the *incremental weights*, are calculated as,

$$\tilde{w}_{t}(X_{t-1}^{(i)}, X_{t}^{(i)}) = \frac{\pi_{t}(X_{t}^{(i)}) L_{t-1}(X_{t}^{(i)}, X_{t-1}^{(i)})}{\pi_{t-1}(X_{t-1}^{(i)}) K_{t}(X_{t-1}^{(i)}, X_{t}^{(i)})}$$
(10)

If π_t is only known up to a normalizing constant, say $\pi_t(x_t) = \gamma_t(x_t)/Z_t$, then we can use the *unnormalized* incremental weights

$$w_t(X_{t-1}^{(i)}, X_t^{(i)}) = \frac{\gamma_t(X_t^{(i)}) L_{t-1}(X_t^{(i)}, X_{t-1}^{(i)})}{\gamma_{t-1}(X_{t-1}^{(i)}) K_t(X_{t-1}^{(i)}, X_t^{(i)})}$$
(11)

for importance sampling. Further, with the previously *normalized* weights $\{W_{t-1}^{(i)}\}_{i=1}^{N}$, we can estimate the ratio of normalizing constant Z_t/Z_{t-1} by

$$\frac{\hat{Z}_t}{Z_{t-1}} = \sum_{i=1}^{N} W_{t-1}^{(i)} w_t(X_{t-1}^{(i)}, X_t^{(i)})$$
(12)

Sequentially, the normalizing constant between initial distribution π_0 and some target π_T , $T \ge 1$ can be estimated. See Del Moral, Doucet, and Jasra (2006b) for details on calculating the incremental weights. In practice, when K_t is invariant to π_t , and an approximated suboptimal backward kernel

$$L_{t-1}(x_t, x_{t-1}) = \frac{\pi(x_{t-1})K_t(x_{t-1}, x_t)}{\pi_t(x_t)}$$
(13)

is used, the unnormalized incremental weights will be

$$w_t(X_{t-1}^{(i)}, X_t^{(i)}) = \frac{\gamma_t(X_{t-1}^{(i)})}{\gamma_{t-1}(X_{t-1}^{(i)})}.$$
(14)

2.3 OTHER SEQUENTIAL MONTE CARLO ALGORITHMS

Some other commonly used sequential Monte Carlo algorithms can be viewed as special cases of algorithms introduced above. The annealed importance sampling (AIS; Neal (2001)) can be viewed as SMC samplers without resampling. Particle filters as seen in the physics and signal processing literature, can also be interpreted as the sequential importance sampling and resampling algorithms. See Doucet and Johansen (2011) for a review of this topic.

3 BASIC USAGE

3.1 CONVENTIONS

All classes that are accessible to users are within the name space vsmc. Class names are in CamelCase and function names, free or class methods, are in small_cases. In the remaining of this guide, we will omit the vsmc:: name space qualifiers. We will use "function" for referring to name space scope functions and "method" for class member functions.

3.2 GETTING AND INSTALLING THE LIBRARY

The library is hosted at GitHub³. This is a header only C++ template library. To install the library just move the contents of the include directory into a proper place, e.g., /usr/local/include on Unix-alike systems. This library requires working C++11, BLAS and LAPACK implementations. Standard C interface headers for the later two (cblas.h and lapacke.h) are required. Intel Threading Building Blocks⁴ (TBB), Intel Math Kernel Library⁵ (MKL) and HDF5⁶ are optional third-party libraries. One need to define the configuration macros VSMC_HAS_TBB, VSMC_HAS_MKL and VSMC_HAS_HDF5 to nonzero values before including any vSMC headers to make their existence known to the library, respectively.

3.3 CONCEPTS

The library is structured around a few core concepts. A sampler is responsible for running an algorithm. It is composed by a particle system and operations on it. A particle system is formed by the states $\{X^{(i)}\}_{i=1}^N$ and weights $\{W^{(i)}\}_{i=1}^N$. This system will also be responsible for resampling. All user defined operations are to be applied to the whole system. These are "initialization" and "moves" which are applied before resampling, and "mcmc" moves which are applied after resampling. These operations do not have to be mcmc kernels. They can be used for any purpose that suits the particular algorithm. Most statistical inferences requires calculation of $\sum_{i=1}^N W^{(i)} \varphi(X^{(i)})$ for some function φ . This can be carried out along each sampler iteration by a monitor. Table 1 lists these concepts and the corresponding classes in the library. Each of them are introduced in detail in the following sections.

³https://github.com/zhouyan/vSMC

⁴https://www.threadingbuildingblocks.org

⁵https://software.intel.com/en-us/intel-mkl

⁶http://www.hdfgroup.org

Concept	Class		
State, $\{X^{(i)}\}_{i=1}^{N}$	T, user defined		
Weight, $\{W^{(i)}\}_{i=1}^{N}$	Weight		
Particle, $\{W^{(i)}, X^{(i)}\}_{i=1}^{N}$	Particle <t></t>		
Single particle, $\{W^{(i)}, X^{(i)}\}$	SingleParticle <t></t>		
Sampler	Sampler <t></t>		
Initialization	Sampler <t>::init_type, user defined</t>		
Move	Sampler <t>::move_type, user defined</t>		
MCMC	Sampler <t>::mcmc_type, user defined</t>		
Monitor	Monitor <t></t>		

Table 1 Core concepts of the library

3.3.1 *State*

The library gives users the maximum flexibility of how the states $\{X^{(i)}\}_{i=1}^N$ shall be stored and structured. Any class type with a constructor that takes a single integer value, the number of particles, as its argument, and a method named copy is acceptable. For example,

```
1 class T
2 {
       public:
3
       T(std::size_t N);
4
       template <typename IntType>
5
       void copy(std::size_t N, IntType *index)
6
7
           for (std::size_t i = 0; i != N; ++i) {
8
                // Let a_i = index[i], set X^{(i)} = X^{(a_i)}
9
           }
10
       }
11
12 };
```

How the state values are actually stored and accessed are entirely up to the user. The method copy is necessary since the library assumes no knowledge of the internal structure of the state. And thus it cannot perform the last step of a resampling algorithm, which makes copies of particles with larger weights and eliminate those with smaller weights.

For most applications, the values can be stored within an N by d matrix, where d is the dimension of the state. The library provides a convenient class template for this situation,

```
1 template <MatrixLayout Layout, std::size_t Dim, typename T>
2 class StateMatrix;
```

where Layout is either RowMajor or ColMajor, which specifies the matrix storage layout; Dim is a non-negative integer value. If Dim is zero, then the dimension may be changed at runtime. If it is positive, then the dimension is fixed and cannot be changed at runtime. The last template parameter T is the C++ type of state space. The following constructs an object of this class,

```
1 StateMatrix<ColMajor, Dynamic, double> sm(N);
```

where Dynamic is just an enumerator with value zero. We can specify the dimension at runtime through the method sm.resize_dim(d). Note that, if the template parameter Dim is positive, then this call results in a compile-time error.

To access X_{ij} , the value of the state of the i^{th} particle at coordinate j, one can use the method sm.state(i, j). The method sm.data() returns a pointer to the beginning of the matrix. If Layout is RowMajor, then the method sm.row_data(i) returns a pointer to the beginning of the i^{th} row. If Layout is ColMajor, then the method sm.col_data(j) returns a pointer to the beginning of the j^{th} column. These methods help interfacing with numerical libraries, such as BLAS.

The StateMatrix class deliberately does not provide a resize method. There are algorithms that change the sample size between iterations. However, such algorithms often change size through resampling or other methods, either deterministically or stochastically. An example of changing size of a sampler is provided in section 7.1.

```
3.3.2 Weight
```

The vector of weights $\{W^{(i)}\}_{i=1}^N$ is abstracted in the library by the Weight class. The following constructs an object of this class,

```
1 Weight w(N);
```

There are a few methods for accessing the weights,

The weights can be manipulated, given a vector of length N, say v,

```
1 w.set(v);  // Set W^{(i)} \propto v^{(i)}

2 w.mul(v);  // Set W^{(i)} \propto W^{(i)} v^{(i)}

3 w.set_log(v);  // Set \log W^{(i)} = v^{(i)} + const.

4 w.add log(v);  // Set \log W^{(i)} = \log W^{(i)} + v^{(i)} + const.
```

The method w.data() returns a pointer to the normalized weights. It is important to note that the weights are always normalized and all mutable methods only allow access to $\{W^{(i)}\}_{i=1}^N$ as a whole.

3.3.3 Particle

A particle system is composed of both the state values, which is of user defined type, say \top , and the weights. The following constructs an object of class Particle< \top >,

```
1 Particle<T> particle(N);
```

The method particle.value() returns the type T object, and particle.weight() returns the type Weight object⁷. They are constructed with the same integer value N when the above constructor is invoked.

As a Monte Carlo algorithm, random number generators (RNG) will be used frequently. The user is free to use whatever RNG mechanism as they see fit. However, one common issue encountered in practice is how to maintain independence of the RNG streams between function calls. For example, consider below a function that manipulates some state values,

```
1 void function(double &x)
2 {
3     std::mt19937 rng;
4     std::normal_distribution<double> rnorm(0, 1);
5     x = rnorm(rng);
6 }
```

Every call of this function will give x exactly the same value. This is hardly what the user intended. One might consider an global RNG or one as class member data. For example,

```
1 std::mt19937 rng;
2 void function(double &x)
3 {
4    std::normal_distribution<double> rnorm(0, 1);
5    x = rnorm(rng);
6 }
```

This will work fine as long as the function is never called by two threads at the same time. However, SMC algorithms are natural candidates to parallelization. Therefore, the user will need to either lock the RNG, which degenerates the performance, or construct different RNGs for different threads. The later, though ensures thread-safety, has other issues. For example, consider

⁷More precisely, it is a Particle<T>::weight_type object, whose exact type depends on the type T. See section 4.2 for more details. If the user does not do something special as shown in that section, then the default type is the class Weight

```
1 std::mt19937 rng1(s1); // For thread i_1 with seed s_1 2 std::mt19937 rng2(s2); // For thread i_2 with seed s_2
```

where the seeds $s_1 \neq s_2$. It is difficult to ensure that the two streams generated by the two RNGs are independent. Common practice for parallel RNG is to use sub-streams or leap-frog algorithms. Without going into any further details, it is sufficient to say that this is perhaps not a problem that most users bother to solve.

The library provides a simple solution to this issue. The method particle.rng(i) returns a reference to an RNG that conforms to the C++11 uniform RNG concept. It can be called from different threads at the same time, for example,

```
1 auto &rng1 = particle.rng(i1); // Called from thread i_1
2 auto &rng2 = particle.rng(i2); // Called from thread i_2
```

If $i_1 \neq i_2$, then the subsequent use of the two RNGs are guaranteed to be thread-safe. In addition, they will produce independent streams. If TBB is available to the library, then it is also thread-safe even if $i_1 = i_2$. One can write functions that process each particle, for example,

And if later this function is called from a parallelized environment, it is still thread-safe and produce desired statistical results. The details of the RNG system are documented later in section 8.

