SSJ User's Guide

Package markovchain

Structures for the Simulation of Simple Markov chains

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This package offers predefined tools for the simulation of simple Markov chains with Monte Carlo and randomized quasi-Monte Carlo methods. It was developed in part to experiment with the array-RQMC method studied in [1, 2]. In this method, an array of n copies of a Markov chain are simulated in parallel, using RQMC at each step of the chain to induce negative dependence across the copies to reduce the variance of the global average of some performance measure.

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MarkovChain

This class defines a generic Markov chain and provides basic tools to simulate it for a given number of steps and recover the performance measure. Chains can be cloned, so one can simulate many replicates in parallel. In a concrete subclass, it suffices to implement the three abstract methods initialState, nextStep, and getPerformance to get things going.

package umontreal.iro.lecuyer.markovchain;

public abstract class MarkovChain implements Cloneable

Abstract methods

```
public abstract void initialState ()
```

Sets the Markov chain to its (deterministic) initial state and initializes the collectors for the performance measure, ready to start a simulation.

```
public abstract void nextStep (RandomStream stream)
```

Simulates one more step of the chain, from its current state, using **stream** for the randomness.

```
public abstract double getPerformance ()
```

Returns the performance measure (total or average cost or gain) so far, for the current simulation run.

Other methods

public Object clone () throws CloneNotSupportedException

Returns a clone of the chain.

```
public void simulSteps (int numSteps, RandomStream stream)
```

Starts a new simulation run and simulates numSteps steps of the Markov chain, using the given stream.

Performs n simulation runs of the chain, for numSteps steps per run, using the given stream. The statistics on the performance for the n runs are placed in statRuns.

```
public void simulRunsWithSubstreams (int n, int numSteps,
RandomStream stream, Tally statRuns)
```

Same as simulRuns, except that the stream is first reset to its initial seed and then reset to the next substream after each run.

Performs m independent replicates of n simulation runs of the chain, using the points of the RQMC point set p, where n is the number of points in p. Each run goes for numSteps steps. For each replicate, a randomization is added to the point set using the stream noise, an iterator is created, and each run uses a different substream of this iterator (i.e., a different point). The statistics on the performance for the m independent replications are placed in statReps.

Same as simulRuns but also returns the results as a formatted string.

Same as simulRunsWithSubstreams but also returns the results as a formatted string.

Same as simulRQMC but also returns the results as a formatted string.

Similar to simulRQMCFormat, but also gives the variance improvement factor with respect to MC, assuming that varMC gives the variance per run for MC.

public String formatResults (Tally stat)

Returns a string that contains the mean, the variance, and a 90% confidence interval for stat.

```
public String formatResultsRQMC (Tally stat, int numPoints)
```

Returns a string that contains the mean, the variance multiplied by numPoints, and a 90% confidence interval for stat.

MarkovChainComparable

A subclass of Markov chain for which there is a total ordering between the states. A list of Markov chains can then be sorted according to their states at a given step, using **sort**.

package umontreal.iro.lecuyer.markovchain;

Methods

```
public abstract int compareTo (Object other)
```

If the current state of this chain is less than that of other, returns -1; if it is equal, returns 0, otherwise returns 1.

MarkovChainComparableStop

A generic Markov chain which can stop at a (possibly random) stopping time. When the stopping time is reached, the **compareTo** method should conclude that the state of this chain is larger than that of any other chain that has not yet reached its stopping time (so the chain should be placed in a special state considered larger than any other state).

Each chain has a boolean indicator **stopped** that should be set to false before starting the simulation and to **true** (in the **nextStep** method) when the stopping time is reached.

package umontreal.iro.lecuyer.markovchain;

public abstract class MarkovChainComparableStop extends MarkovChainComparable

Methods

public void simulSteps (int numSteps, RandomStream stream)

Starts a new simulation run and stops whenever numSteps are done or the stopping time is reached (the first of these two events), using the given stream.

```
public void simulSteps (RandomStream stream)
```

Starts a new simulation run and simulates until the stopping time is reached, using the given stream. Same as simulSteps (Integer.MAX_VALUE, stream)

MarkovChainDouble

A special kind of Markov chain whose state space is a subset of the real numbers. Methods are provided to initialize the chain, advance by a number of steps, and get statistics on the performance. The method **nextStepDouble** makes it possible to simulate several copies of this chain in parallel without cloning and without maintaining the state of the chain in a local variable. The states can be maintained in an external vector and at each step, one passes the current state to the method **nextStepDouble**, which returns the next state. This is exploited in the implementation of **ArrayOfDoubleChains**.