3.3.4 Single particle

It is often easier to define a function $f(X^{(i)})$ than $f(X^{(1)},\ldots,X^{(N)})$. However, Particle<T> only provides access to $\{X^{(i)}\}_{i=1}^N$ as a whole through particle.value(). To allow direct access to $X^{(i)}$, the library uses a class template SingeParticle<T>. An object of this class is constructed from the index i of the particle, and a pointer to the particle system it belongs to,

```
1 SingleParticle<T> sp(i, &particle);
  or more conveniently,
1 auto sp = particle.sp(i);
```

In its most basic form, it has the following methods,

```
1 sp.id();  // Get the value i that sp was constructed with
2 sp.particle(); // Get a reference to the Particle<T> object sp belongs to
3 sp.rng();  // => sp.particle().rng(sp.id());
```

If T is a subclass of StateMatrix, then it has two additional methods,

```
1 sp.dim();  // => sp.particle().value().dim();
2 sp.state(j); // => sp.particle().value().state(sp.id(), j);
```

It is clear now that the interface of SingleParticle<T> depends on the type T. Later in section ?? we will show how to insert additional methods into this class.

3.3.5 Sampler

A sampler can be constructed in a few ways,

```
1 Sampler<T> sampler(N);
```

constructs a sampler that is never resampled, while

```
1 Sampler<T> sampler(N, Multinomial);
```

constructs a sampler that is resampled every iteration, using the multinomial algorithm. Other resampling schemes are also implemented, see section 7. Last, one can also construct a sampler that is only resampled when ESS $< \alpha N$, where $\alpha \in [0, 1]$, by the following,

```
1 Sampler<T> sampler(N, Multinomial, alpha);
```

If $\alpha > 1$, then it has the same effect as the first constructor, since ESS $\leq N$. If $\alpha < 0$, then it has the same effect as the second constructor, since ESS > 0.

In summary, if one does not tell the constructor which resampling scheme to use, then it is assumed one does not want to do resampling. If one specify the resampling scheme without a threshold for Ess, then it is assumed it need to be done at every step.

The method sampler.particle() returns a reference to the particle system. A sampler can be initialized by user defined object that is convertible to the following type,

```
1 using init_type = std::function<std::size_t(Particle<T> &, void *)>;
  For example,
1 auto init = [](Particle<T> &particle, void *param) {
2     // Process initialization parameter
3     // Initialize the particle system
4 };
```

is a C++11 lambda expression that can be used for this purpose. One can add it to a sampler by calling sampler.init(init). Upon calling sampler.initialize(param), the user defined function init will be called and the argument param will be passed to it.

Similarly, after initialization, at each iteration, the particle system can be manipulated by user defined callable objects that is convertible to the following types,

```
1 using move_type = std::function<std::size_t(std::size_t, Particle<T> &)>;
2 using mcmc_type = std::function<std::size_t(std::size_t, Particle<T> &)>;
```

Multiple moves can be added to a sampler. The call sampler.move(move, append) adds a move_type object to the sampler, where append is a boolean value. If it is false, it will clear any moves that were added before. If it is true, then move is appended to the end of an existing sequence of moves. Each move will be called one by one upon calling sampler.iterate(). A similar sequence of MCMC moves can also be added to a sampler. The call sampler.iterate() will call user defined moves first, then perform the possible resampling, and then the sequence of MCMC moves.

Note that the possible resampling will also be performed after the user defined initialization function is called by sampler.initialize(param). And after that, the sequence of MCMC moves will be called. If it desired not to perform mutations during initialization, then following can be used,

```
1 sampler.init(init).initialize(param);
2 sampler.move(move, false).mcmc(mcmc, false).iterate(n);
```

The above code also demonstrates that most methods of Sampler<T> return a reference to the sampler itself and thus method calls can be chained. In addition, method sampler.iterate(n) accepts an optional argument that specifies the number of iterations. It is a shortcut for

```
1 for (std::size_t i = 0; i != n; ++i)
2     sampler.iterate();
3.3.6     Monitor
```

Inferences using a SMC algorithm usually require the calculation of the quantity $\sum_{i=1}^{N} W^{(i)} \varphi(X^{(i)})$ at each iteration for some function φ . One can define callable object that is convertible to the following type,

```
1 using eval_type =
2    std::function<void(std::size_t, std::size_t, Particle<T> &, double *);
    For example,
1 void eval(std::size_t iter, std::size_t d, Particle<T> &particle, double *r)
2 {
```

```
for (std::size_t i = 0; i != particle.size(); ++i, r += dim) {
    auto sp = particle.sp(i);
    r[0] = /* \varphi_1(X^{(i)}) */;
    // ...
    r[d - 1] = /* \varphi_d(X^{(i)}) */;
    }
}
```

The argument d is the dimension of the vector function φ . The output is an N by d matrix in row major layout, with each row corresponding to the value of $\varphi(X^{(i)})$. Then one can add this function to a sampler by calling,

```
1 sampler.monitor("name", d, eval);
```

where the first argument is the name for the monitor, the second its dimension, and the third the evaluation function. At each iteration, after all the initialization, possible resampling, moves and MCMC moves are done, the sampler will calculate $\sum_{i=1}^{N} W^{(i)} \varphi(X^{(i)})$. This method has two optional arguments. First is a boolean value record only. If it is true, it is assumed that no summation is needed. For example,

In this case, the monitor acts merely as a storage facility. The second optional argument is stage which specifies at which point the monitoring shall happen. It can be MonitorMove, which specifies that the monitoring happens right after the moves and before resampling. It can also be MonitorResample, which specifies that the monitoring happens right after the resampling and before the MCMC moves. Last, the default is MonitorMCMC, which specifies that the monitoring happens after everything.

The output of a sampler, together with the records of any monitors it has can be output in plain text forms through a C++ output stream. For example,

```
1 std::cout << sampler;</pre>
```

We will see how this works later with a concrete particle filter example. If the HDF5 library is available, it is also possible to write such output to HDF5 format, for example,

```
1 hdf5store(sampler, file_name, data_name);
```

Details can be found in section 9.3.

3.4.1 Model and algorithm

This is an example used in Johansen (2009). Through this example, we will show how to re-implement a simple particle filter in vSMC. It shall walk one through the basic features of the library introduced above.

The state space model, known as the almost constant velocity model in the tracking literature, provides a simple scenario. The state vector X_t contains the position and velocity of an object moving in a plane. That is, $X_t = (X_{pos}^t, Y_{pos}^t, X_{vel}^t, Y_{vel}^t)^T$. Imperfect observations $Y_t = (X_{obs}^t, Y_{obs}^t)^T$ of the positions are possible at each time instance. The state and observation equations are linear with additive noises,

$$X_t = AX_{t-1} + V_t$$
$$Y_t = BX_t + \alpha W_t$$

where

$$A = \begin{pmatrix} 1 & \Delta & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{pmatrix} \qquad B = \begin{pmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \end{pmatrix} \qquad \alpha = 0.1$$

and we assume that the elements of the noise vector V_t are independent Gaussian with variance 0.02 and 0.001 for position and velocity, respectively. The observation noise, W_t comprises independent, identically distributed t-distributed random variables with degree of freedom v=10. The prior at time 0 corresponds to an axis-aligned Gaussian with variance 4 for the position coordinates and 1 for the velocity coordinates. The particle filter algorithm is shown in algorithm 1.

3.4.2 *Implementations*

The complete program is shown in appendix a.1. In this section we show the outline of the implementation.

The main program

```
1 vsmc::Sampler<PFState> sampler(N, vsmc::Multinomial, 0.5);
2 sampler.init(PFInit()).move(PFMove(), false).monitor("pos", 2, PFMEval());
3 sampler.initialize(const_cast<char *>("pf.data")).iterate(n - 1);
4 std::ofstream output("pf.out");
5 output << sampler;
6 output.close();</pre>
```

```
Initialization  \text{Set } t \leftarrow 0. \\ \text{Sample } X_{\text{pos}}^{(0,i)}, Y_{\text{pos}}^{(0,i)} \sim \mathcal{N}(0,4) \text{ and } X_{\text{vel}}^{(0,i)}, Y_{\text{vel}}^{(0,i)} \sim \mathcal{N}(0,1). \\ \text{Weight } W_0^{(i)} \propto \exp \ell(X_0^{(i)}|Y_0) \text{ where } \ell \text{ is the likelihood function.} \\ \text{Iteration} \\ \text{Set } t \leftarrow t+1. \\ \text{Sample} \\ X_{\text{pos}}^{(t,i)} \sim \mathcal{N}(X_{\text{pos}}^{(t-1,i)} + \Delta X_{\text{vel}}^{(t-1,i)}, 0.02) \qquad X_{\text{vel}}^{(t,i)} \sim \mathcal{N}(X_{\text{vel}}^{(t-1,i)}, 0.001) \\ Y_{\text{pos}}^{(t,i)} \sim \mathcal{N}(Y_{\text{pos}}^{(t-1,i)} + \Delta Y_{\text{vel}}^{(t-1,i)}, 0.02) \qquad Y_{\text{vel}}^{(t,i)} \sim \mathcal{N}(Y_{\text{vel}}^{(t-1,i)}, 0.001) \\ \text{Weight } W_t^{(i)} \propto W_{t-1}^{(i)} \exp \ell(X_t^{(i)}|Y_t). \\ \text{Repeat the Iteration } \text{step until all data are processed.} \\ \end{cases}
```

Algorithm 1 Particle filter algorithm for the almost constant velocity model.

Sampler<PFState> object is constructed first. Then the initialization PFInit, move PFMove and a monitor PFMEval that records X_{pos}^t and Y_{pos}^t are added to the sampler. The monitor is named "pos". Then it is initialized with the name of the data file "pf.data", and iterated n-1 times, where n is the number of data points. At last, the output is written into a text file "pf.out". Below is a short R^8 script that can be used to process the output

```
1 library(ggplot2)
2 pf <- read.table("pf.out", header = TRUE)
3 sink("pf.rout")
4 print(pf[1:5,])
5 sink()
6 obs <- read.table("pf.data", header = FALSE)
7 dat <- data.frame(
8 X = c(pf[["pos.0"]], obs[,1]),
9 Y = c(pf[["pos.1"]], obs[,2]))
10 dat[["Source"]] <- rep(c("Estimate", "Observation"), each = dim(obs)[1])</pre>
```

8http://r-project.org

```
11 plt <- qplot(x = X, y = Y, data = dat, geom = "path")
12 plt <- plt + aes(group = Source, color = Source, linetype = Source)
13 plt <- plt + theme_bw() + theme(legend.position = "top")
14 pdf("pf.pdf")
15 print(plt)
16 dev.off()</pre>
```

The print statement shows the first five lines of the output,

```
Size Resampled Accept.0
                                  ESS
                                         pos.0
                                                 pos.1
2 1 1000
                 1
                               2.9204 -1.21951 3.16397
3 2 1000
                          0 313.6830 -1.15602 3.22770
                          0 33.0421 -1.26451 3.04031
4 3 1000
                 1
5 4 1000
                 1
                          0 80.1088 -1.45922 3.37625
6 5 1000
                          0 382.8820 -1.47299 3.49230
```

The column Size shows the sample size at each iteration. The library does not provide direct support of changing the sample size. However, it is possible and an example is shown in section 7.1. The column Resampled shows nonzero values if resampling were performed and zero otherwise. For each moves and MCMC steps, an acceptance count will be recorded. In this particular example, it is irrelevant. Next the column ESS shows the value of ESS. The last two columns show the importance sampling estimates of the positions recorded by the monitor named "pos". The graphical representation of the output is shown in figure 1.

Before diving into the details of the implementation of PFState, etc., we will first define a few constant and types. The state space is of dimension 4. And it is natural to use a StateMatrix as the base class of PFState,

```
using PFStateBase = vsmc::StateMatrix<vsmc::RowMajor, 4, double>;
```

The numbers of particles and data points are also defined as constants in this simple example,

```
1 static constexpr std::size_t N = 1000; // Number of particles
2 static constexpr std::size_t n = 100; // Number of data points
```

Last, we define the following constants as the indices of each state component.

```
1 static constexpr std::size_t PosX = 0;
2 static constexpr std::size_t PosY = 1;
3 static constexpr std::size_t VelX = 2;
4 static constexpr std::size t VelY = 3;
```

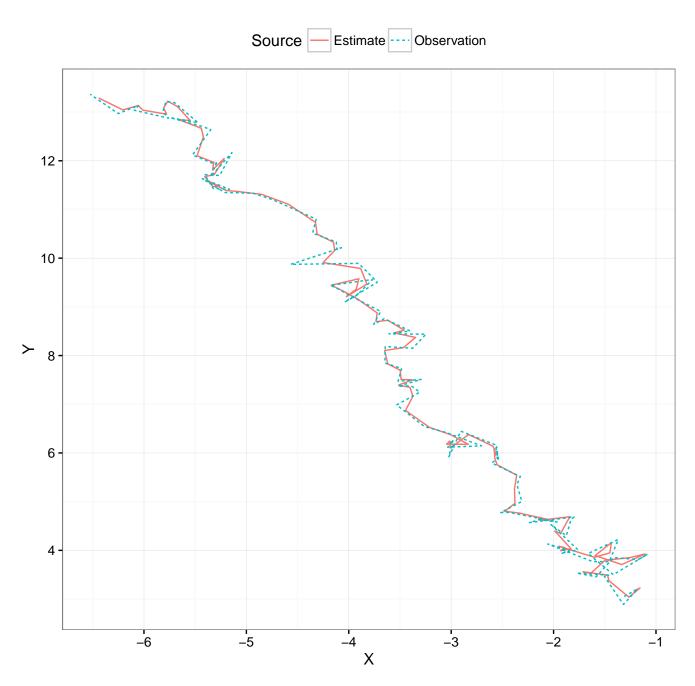


Figure 1 A simple particle filter

State: PFState As noted earlier, StateMatrix will be used as the base class of PFState. Since the data will be shared by all particles, we also store the data within this class. And methods will be provided to read the data from an external file, and compute the log-likelihood $\ell(X^{(i)})$, which accesses the data. Below the declaration of the class PFState is shown,

1 class PFState : public PFStateBase

```
2 {
       public:
3
       using PFStateBase::PFStateBase;
4
      // Return \ell(X_t^{(i)}|Y_t)
5
       double log_likelihood(std::size_t t, size_type i) const;
6
       // Read data from an external file
7
       void read data(const char *param);
8
       private:
9
       vsmc::Vector<double> obs_x_;
10
       vsmc::Vector<double> obs_y_;
11
12 };
   Initialization: PFInit The initialization step is implemented as below,
1 class PFInit
2 {
       public:
3
       std::size_t operator()(vsmc::Particle<PFState> &particle, void *param)
4
5
           eval_param(particle, param);
6
           eval_pre(particle);
7
           std::size t acc = 0;
8
           for (std::size t i = 0; i != particle.size(); ++i)
9
                acc += eval_sp(particle.sp(i));
10
           eval_post(particle);
11
           return acc;
12
       }
13
       void eval_param(vsmc::Particle<PFState> &particle, void *param)
14
```

```
{
15
           particle.value().read_data(static_cast<const char *>(param));
16
       }
17
      void eval pre(vsmc::Particle<PFState> &particle)
18
19
           w_.resize(particle.size());
20
       }
21
       std::size t eval sp(vsmc::SingleParticle<PFState> sp)
22
       {
23
           vsmc::NormalDistribution<double> norm pos(0, 2);
24
           vsmc::NormalDistribution<double> norm vel(0, 1);
25
           sp.state(PosX) = norm_pos(sp.rng());
26
           sp.state(PosY) = norm pos(sp.rng());
27
           sp.state(VelX) = norm_vel(sp.rng());
28
           sp.state(VelY) = norm_vel(sp.rng());
29
           w_[sp.id()] = sp.particle().value().log_likelihood(0, sp.id());
30
           return 0;
31
       }
32
      void eval post(vsmc::Particle<PFState> &particle)
33
       {
34
           particle.weight().set_log(w_.data());
35
36
       }
       private:
37
38
       vsmc::Vector<double> w ;
39 };
```

An object of this class is convertible to Sampler<PFState>::init_type. In the main method, operator(), eval_param is called first to initialize the data. Then eval_pre is called to allocated any resource this class need before calling any eval_sp. In this case, it allocate the vector w_ for storing weights computed later. Next, the main loop initializes each state component with the respective Gaussian distribution, computes the log-likelihood and store them in the vector allocated in the last step. This is done by calling the eval_sp method. After all particles have been initialized, we set the weights of the system in eval_post. Later in section 3.5 it will become clear why we structured the implementation this way.

Move: PFMove The move step is similar to the initialization. We show the declaration here,

```
1 class PFMove
2 {
      public:
3
       std::size_t operator()(std::size_t t, vsmc::Particle<PFState> &particle);
4
      void eval pre(std::size t t, vsmc::Particle<PFState> &particle);
5
      std::size_t eval_sp(std::size_t t, vsmc::SingleParticle<PFState> sp);
6
      void eval_post(std::size_t t, vsmc::Particle<PFState> &particle);
7
      private:
8
9
       vsmc::Vector<double> w ;
10 };
  Monitor: PFMEval Last we define PFMEval, which simply copies the values of the positions.
1 class PFMEval
2 {
      public:
3
      void operator()(std::size_t t, std::size_t dim,
4
           vsmc::Particle<PFState> &particle, double *r)
5
6
      {
           eval_pre(t, particle);
7
           for (std::size_t i = 0; i != particle.size(); ++i, r += dim)
8
               eval_sp(t, dim, particle.sp(i), r);
9
10
           eval_post(t, particle);
      }
11
      void eval_pre(std::size_t t, vsmc::Particle<PFState> &particle) {}
12
      void eval_sp(std::size_t t, std::size_t dim,
13
           vsmc::SingleParticle<PFState> sp, double *r)
14
15
           r[0] = sp.state(PosX);
16
           r[1] = sp.state(PosY);
17
       }
18
       void eval post(std::size t t, vsmc::Particle<PFState> &particle) {}
19
20 };
```

3.5 SYMMETRIC MULTIPROCESSING

The above example is implemented in a sequential fashion. However, the loops inside PFInit, PFMove and PFMEval clearly can be parallelized. The library provides basic support of multicore parallelization through its SMP module. Two widely used backends, OpenMP and TBB are available. Here we demonstrate how to use the TBB backend. First we will declare the implementation classes as subclasses as below,

```
1 class PFInit : public InitializationTBB<PFState>;
2 class PFMove : public MoveTBB<PFState>;
3 class PFMEval : public MonitorEvalTBB<PFState>;
```

And remove operator() from their implementations. After these changes, the implementation will be parallelized using TBB. The complete program can be found in section The complete program is shown in appendix a.2.

It works as if InitializationTBB<PFState> has an implementation of operator() as we did before, except it is parallelized. Now it is clear that, method such as eval_pre and eval_post are called before and after the main loop. Method eval_sp is called within the loop and it need to be thread-safe if called with different arguments. This is the main reason we constructed the NormalDistribution objects within eval_sp instead of as member data, even though they are constructed in exactly the same way for each particle. This is because NormalDistribution::operator() is a mutable method and thus not thread-safe. If any of these member functions does not do anything, then it does not have to be defined in the derived class.

Apart from the three base classes we have shown here, there are also InitializationOMP, etc., for using the OpenMP backend. And InitializationSEQ, etc., for implementation without parallelization. The later works in exactly the same way as our implementation in the last section. It is often easier to debug a single-threaded program than a parallelized one. And thus one may develop the algorithm with the sequential backend and obtain optimal performance latter by only changing the name of a few base class names. This can usually be done automatically through a build system.

3.5.1 *Performance consideration*

The base classes dispatch calls to eval_pre, eval_sp, etc., through the virtual function mechanism. The performance impact is minimal for eval_pre and eval_post, since they are called only once in each iteration and we expect the computational cost will be dominated by eval_sp in most cases. However, the dynamic dispatch can cause considerable performance degenerating if the cost of a single call to eval_sp is small while the number of particles is large. Modern optimizing compilers can usually devirtualize the method calls in trivial situations. However, it is not always possible. In this situation, the library will need a little help from the user to make compile-time dispatch. For each implementation class, we will declare it in the following way,

```
1 class PFInit : public InitializationTBB<PFState, PFInit>;
2 class PFMove : public MoveTBB<PFState, PFMove>;
3 class PFMEval : public MonitorEvalTBB<PFState, PFMEval>;
```

The second template argument of the base class need to be exactly the same as the derived class. For interested users, this is called Curiously Recurring Template Pattern⁹ (CRTP). This usage of the library's base classes also provides other flexibility. The methods eval_pre etc., can be either const or mutable. They can also be static.

 $^{^9} https://en.wikipedia.org/wiki/Curiously_recurring_template_pattern$

4 ADVANCED USAGE

4.1 CLONING OBJECTS

The Sampler<T> and Particle<T> objects have copy constructors, assignment operators, move constructors, and move assignment operators that behaves exactly the way as C++ programmers would expect. However, these behaviors are not always desired. For example, in Beskos et al. (2014) a stable particle filter in high-dimensions was developed. Without going into the details, the algorithm consists of a particle system where each particle is itself a particle filter. And thus when resampling the global system, the Sampler<T> object will be copied, together with all of its sub-objects. This include the RNG system within the Particle<T> object. Even if the user does not use this RNG system for random number generating within user defined operations, one of these RNG will be used for resampling by the Particle<T> object. Direct copying the Sampler<T> object will lead to multiple local filters start to generating exactly the same random numbers in the next iteration. This is an undesired side effects. In this situation, one can clone the sampler with the following method,

```
1 auto new_sampler = sampler.clone(new_rng);
```

where new_rng is a boolean value. If it is true, then an exact copy of sampler will be returned, except it will have the RNG system re-seeded. If it is false, then the above assignemnt behaves exactly the same as

```
1 auto new_sampler = sampler;
```

Alternatively, the contents of an existing Sampler<T> object can be replaced from another one by the following method,

```
1 sampler.clone(other_sampler, retain_rng);
```

where retain_rng is a boolean value. If it is true, then the RNG system of other_sampler is not copied and its own RNG system is retained. If it is false, then the above call behaves exactly the same as

```
1 sampler = other_sampler;
```

The above method also supports move semantics. Similar clone methods exist for the Particle<T> class.

4.2 CUSTOMIZING MEMBER TYPES

The Particle<T> class has a few member types that can be replaced by the user. If the class T has the corresponding types, then the member type of Particle<T> will be replaced. For example, given the following declarations inside class T,

```
1 class T
2 {
3    public:
4    using size_type = int;
5    using weight_type = /* User defined type */;
6    using rng_set_type = RNGSetTBB<AES256_4x32>;
7 };
```

The corresponding Particle<T>::size_type, etc., will have their defaults replaced with the above types. A note on weight_type, it needs to provide the following method,

```
1 w.ess(); // Get ESS 
2 w.set_equal(); // Set W^{(i)}=1/N 
3 w.resample_size(); // Get the size N. 
4 w.resample_data(); // Get a pointer to normalized weights
```

For the library's default class Weight, the last two calls are the same as w.size() and w.data(). However, this does not need to be so. For example, below is the outline of an implementation of weight_type for distributed systems, assuming each computing node has been allocated N_r particles.

```
1 class WeightMPI
2 {
       public:
3
       double ess()
4
5
           double local = /* \sum_{i=1}^{N_r} (W^{(i)})^2 */;
6
           double global = /* Gather local from each all nodes */;
7
           // Broadcast the value of global
8
           return 1 / global;
9
10
       }
       std::size_t resample_size() { return /* \sum N_r */; }
11
       double *resample_data()
12
       {
13
           if (rank == 0) {
14
               // Gather all normalized weights into a member data on this node
15
                // Say resample weight
16
                return resample weight .data();
17
```

```
} else {
18
                      return nullptr;
19
20
          }
21
         void set equal()
22
          {
23
                // Set all weights to 1/\sum N_r
24
                // Synchronization
25
          }
26
         void set(const double *v)
27
28
               // Set W^{(i)}=v_i for i=1,\ldots,N_r // Compute S_r=\sum_{i=1}^{N_r}W^{(i)} // Gathering S_r, compute S=\sum S_r
29
30
31
               // Broadcast S
32
               // Set W^{(i)} = W^{(i)}/S for i = 1, ..., N_r
33
          }
34
35 };
```

When Particle<T> performs resampling, it checks if the pointer returned by w.resample_data() is a null pointer. It will only generate the vector $\{a_i\}_{i=1}^N$ (see section 3.3.1) when it is not a null pointer. And then a pointer to this vector is passed to T::copy. Of course, the class T also needs to provide a suitable method copy that can handle the distributed system. By defining suitable WeightMPI and T::copy, the library can be extended to handle distributed systems.

4.3 EXTENDING SingleParticle<T>

The SingleParticle<T> can also be extended by the user. We have already see in section 3.3.1 that if class T is a subclass of StateMatrix, SingleParticle<T> can have additional methods to access the state. This class can be extended by defining a member class template inside class T. For example, for the simple particle filter in section 3.4, we can redefine the PFState as the following,

```
1 using PFStateBase = vsmc::StateMatrix<vsmc::RowMajor, 4, double>;
2 template <typename T>
3 using PFStateSPBase = PFStateBase::single_particle_type<T>;
```

```
4 class PFState : public PFStateBase
5 {
      public:
6
       using PFStateBase::StateMatrix;
7
      template <typename S>
8
       class single_particle_type : public PFStateSPBase<S>
9
       {
10
           public:
11
           using PFStateSPBase<S>::single_particle_type;
12
           double &pos_x() { return this->state(0); }
13
           double &pos y() { return this->state(1); }
14
           double &vel_x() { return this->state(2); }
15
16
           double &vel_y() { return this->state(3); }
          // Return $\ell(X_t^{(i)}|Y_t)$
17
           double log_likelihood(std::size_t t);
18
19
      };
      void read_data(const char *param);
20
      private:
21
       vsmc::Vector<double> obs_x_;
22
      vsmc::Vector<double> obs_y_;
23
24 };
  And later, we can use these methods when implement PFInit etc.,
1 class PFInit : public vsmc::InitializeTBB<PFState, PFInit>
2 {
      public:
3
      void eval_param(vsmc::Particle<PFState> &particle, void *param);
      void eval_pre(vsmc::Particle<PFState> &particle);
5
       std::size_t eval_sp(vsmc::SingleParticle<PFState> sp)
6
7
           vsmc::NormalDistribution<double> norm_pos(0, 2);
8
```

```
vsmc::NormalDistribution<double> norm_vel(0, 1);
9
           sp.pos_x() = norm_pos(sp.rng());
10
           sp.pos_y() = norm_pos(sp.rng());
11
           sp.vel_x() = norm_vel(sp.rng());
12
           sp.vel_y() = norm_vel(sp.rng());
13
           w_{sp.id()} = sp.log_likelihood(0);
14
           return 0;
15
16
      }
      void eval_post(vsmc::Particle<PFState> &particle);
17
      private:
18
      vsmc::Vector<double> w_;
19
20 };
```

It shall be noted that, it is important to keep single_particle_type small and copying the object efficient. The library will frequently pass argument of SingleParticle<T> type by value.

5 CONFIGURATION MACROS

The library has a few configuration macros. All these macros can be overwritten by the user by defining them with proper values before including any of the library's headers. Most of the macros have the prefix VSMC_HAS_, which specify if a feature is available to the library. A few have the prefix VSMC_USE_, which specify if a feature shall be used in the case that it is available. A handful remaining ones define some constants or types used by the library. These macros are listed in table 2.

Macro	Default	Description
VSMC_INT64	Platform dependent	The 64-bits integer type used by x86 intrinsics
		functions
VSMC_HAS_INT128	Platform dependent	Support for 128-bits integers
VSMC_INT128	Platform dependent	The 128-bits integer type
VSMC_HAS_SSE2	Platform dependent	Support for sse2 intrinsic functions
VSMC_HAS_AVX2	Platform dependent	Support for AVX2 intrinsic functions
VSMC_HAS_AES_NI	Platform dependent	Support for AES-NI intrinsic functions
VSMC_HAS_RDRAND	Platform dependent	Support for RDRAND intrinsic functions
VSMC_HAS_X86	Platform dependent	If we are using x86
VSMC_HAS_X86_64	Platform dependent	If we are using x86-64
VSMC_HAS_POSIX	Platform dependent	If we are on a POSIX platform
VSMC_HAS_OMP	Platform dependent	Support for OpenMP 3.0 or higher
VSMC_HAS_TBB	0	Support for твв 4.0 or higher
VSMC_HAS_TBB_MALLOC	VSMC_HAS_TBB	Support for TBB scalable memory allocation
VSMC_HAS_HDF5	0	Support for HDF5 1.8.6 or higher
VSMC_HAS_MKL	0	Support for мкl 11 or higher
VSMC_USE_MKL_CBLAS	VSMC_HAS_MKL	Use мкl header mkl_cblas.h instead of the
		standard cblas.h
VSMC_USE_MKL_LAPACKE	VSMC_HAS_MKL	Use мкц header mkl_lapacke.h instead of the
		standard lapacke.h
VSMC_USE_MKL_VML	VSMC_HAS_MKL	Use MKL vector mathematical functions (VML)
VSMC_USE_MKL_VSL	VSMC_HAS_MKL	Use MKL statistical functions (VSL)
VSMC_CBLAS_INT_TYPE	int	The default integer type of BLAS routines
VSMC_ALIGNMENT	32	Default alignment of AlignedAllocator

Table 2 Configuration macros

6 MATHEMATICAL OPERATIONS

6.1 CONSTANTS

The library defines some mathematical constants in the form of constexpr functions. For example, to get the value of π with a desired precision, one can call the following.

```
1 auto pi_f = const_pi<float>();
2 auto pi_d = const_pi<double>();
3 auto pi_l = const_pi<long double>();
```

The compiler will evaluate these values at compile-time and thus there is no performance difference from hard-coding the constants in the program, while the readability is improved. All defined constants are listed in table 3. Note that all functions has a prefix const_, which is omitted in the table.

Function	Value	Function	Value	Function	Value
pi	π	pi_2	2π	pi_inv	$1/\pi$
pi_sqr	π^2	pi_by2	$\pi/2$	pi_by3	$\pi/3$
pi_by4	$\pi/4$	pi_by6	$\pi/6$	pi_2by3	$2\pi/3$
pi_3by4	$3\pi/4$	pi_4by3	$4\pi/3$	sqrt_pi	$\sqrt{\pi}$
sqrt_pi_2	$\sqrt{2\pi}$	sqrt_pi_inv	$\sqrt{1/\pi}$	sqrt_pi_by2	$\sqrt{\pi/2}$
sqrt_pi_by3	$\sqrt{\pi/3}$	sqrt_pi_by4	$\sqrt{\pi/4}$	sqrt_pi_by6	$\sqrt{\pi/6}$
sqrt_pi_2by3	$\sqrt{2\pi/3}$	sqrt_pi_3by4	$\sqrt{3\pi/4}$	sqrt_pi_4by3	$\sqrt{4\pi/3}$
ln_pi	$\ln \pi$	ln_pi_2	$\ln 2\pi$	ln_pi_inv	$\ln 1/\pi$
ln_pi_by2	$\ln \pi/2$	ln_pi_by3	$\ln \pi/3$	ln_pi_by4	$\ln \pi/4$
ln_pi_by6	$\ln \pi/6$	ln_pi_2by3	$\ln 2\pi/3$	ln_pi_3by4	$\ln 3\pi/4$
ln_pi_4by3	$\ln 4\pi/3$	e	e	e_inv	1/e
sqrt_e	\sqrt{e}	sqrt_e_inv	$\sqrt{1/e}$	sqrt_2	$\sqrt{2}$
sqrt_3	$\sqrt{3}$	sqrt_5	$\sqrt{5}$	sqrt_10	$\sqrt{10}$
sqrt_1by2	$\sqrt{1/2}$	sqrt_1by3	$\sqrt{1/3}$	sqrt_1by5	$\sqrt{1/5}$
sqrt_1by10	$\sqrt{1/10}$	ln_2	ln 2	ln_3	ln 3
ln_5	ln 5	ln_10	ln 10	ln_inv_2	1/ln 2
ln_inv_3	$1/\ln 3$	ln_inv_5	$1/\ln 5$	ln_inv_10	1/ln 10
ln_ln_2	ln ln 2				

Table 3 Mathematical constants. Note: All functions are prefixed by const_.

6.2 VECTORIZED OPERATIONS

The library provides a set of functions for vectorized mathematical operations. For example,

```
1 std::size_t n = 1000;
2 vsmc::Vector<double> a(n), b(n), y(n);
3 // Fill vectors a and b
4 add(n, a.data(), b.data(), y.data());
performs addition for vectors. It is equivalent to
1 for (std::size_t i = 0; i != n; ++i)
2  y[i] = a[i] + b[i];
```

The functions defined are listed in table 4.

Function	Operation	Function	Operation	Function	Operation
add	a+b	sub	a-b	sqr	a^2
mul	ab	abs	a	fma	ab + c
inv	1/a	div	a/b	sqrt	\sqrt{a}
invsqrt	$1/\sqrt{a}$	cbrt	$\sqrt[3]{a}$	invcbrt	$1/\sqrt[3]{a}$
pow2o3	$a^{2/3}$	pow3o2	$a^{3/2}$	ром	a^b
hypot	$\sqrt{a^2+b^2}$	exp	e^a	exp2	2^a
exp10	10^a	expm1	$e^a - 1$	log	ln(a)
log2	$\log_2(a)$	log10	$\log_{10}(a)$	log1p	ln(a+1)
cos	$\cos(a)$	sin	sin(a)	sincos	sin(a) and $cos(a)$
tan	tan(a)	acos	arccos(a)	asin	arcsin(a)
atan	arctan(a)	acos	arccos(a)	atan2	arctan(a/b)
cosh	$\cosh(a)$	sinh	sinh(a)	tanh	tanh(a)
acosh	$\operatorname{arc} \cosh(a)$	asinh	arc sinh(a)	atanh	arc tanh(a)
erf	erf(a)	erfc	erfc(a)	cdfnorm	$1 - \operatorname{erfc}(a/\sqrt{2})/2$
lgamma	$\ln \Gamma(a)$	tgamma	$\Gamma(a)$		

Table 4 Vectorized mathematical operations

For each function, the first parameter is always the length of the vector, and the last is a pointer to the output vector (except sincos which has two output parameters). For all functions, the output is always a

vector. If there are more than one input parameters, then some of them, but not all, can be scalars. The order of the input parameters are as they appear in the mathematical expressions. For example, fma perform the operation ab + c, then the function shall be called as,

```
1 fma(n, a, b, c, y);
```

where a, b, c are input parameters, and some of them, not all, can be scalars instead of pointers. And y is the output parameter, which has to be pointer to a length n vector.

7 RESAMPLING

The library supports resampling in a more general way than the algorithm described in section 2. Recall that, given a particle system $\{W^{(i)}, X^{(i)}\}_{i=1}^N$, a new system $\{\bar{W}^{(i)}, \bar{X}^{(i)}\}_{i=1}^M$ is generated. Regardless other statistical properties, in practice, such an algorithm can be decomposed into three steps. First, a vector $\{r_i\}_{i=1}^N$ is generated such that $\sum_{i=1}^N r_i = M$. Then a vector $\{a_i\}_{i=1}^M$ is generated such that, $\sum_{i=1}^M \mathbb{I}_{\{j\}}(a_i) = r_j$. And last, set $\bar{X}^{(i)} = X^{(a_i)}$.

The first step determines the statistical properties of the resampling algorithm. The library defines all algorithms discussed in Douc, Cappé, and Moulines (2005). Samplers can be constructed with builtin schemes as seen in section 3.4.2. In addition, samplers can also be constructed with user defined resampling operations. Below is the signature,

```
1 template <typename IntType, typename RNGType>
2 void resample(std::size_t M, std::size_t N, RNGType &rng,
3 const double *weight, IntType *replication);
```

The last parameter is the output vector $\{r_i\}_{i=1}^N$. The builtin schemes are implemented as classes with operator() conforms to the above signature. For example, ResampleMultinomial implements the multinomial resampling algorithm.

To transform $\{r_i\}_{i=1}^N$ into $\{a_i\}_{i=1}^M$, one can call the following function,

```
1 template <typename IntType1, typename IntType2>
2 void resample_trans_rep_index(std::size_t M, std::size_t N,
3 const IntType1 *replication, IntType2 *index);
```

where the last parameter is the output vector $\{a_i\}_{i=1}^M$. This function guarantees that $a_i = i$ if $r_i > 0$. However, its output may not be optimal for all applications. The last step of a resampling operation, the copying of particles can be the most time consuming one, especially on distributed systems. The topology of the system will need to be taking into consideration to achieve optimal performance. In those situations, it is best to use ResampleMultinomial etc., to generate the replication numbers, and manually perform the rest of the resampling algorithm.

7.1 RESIZING A SAMPLER

Now, we provide an example of changing sampler size,

```
1 // sampler is an existing Sampler<T> object
2 auto N = sampler.size();
3 auto &rng = sampler.particle().rng();
4 auto weight = sampler.particle().weight().data();
```

```
5 Vector<std::size_t> rep(N);
6 Vector<std::size_t> idx(M);
7 ResampleMultinomial resample;
8 resample(M, N, rng, weight, rep.data());
9 resample_trans_rep_index(M, N, rep.data(), idx.data());
10 Particle<T> particle(M);
11 for (std::size_t i = 0; i != M; ++i) {
    auto sp_dst = particle.sp(i);
12
    auto sp_src = sampler.partice().sp(idx[i]);
13
   // Assuming T is a subclass of StateMatrix
14
   for (std::size_t d = 0; d != sp_dst.dim(); ++d)
15
   sp_dst.state(d) = sp_src.state(d);
16
17 }
18 // Copy other data of class T if any
19 sampler.particle() = std::move(particle);
```

8 RANDOM NUMBER GENERATING

The library has a comprehensive RNG system to facilitate implementation of Monte Carlo algorithms.

8.1 SEEDING

The singleton class template SeedGenerator can be used to generate distinctive seed sequentially. For example,

```
1 auto &seed = SeedGenerator<void, unsigned>::instance();
2 RNG rng1(seed.get()); // Construct rng1
3 RNG rng2(seed.get()); // Construct rng2 with another seed
```

The first argument to the template can be any type. For different types, different instances of SeedGenerator will be created. Thus, the seeds generated by SeedGenerator<T1> and SeedGenerator<T2> will be independent. The second parameter is the type of the seed values. It can be an unsigned integer type. Classes such as Particle<T> will use the generator of the following type,

```
1 using Seed = SeedGenerator<NullType, VSMC SEED RESULT TYPE>;
```

where VSMC SEED RESULT TYPE is a configuration macro which is defined to unsigned by default.

One can save and set the seed generator using standard C++ streams. For example

```
1 std::ifstream seed_txt("seed.txt");
2 if (seed_txt.good())
3 seed_txt >> Seed::instance(); // Read seed from a file
4 else
5 Seed::instance().set(101); // The default seed
6 seed_txt.close();
7 // The program
8 std::ofstream seed_txt("seed.txt");
9 seed_txt << Seed::instance(); // Write the seed to a file
10 seed txt.close();</pre>
```

This way, if the simulation program need to be repeated multiple times, each time is will use a different set of seeds.

A single seed generator is enough for a single computer program. However, it is more difficult to ensure that each computing node has a distinctive set of seeds in a distributed system. A simple solution is to use the modulo method of SeedGenerator. For example,

```
1 Seed::instance().modulo(n, r);
```

where n is the number of processes and r is the rank of the current node. After this call, all seeds generated will belong to the equivalent class $s \equiv r \mod n$. Therefore, no two nodes will ever generate the same seeds.

8.2 COUNTER-BASED RNG

The standard library provides a set of RNG classes. Unfortunately, none of them are suitable for parallel computing without considerable efforts.

The development by Salmon et al. (2011) made high performance parallel RNG much more accessible than it was before. In the author's personal opinion, it is the most significant development for parallel Monte Carlo algorithms in recent memory. See the paper for more details. Here, it is sufficient to mention that, the RNG introduced in the paper use a deterministic function f_k , such that, for any sequence $\{c_i\}_{i>0}$, the sequence $\{y_i\}_{i>0}$, $y_i = f_k(c_i)$, appears as random. In addition, for $k_1 \neq k_2$, f_{k_1} and f_{k_2} will generate two sequences that appear statistically independent. Compared to more conventional RNGs which use recursions $y_i = f(y_{i-1})$, these counter-based RNGs are much easier to setup in a parallelized environment.

If c, the counter, is an unsigned integer with b bits, and k, the key, is an unsigned integer with d bits. Then for each k, the RNG has a period 2^b . And there can be at most 2^d independent streams. Table 5 lists all counter-based RNGs implemented in this library, along with the bits of the counter and the key. They all conform to the C++11 uniform RNG concept. All RNGs in Salmon et al. (2011) are implemented along with a few additions. Note that, the actual period of an RNG can be longer. For example, Philox4x64 has a 256-bits counter but output 64-bits integers. And thus it has a 2^{1024} period. Such period length may seems very small compared to many well known RNGs. For example, the famous Mersenne-Twister generator (std::mt19937) has a period $2^{19937} - 1$. However, combined with 2^{256} independent streams, only the most demanding programs will find these counter-base RNGs insufficient.

8.2.1 AES-NI intrinsics based RNG

The AES-NI intrinsics based RNGs in Salmon et al. (2011) are implemented in a more general form,

```
1 template <typename ResultType, typename KeySeqType, std::size_t Rounds,
2 std::size_t Blocks>
3 using AESNIEngine =
4 CounterEngine<AESNIGenerator<ResultType, KeySeqType, Rounds, Blocks>>;
```

where KeySeqType is the class used to generate the sequence of round keys; Rounds is the number of rounds of AES encryption to be performed. See the reference manual for details of how to define the key sequence class. The AES-NI intrinsics have a latency of seven or eight cycles, while they can be issued at every cycle. Therefore better performance can be achieved if multiple 128-bits random integers are generated at the same time. This is specified by the template parameter Blocks. Larger blocks, up to eight, can improve runtime performance but this is at the cost of larger state size.

		Bits	
Class	Result type	Counter	Key
AES128_Bx32	std::uint32_t	128	128
AES128_ B x64	std::uint64_t	128	128
AES192_ B x32	std::uint32_t	128	192
AES192_ B x64	std::uint64_t	128	192
AES256_ B x32	std::uint32_t	128	256
AES256_ B x64	std::uint64_t	128	256
ARS_Bx32	std::uint32_t	128	128
ARS_Bx64	std::uint64_t	128	128
Philox2x32V	std::uint32_t	64	64
Philox2x64V	std::uint64_t	128	128
Philox4x32V	std::uint32_t	128	128
Philox4x64V	std::uint64_t	256	256
Threefry2x32 <mark>V</mark>	std::uint32_t	64	64
Threefry2x64 <mark>V</mark>	std::uint64_t	128	128
Threefry4x32 <mark>V</mark>	std::uint32_t	128	128
Threefry4x64 <mark>V</mark>	std::uint64_t	256	256

Table 5 Counter-based RNG; B: either 1, 2, 4, or 8; V: either empty, SSE2, or AVX2.

Four types of key sequences are implemented by the library, corresponding to the ARS algorithm in Salmon et al. (2011) and the AES-128, AES-192, and AES-256 algorithms. The following RNG engines are defined.

```
template <typename ResultType, std::size_t Rounds = VSMC_RNG_ARS_ROUNDS,
std::size_t Blocks = VSMC_RNG_ARS_BLOCKS>
using ARSEngine =
AESNIEngine<ResultType, ARSKeySeq<ResultType>, Rounds, Blocks>;

template <typename ResultType, std::size_t Blocks = VSMC_RNG_AES_BLOCKS>
using AES128Engine =
AESNIEngine<ResultType, AES128KeySeq<ResultType, 10>, 10, Blocks>;

template <typename ResultType, std::size_t Blocks = VSMC_RNG_AES_BLOCKS>
using AES192Engine =
```

Macro	Default
VSMC_RNG_AES_BLOCKS	4
VSMC_RNG_ARS_ROUNDS	5
VSMC_RNG_ARS_BLOCKS	4
VSMC_RNG_PHILOX_ROUNDS	10
VSMC_RNG_PHILOX_VECTOR_LENGTH	4
VSMC_RNG_THREEFRY_ROUNDS	20
VSMC_RNG_THREEFRY_VECTOR_LENGTH	4

Table 6 Configuration macros for the counter-based RNG

```
10 AESNIEngine<ResultType, AES192KeySeq<ResultType, 12>, 12, Blocks>;
11 template <typename ResultType, std::size_t Blocks = VSMC_RNG_AES_BLOCKS>
12 using AES256Engine =
13 AESNIEngine<ResultType, AES256KeySeq<ResultType, 14>, 14, Blocks>;
```

The default template arguments can be changed by configuration macros listed in table 6. Shortcuts are also defined, as listed in table 5. For example, ARS_4x32 is ARSEngine with result type std::uint32_t, four blocks, and the default number of rounds.

8.2.2 Philox

The Philox algorithm in Salmon et al. (2011) is implemented in a more general form,

```
1 template <typename ResultType, std::size_t K = VSMC_RNG_PHILOX_VECTOR_LENGTH,
2 std::size_t Rounds = VSMC_RNG_PHILOX_ROUNDS>
3 using PhiloxEngine = CounterEngine<PhiloxGenerator<ResultType, K, Rounds>>;
```

The default vector length and the number of rounds can be changed by configuration macros listed in table ??. Shortcuts are also defined, as listed in table 5. For example, Philox4x32 is PhiloxEngine with result type std::uint32_t, vector length four, and the default number of rounds.

8.2.3 *Threefry*

The Threefry algorithm in Salmon et al. (2011) is implemented in a more general form,

```
1 template <typename ResultType, std::size_t K = VSMC_RNG_THREEFRY_VECTOR_LENGTH,
2 std::size_t Rounds = VSMC_RNG_THREEFRY_ROUNDS>
```

```
3 using ThreefryEngine = CounterEngine<ThreefryGenerator<ResultType, K, Rounds>>;
```

The default vector length and the number of rounds can be changed by configuration macros listed in table ??. Shortcuts are also defined, as listed in table 5. For example, Threefry4x32 is ThreefryEngine with result type std::uint32_t, vector length four, and the default number of rounds.

If ssE2 intrinsics are supported, then a more optimized version is also implemented. This implementation can have higher performance at the cost of larger state size. If AVX2 intrinsics are supported, then a even more optimized, with even larger state size version is also implemented.

```
template <typename ResultType, std::size_t K = VSMC_RNG_THREEFRY_VECTOR_LENGTH,
std::size_t Rounds = VSMC_RNG_THREEFRY_ROUNDS>
using ThreefryEngineSSE2 =
CounterEngine<ThreefryGeneratorSSE2<ResultType, K, Rounds>>;

template <typename ResultType, std::size_t K = VSMC_RNG_THREEFRY_VECTOR_LENGTH,
std::size_t Rounds = VSMC_RNG_THREEFRY_ROUNDS>
using ThreefryEngineAVX2 =
CounterEngine<ThreefryGeneratorAVX2<ResultType, K, Rounds>>;
```

8.2.4 Default RNG

Note that, not all RNGs defined by the library is available on all platforms. The library also defines a type alias RNG which is one of the RNGs listed in table 5. More specifically, if the AES-NI intrinsics are supported,

```
1 using RNG = ARS_4x32;
  otherwise if Avx2 intrinsics are supported,
1 using RNG = Threefry4x32AVX2;
  otherwise if SSE2 intrinsics are supported,
1 using RNG = Threefry4x32SSE2;
  and last, on all other platforms,
1 using RNG = Threefry4x32;
```

This can be changed by the configuration macro VSMC_RNG_TYPE.

```
8.3 NON-DETERMINISTIC RNG
```

If the RDRAND intrinsics are supported, the library also implement three RNGs, RDRAND16, RDRAND32 and RDRAND64. They output 16-, 32-, and 64-bits random unsigned integers, respectively.

Class	MKL BRNG
MKL_MCG59	VSL_BRNG_MCG59
MKL_MT19937	VSL_BRNG_MT19937
MKL_MT2203	VSL_BRNG_MT2203
MKL_SFMT19937	VSL_BRNG_SFMT19937
MKL_NONDETERM	VSL_BRNG_NONDETERM
MKL_ARS5	VSL_BRNG_ARS5
MKL_PHILOX4X32X10	VSL_BRNG_PHILOX4X32X10

Table 7 MKL RNG. Note: all classes can have a suffix _64.

8.4 MKL RNG

The MKL library provides some high performance RNGs. The library implement a wrapper class MKLEngine that makes them accessible as C++11 generators. They are listed in table 7. Note that, MKL RNGs performs best when they are used to generate vectors of random numbers. These wrappers use a buffer to store such vectors. And thus they have much larger state space than usual RNGs.

8.5 MULTIPLE RNG STREAMS

Earlier in section 3.3.3 we introduced that particle.rng(i) returns an independent RNG instance. This is actually done through a class template called RNGSet. Three of them are implemented in the library. They all have the same interface,

```
1 RNGSet<RNG> rng_set(N); // A set of N RNGs
2 rng_set.resize(n); // Change the size of the set
3 rng_set.seed(); // Seed each RNG in the set with Seed::instance()
4 rng set[i]; // Get a reference to the i-th RNG
```

The first implementation is RNGSetScalar. As its name suggests, it is only a wrapper of a single RNG. All calls to rng_set[i] returns a reference to the same RNG. It is only useful when an RNGSet interface is required while the thread-safety and other issues are not important.

The second implementation is RNGSetVector. It is an array of RNGs with length N. It has memory cost O(N). Many of the counter-based RNGs have small state size and thus for moderate N, this cost is not an issue. The method calls rng_set[i] and rng_set[j] return independent RNGs if $i \neq j$.

Last, if TBB is available, there is a third implementation RNGSetTBB, which uses thread-local storage (TLS). It has much smaller memory footprint than RNGSetVector while maintains better thread-safety. The performance impact of using TLS is minimal unless the computation at the calling site is trivial. For example,

```
1 std::size_t eval_pre(SingleParticle<T> sp)
2 {
3   auto &rng = sp.rng();
4   // using rng to initialize state
5   // do some computation, likely far more costly than TLS
6 }
```

The type alias RNGSet is defined to be RNGSetTBB if TBB is available. Otherwise it is defined to be RNGSetVector. It is used by the Particle class template. One can replace the type of RNG set used by Particle<T> with a member type of T. For example,

```
1 class T
2 {
3  using rng_set_type = /* User defined type */;
4 };
```

will make the RNG set used by Particle<T> replaced by the user defined type.

8.6 DISTRIBUTIONS

The library also provides implementations of some common distributions. They all conforms to the C++11 random number distribution concepts. Some of them are the same as those in the C++11 standard library, with CamelCase names. For example, NormalDistribuiton can be used as an drop-in replacement for std::normal_distribuiton. This includes all of the continuous distributions defined in the standard library. Their benefits compared to the standard library will be discussed later. Table 8 lists all the additional distributions implemented.

The last, the library also implement the multivariate Normal distribution. Its usable is summarized by the following.

```
1 double mean[2] = { /* the mean vector */ };
2 double cov[4] = { / * the covariance matrix */ };
3 double chol[3];
4 double r[2];
5 // Compute the Lower triangular of the Cholesky decomposition
6 cov_chol(2, cov, chol);
7 RNG rng;
8 NormalMVDistribution<double, 2> norm2(mean, chol); // Bivariate Normal
9 NormalMVDistribution<double, Dynamic> normd(2, mean, chol); // Same as above
10 norm2(rng, r); // Generate a bivariate Normal
11 normd(rng, r); // Same as above
```

Class	Notes	
UniformBits	No parameters, uniform on the set $\{0, \dots, 2^b - 1\}$, where <i>b</i> is the number of bits	
	of the result type, which has to be an unsigned integer type.	
U01	No parameters, uniform on [0, 1)	
U01CC	No parameters, uniform on [0, 1]	
U01CO	No parameters, uniform on [0, 1)	
U010C	No parameters, uniform on (0, 1]	
U0100	No parameters, uniform on (0, 1)	
Laplace	Parameters: location a; scale b	
Levy	Parameters: location a; scale b	
Pareto	Parameters: shape a; scale b	
Rayleigh	Parameters: scale sigma	

Table 8 Random number distributions. Note: all class names have a suffix Distribution which is omitted in the table

We shall mention here that the static form, where the dimension is specified as a template parameter is more efficient.

8.7 VECTORIZED RANDOM NUMBER GENERATING

The RNGs and distributions implemented by this library provides vectorized operations. For example,

```
1 std::size_t n = 1000;
2 RNG rng;
3 NormalDistribution<double> norm(0, 1);
4 Vector<RNG::result_type> u(n);
5 Vector<double> r(n);
6 rng(n, u.data());  // Generate n random unsigned integers
7 rng_rand(rng, n, u.data()); // Same as above
8 norm(rng, n, r.data()); // Generate n Normal random numbers
9 normal_distribution(rng, n, r.data(), 0.0, 1.0);  // Same as above
10 normal_distribution(rng, n, r.data(), norm.param()); // Same as above
11 rng_rand(rng, norm, n, r.data());  // Same as above
```

Note that these functions will be specialized to use MKL routines if rng is one of the engines listed in table 7.

8.8 RANDOM WALK

9 UTILITIES

The library provides some utilities for writing Monte Carlo simulation programs. For some of them, such as command line option processing, there are more advanced, dedicated libraries out there. The library only provides some basic functionality that is sufficient for most simple cases.

9.1 ALIGNED MEMORY ALLOCATION

The standard library class std::allocator is used by containers to allocate memory. It works fine in most cases. However, sometime it is desired to allocate memory aligned by a certain boundary. The library provides the class template,

where the configuration macro VSMC_ALIGNMENT is defined to be 32 by default. For the requirement of the parameter type Memory, see the reference manual. It is sufficient to mention here that the default implementation works best if TBB is available. This class can be used as a drop-in replacement of std::allocator<T>. In fact, this library defines a type alias Vector<T> which is std::vector<T, AlignedAllocator<T>> if T is a scalar type, and std::vector<T> otherwise.

9.2 SAMPLE COVARIANCE ESTIMATING

The library provides some basic functionality to estimate sample variance. For example,

```
1 constexpr std::size_t d = /* Dimension */;
2 using T = StateMatrix<RowMajor, d, double>;
3 Sampler<T> sampler(N);
4 // operations on the sampler
5 double mean[d];
6 double cov[d * d];
7 Covariance eval;
8 auto x = sampler.particle().value().data();
9 auto w = sampler.particle().weight().data();
10 eval(RowMajor, N, d, x, w, mean, cov);
```

The sample covariance matrix will be computed and stored in cov. The mean vector is stored in mean. Note that, if any of them is a null pointer, then the corresponding output is not computed. The sample \times is assumed to be stored in an N by d matrix. The first argument passed to eval is the storage layout of

this matrix. If x is a null pointer, then no computation will be done. If w is a null pointer, then the weight is assumed to be equal for all samples. The method has three additional optional parameters. The first is cov_layout, which specifies the covariance matrix storage layout. The second is cov_upper and the third cov_packed, both are false by default. If the later is cov_packed, a packed vector of length d(d + 1)/2 is written into cov. If cov_upper is false, then the upper triangular is packed, otherwise the lower triangular is packed.

The estimated covariance matrix is often used to construct multivariate Normal distribution for the purpose of generating random walk scales. The NormalMVDistribution in section 8.6 accepts the lower triangular of the Cholesky decomposition of the covariance instead of the covariance matrix itself. The following function will compute this decomposition,

```
1 double chol[d * (d + 1) / 2];
2 cov_chol(d, cov, chol);
```

The output chol is a packed vector in row major storage. This function also has three optional parameters, which are the same as those of Covariance::operator(), except that they are now used to specify the storage scheme of the input parameter cov.

```
9.3 STORING OBJECTS IN HDF5
```

If the HDF5 library is available, it is possible to store Sampler<T> objects, etc., in the HDF5 format. For example,

```
1 hdf5store(sampler, "pf.h5", "sampler");
```

create a HDF5 file with the sampler stored as a list. In R it can be processed as the following,

```
1 library(rhdf5)
2 pf <- as.data.frame(h5read("pf.h5", "sampler"))</pre>
```

This creates a data. frame similar to that shown in section 3.4.2. Other types of objects can also be store, see the reference manual for details.

```
9.4 RAII CLASSES FOR MKL POINTERS
```

The library provides a few classes to manage MKL pointers. It provides Resource Acquisition Is Initialization (RAII) idiom on top of the MKL C interface. For example,

```
1 // VSLSSTaskPtr ptr;
2 // vsldSSNewTask(&ptr, &p, &n, &xstorage, x, w, indices);
3 MKLSSTask<double> task(&p, &n, &xstorage, x, w, indices);
```

Class	MKL pointer type
MKLStream	VSLStreamStatePtr
MKLSSTask	VSLSSTaskPtr
MKLConvTask	VSLConvTask
MKLCorrTask	VSLCorrTask
MKLDFTask	DFTaskPtr

Table 9 RAII classes for MKL pointers

```
4 // vsldSSEditMoments(ptr, mean, r2m, r3m, r4m, c2m, c3m, c4m);
5 task.edit_moments(mean, r2m, r3m, r4m, c2m, c3m, c4m);
6 // vsldSSCompute(ptr, estimates, method);
7 task.compute(estimates, method)
8 // vslSSDeleteTask(&ptr);
```

In the above snippets, MKLSSTask manages a VSLSSTaskPtr task pointer. All C functions that operates on the pointer, is also defined as methods in the class. Table 9 lists the classes defined by the library and their corresponding MKL pointers.

9.5 PROGRAM OPTIONS

The library provides some basic support of processing command line program options. Here we show a minimal example. The complete program is shown in appendix a.3. First, one need to allocated variables to store the options to be processed.

```
1 int n;
2 std::string str;
3 std::vector<double> vec;
```

All types that support standard library IO operations are supported. In addition, std::vector<T>, where T is a type that supports standard library IO operations is also supported. Then one constructs a ProgramOptionMap object,

```
1 ProgramOptionMap option_map;
```

Options can be added to the map,

```
1 option_map
2 .add("str", "A string option with a default value", &str, "default")
```

```
3 .add("n", "An integer option", &n)
4 .add("vec", "A vector option", &vec);
```

The first argument is the name of the option, the second is a description, and the third is a pointer to where the option's value shall be stored. The last optional argument is a default value. The options on the command line can be processed as the following,

```
1 option_map.process(argc, argv);
```

where argc and argv are the arguments of the main function. When the program is invoked, each option can be passed like below,

```
1 ./program_option --vec 1 2 1e-1 --str "abc" --vec 8 9 --str "def hij" --n 2 4
```

The results of the option processing is displayed below,

```
1 n: 4
2 str: def hij
3 vec: 1 2 0.1 8 9
```

To summarize these output, the same option can be specified multiple times. If it is a scalar option, the last one is used (--str, --n). A string option's value can be grouped by quotes. For a vector option (--vec), all values are gather together and inserted into the vector.

9.6 PROGRAM PROGRESS

Sometime it is desirable to see how much progress of a program has been made. The library provide a Progress class for this purpose. Here we show a minimal example. The complete program is shown in appendix a.4.

```
1 vsmc::Progress progress;
2 progress.start(n * n);
3 for (std::size t i = 0; i != n; ++i) {
      std::stringstream ss;
      ss << "i = " << i;
5
      progress.message(ss.str());
6
      for (std::size_t j = 0; j != n; ++j) {
7
          // Do some computation
8
          progress.increment();
9
      }
10
11 }
12 progress.stop();
```

When invoked, the program output something similar the below

```
1 [ 4%][00:07][ 49019/1000000][i = 49]
```

The method progress.start(n * n) starts the printing of the progress. The argument specifies how many iterations there will be before it is stopped. The method progress.message(ss.str()) direct the program to print a message. This is optional. Each time after we finish n iterations, we increment the progress count by calling progress.increment(). And after everything is finished, the method progress.stop() is called.

```
9.7 TIMING
```

Performance can only be improved after it is first properly benchmarked. There are advanced profiling programs for this purpose. However, sometime simple timing facilities are enough. The library provides a simple class StopWatch for this purpose. As its name suggests, it works much like a physical stop watch. Here is a simple example

```
1 StopWatch watch;
2 for (std::size_t i = 0; i != n; ++i) {
3     // Some computation
4     watch.start();
5     // Computation to be benchmarked;
6     watch.stop();
7     // Some other computation
8 }
9 double t = watch.seconds(); // The time in seconds
```

The above example demonstrate that timing can be accumulated between loop iterations, function calls, etc. It shall be noted that, the time is only accurate if the computation between watch.start() and watch.stop() is non-trivial.

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A SOURCE CODE OF COMPLETE PROGRAMS

A.1 SEQUENTIAL IMPLEMENTATION OF A SIMPLE PARTICLE FILTER

```
1 #include <vsmc/vsmc.hpp>
2 static constexpr std::size_t N = 1000; // Number of particles
3 static constexpr std::size_t n = 100; // Number of data points
4 static constexpr std::size_t PosX = 0;
5 static constexpr std::size_t PosY = 1;
6 static constexpr std::size_t VelX = 2;
7 static constexpr std::size_t VelY = 3;
8 using PFStateBase = vsmc::StateMatrix<vsmc::RowMajor, 4, double>;
9 class PFState : public PFStateBase
10 {
11
      public:
12
      using PFStateBase::PFStateBase;
      double log_likelihood(std::size_t t, size_type i) const
13
14
          double llh_x = 10 * (this->state(i, PosX) - obs_x_[t]);
15
           double llh_y = 10 * (this->state(i, PosY) - obs_y_[t]);
16
          llh_x = std::log(1 + llh_x * llh_x / 10);
17
          llh_y = std::log(1 + llh_y * llh_y / 10);
18
          return -0.5 * (10 + 1) * (llh_x + llh_y);
19
20
      }
      void read data(const char *param)
21
22
          if (param == nullptr)
23
               return;
24
          obs_x_.resize(n);
25
          obs_y_.resize(n);
26
           std::ifstream data(param);
27
```

```
28
           for (std::size_t i = 0; i != n; ++i)
               data >> obs_x_[i] >> obs_y_[i];
29
           data.close();
30
       }
31
      private:
32
      vsmc::Vector<double> obs x ;
33
      vsmc::Vector<double> obs_y_;
34
35 };
36 class PFInit
37 {
      public:
38
       std::size_t operator()(vsmc::Particle<PFState> &particle, void *param)
39
40
           eval_param(particle, param);
41
           eval_pre(particle);
42
           std::size_t acc = 0;
43
           for (std::size_t i = 0; i != particle.size(); ++i)
44
               acc += eval_sp(particle.sp(i));
45
           eval_post(particle);
46
           return acc;
47
      }
48
      void eval_param(vsmc::Particle<PFState> &particle, void *param)
49
50
           particle.value().read_data(static_cast<const char *>(param));
51
52
      }
      void eval_pre(vsmc::Particle<PFState> &particle)
53
54
      {
           w_.resize(particle.size());
55
      }
56
      std::size_t eval_sp(vsmc::SingleParticle<PFState> sp)
57
58
           vsmc::NormalDistribution<double> norm_pos(0, 2);
59
```

```
60
           vsmc::NormalDistribution<double> norm_vel(0, 1);
           sp.state(PosX) = norm_pos(sp.rng());
61
           sp.state(PosY) = norm_pos(sp.rng());
62
           sp.state(VelX) = norm_vel(sp.rng());
63
           sp.state(VelY) = norm_vel(sp.rng());
64
           w_[sp.id()] = sp.particle().value().log_likelihood(0, sp.id());
65
           return 0;
66
       }
67
68
       void eval_post(vsmc::Particle<PFState> &particle)
69
       {
           particle.weight().set log(w .data());
70
       }
71
72
       private:
       vsmc::Vector<double> w_;
73
74 };
75 class PFMove
76 {
       public:
77
       std::size_t operator()(std::size_t t, vsmc::Particle<PFState> &particle)
78
79
           eval_pre(t, particle);
80
           std::size_t acc = 0;
81
           for (std::size_t i = 0; i != particle.size(); ++i)
82
               acc += eval_sp(t, particle.sp(i));
83
           eval_post(t, particle);
84
           return 0;
85
86
       }
       void eval_pre(std::size_t t, vsmc::Particle<PFState> &particle)
87
88
       {
           w_.resize(particle.size());
89
90
```

```
std::size_t eval_sp(std::size_t t, vsmc::SingleParticle<PFState> sp)
91
 92
           vsmc::NormalDistribution<double> norm_pos(0, std::sqrt(0.02));
 93
            vsmc::NormalDistribution<double> norm_vel(0, std::sqrt(0.001));
 94
            sp.state(PosX) += norm_pos(sp.rng()) + 0.1 * sp.state(VelX);
 95
           sp.state(PosY) += norm pos(sp.rng()) + 0.1 * sp.state(VelY);
96
           sp.state(VelX) += norm_vel(sp.rng());
97
           sp.state(VelY) += norm vel(sp.rng());
 98
           w_[sp.id()] = sp.particle().value().log_likelihood(t, sp.id());
99
           return 0;
100
       }
101
       void eval_post(std::size_t t, vsmc::Particle<PFState> &particle)
102
103
           particle.weight().add_log(w_.data());
104
       }
105
106
       private:
       vsmc::Vector<double> w ;
107
108 };
109 class PFMEval
110 {
       public:
111
       void operator()(std::size_t t, std::size_t dim,
112
            vsmc::Particle<PFState> &particle, double *r)
113
       {
114
115
           eval pre(t, particle);
           for (std::size t i = 0; i != particle.size(); ++i, r += dim)
116
                eval sp(t, dim, particle.sp(i), r);
117
           eval_post(t, particle);
118
       }
119
       void eval_pre(std::size_t t, vsmc::Particle<PFState> &particle) {}
120
       void eval_sp(std::size_t t, std::size_t dim,
121
            vsmc::SingleParticle<PFState> sp, double *r)
122
```

```
{
123
           r[0] = sp.state(PosX);
124
           r[1] = sp.state(PosY);
125
126
       }
       void eval_post(std::size_t t, vsmc::Particle<PFState> &particle) {}
127
128 };
129 int main()
130 {
131
       vsmc::Sampler<PFState> sampler(N, vsmc::Multinomial, 0.5);
       sampler.init(PFInit()).move(PFMove(), false).monitor("pos", 2, PFMEval());
132
       sampler.initialize(const cast<char *>("pf.data")).iterate(n - 1);
133
134
       std::ofstream output("pf.out");
       output << sampler;</pre>
135
       output.close();
136
137
       return 0;
138 }
   A.2 PARALLELIZED IMPLEMENTATION OF A SIMPLE PARTICLE FILTER
 1 #include <vsmc/vsmc.hpp>
 2 static constexpr std::size_t N = 1000; // Number of particles
 3 static constexpr std::size_t n = 100; // Number of data points
 4 using PFStateBase = vsmc::StateMatrix<vsmc::RowMajor, 4, double>;
 5 template <typename T>
 6 using PFStateSPBase = PFStateBase::single_particle_type<T>;
 7 class PFState : public PFStateBase
 8 {
       public:
 9
       using PFStateBase::StateMatrix;
 10
```

```
template <typename S>
11
       class single_particle_type : public PFStateSPBase<S>
12
       {
13
           public:
14
           using PFStateSPBase<S>::single_particle_type;
15
           double &pos_x() { return this->state(0); }
16
           double &pos_y() { return this->state(1); }
17
           double &vel_x() { return this->state(2); }
18
           double &vel_y() { return this->state(3); }
19
           double log likelihood(std::size t t)
20
           {
21
               double llh_x = 10 * (pos_x() - obs_x(t));
22
               double llh_y = 10 * (pos_y() - obs_y(t));
23
               llh_x = std::log(1 + llh_x * llh_x / 10);
24
               llh_y = std::log(1 + llh_y * llh_y / 10);
25
               return -0.5 * (10 + 1) * (llh_x + llh_y);
26
27
           }
           private:
28
           double obs_x(std::size_t t)
29
30
               return this->particle().value().obs_x_[t];
31
32
           double obs_y(std::size_t t)
33
34
               return this->particle().value().obs y [t];
35
           }
36
      };
37
      void read_data(const char *param)
38
       {
39
           if (param == nullptr)
40
               return;
41
```

```
obs_x_.resize(n);
42
           obs_y_.resize(n);
43
           std::ifstream data(param);
44
           for (std::size_t i = 0; i != n; ++i)
45
               data >> obs_x_[i] >> obs_y_[i];
46
           data.close();
47
       }
48
       private:
49
       vsmc::Vector<double> obs_x_;
50
       vsmc::Vector<double> obs_y_;
51
52 };
53 class PFInit : public vsmc::InitializeTBB<PFState, PFInit>
54 {
       public:
55
       void eval_param(vsmc::Particle<PFState> &particle, void *param)
56
57
           particle.value().read_data(static_cast<const char *>(param));
58
59
       void eval_pre(vsmc::Particle<PFState> &particle)
60
       {
61
           w_.resize(particle.size());
62
       }
63
       std::size_t eval_sp(vsmc::SingleParticle<PFState> sp)
64
65
       {
           vsmc::NormalDistribution<double> norm pos(0, 2);
66
           vsmc::NormalDistribution<double> norm vel(0, 1);
67
           sp.pos_x() = norm_pos(sp.rng());
68
           sp.pos_y() = norm_pos(sp.rng());
69
           sp.vel_x() = norm_vel(sp.rng());
70
           sp.vel_y() = norm_vel(sp.rng());
71
           w_[sp.id()] = sp.log_likelihood(0);
72
           return 0;
73
```

```
74
       }
       void eval post(vsmc::Particle<PFState> &particle)
 75
 76
       {
            particle.weight().set_log(w_.data());
 77
       }
 78
       private:
79
       vsmc::Vector<double> w ;
 80
 81 };
82 class PFMove : public vsmc::MoveTBB<PFState, PFMove>
 84
       public:
 85
       void eval_pre(std::size_t t, vsmc::Particle<PFState> &particle)
 86
            w_.resize(particle.size());
 87
 88
       }
 89
       std::size_t eval_sp(std::size_t t, vsmc::SingleParticle<PFState> sp)
       {
 90
            vsmc::NormalDistribution<double> norm_pos(0, std::sqrt(0.02));
 91
            vsmc::NormalDistribution<double> norm_vel(0, std::sqrt(0.001));
 92
            sp.pos_x() += norm_pos(sp.rng()) + 0.1 * sp.vel_x();
 93
            sp.pos_y() += norm_pos(sp.rng()) + 0.1 * sp.vel_y();
 94
            sp.vel_x() += norm_vel(sp.rng());
 95
            sp.vel_y() += norm_vel(sp.rng());
96
            w_[sp.id()] = sp.log_likelihood(t);
 97
            return 0;
98
       }
99
       void eval_post(std::size_t t, vsmc::Particle<PFState> &particle)
100
       {
101
            particle.weight().add_log(w_.data());
102
       }
103
       private:
104
```

```
105
       vsmc::Vector<double> w_;
106 };
107 class PFMEval : public vsmc::MonitorEvalTBB<PFState, PFMEval>
108 {
       public:
109
       void eval_sp(std::size_t t, std::size_t dim,
110
            vsmc::SingleParticle<PFState> sp, double *r)
111
112
            r[0] = sp.pos_x();
113
            r[1] = sp.pos_y();
114
       }
115
116 };
117 int main()
118 {
       vsmc::Sampler<PFState> sampler(N, vsmc::Multinomial, 0.5);
119
       sampler.init(PFInit()).move(PFMove(), false).monitor("pos", 2, PFMEval());
120
       sampler.initialize(const_cast<char *>("pf.data")).iterate(n - 1);
121
       std::ofstream output("pf.out");
122
       output << sampler;</pre>
123
       output.close();
124
       return 0;
125
126 }
   A.3 PROCESSING COMMAND LINE PROGRAM OPTIONS
 1 #include <vsmc/vsmc.hpp>
 2 int main(int argc, char **argv)
 3 {
       int n;
 4
       std::string str;
 5
       std::vector<double> vec;
 6
```

```
7
      vsmc::ProgramOptionMap option_map;
8
      option map
           .add("str", "A string option with a default value", &str, "default")
9
           .add("n", "An integer option", &n)
10
           .add("vec", "A vector option", &vec);
11
      option_map.process(argc, argv);
12
      std::cout << "n: " << n << std::endl;</pre>
13
      std::cout << "str: " << str << std::endl;</pre>
14
      std::cout << "vec: ";</pre>
15
16
      for (auto v : vec)
           std::cout << v << ' ';
17
      std::cout << std::endl;</pre>
18
19
      return 0;
20 }
  A.4 DISPLAY PROGRAM PROGRESS
1 #include <vsmc/vsmc.hpp>
2 int main()
3 {
      vsmc::RNG rng;
4
      vsmc::FisherFDistribution<double> dist(10, 20);
5
      std::size t n = 1000;
6
      double r = 0;
7
      vsmc::Progress progress;
8
      progress.start(n * n);
9
      for (std::size t i = 0; i != n; ++i) {
10
           std::stringstream ss;
11
           ss << "i = " << i;
12
           progress.message(ss.str());
13
           for (std::size_t j = 0; j != n; ++j) {
14
               for (std::size_t k = 0; k != n; ++k)
15
                   r += dist(rng);
16
```

progress.increment();

17