The methods initialState, nextStep, getPerformance and compareTo, which are abstract in MarkovChainComparable, all have a default implementation that uses internal variables to memorize the *state*, *step number* and *total number of steps*. They update and use these variables by invoking the initialStateDouble, nextStepDouble, and getPerformance(numSteps) methods implemented in subclasses. This is used internally by other methods in MarkovChain. On the other hand, the abstract methods specified in the present class do not necessarily update these variables.

package umontreal.iro.lecuyer.markovchain;

public abstract class MarkovChainDouble extends MarkovChainComparable

Abstract Methods

public abstract double initialStateDouble ()

Same as initialState() but also returns the initial state.

Advances the chain by one step, from state s to the next state and using the current stream, assuming that we are at step step. Saves and returns the new state, but not necessarily the step number. The first call should be with step = 0.

```
public abstract void initStats ()
```

Initializes the statistics (collector) for this chain.

```
public abstract double getPerformance (int numSteps)
```

Returns the performance measure accumulated so far, which may depend on the number of steps.

Other Methods

public double simulStepsDouble (int numSteps, RandomStream stream)

After invoking InitStats starts a new simulation run, simulates numSteps steps of the Markov chain using the given stream, and returns the state. The simulSteps method does the same, but returns nothing.

MarkovChainDoubleStop

Defines a generic Markov chain over the real numbers, where the number of steps of the chain is a (possibly random) stopping time. When the stopping time is reached, the state of the chain is set to Double.POSITIVE_INFINITY.

package umontreal.iro.lecuyer.markovchain;

public abstract class MarkovChainDoubleStop extends MarkovChainDouble

Methods

public double getPerformance ()

Can be used when the performance does not depend on the number of steps.

public double simulStepsDouble (int numSteps, RandomStream stream)

Starts a new simulation run and stops whenever numSteps are done or the stopping time is reached (the first of these two events), using the given stream.

public void simulSteps (int numSteps, RandomStream stream)

Same as simulStepsDouble (numSteps, stream), but returns nothing.

public void simulSteps (RandomStream stream)

Starts a new simulation run and simulates until the stopping time is reached, using the given stream. Same as simulSteps (Integer.MAX_VALUE, stream).

Same as MarkovChain.simulRQMC, except the simulation of any given chain stops whenever it reaches its stopping time.

ArrayOfComparableChains

Permits one to simulate an array of comparable Markov chains using the array-RQMC method of [2], where n copies of the chain are simulated in parallel, sorted in increasing order of their states at each step of the chain, and where the transitions of the n chains at any given step are determined from the n points of a d-dimensional RQMC point set, where d is the number of uniforms required at each step of the chain.

package umontreal.iro.lecuyer.markovchain;

public class ArrayOfComparableChains

Constructor

```
public ArrayOfComparableChains (MarkovChainComparable baseChain)
```

Creates an array of the comparable chain baseChain. The method makeCopies must be called to make the copies.

Methods

```
public void makeCopies (int n)
```

Creates n copies (clones) of the chain baseChain and put them in a list, ready for the array RQMC simulation.

Simulates the n copies of the chain, numSteps steps for each copy, using point set p, where n is the current number of copies (clones) of the chain and is *assumed* to equal the number of points in p. At each step, the points are randomized using noise. The dimension of p must be at least as large as the number of uniforms required to simulate one step of the chain. Returns the average performance per run.

Performs m independent replications of an array-RQMC simulation as in simulArrayRQMC. The statistics on the m corresponding averages are collected in statReps and the results are also returned in a string.

Similar to simulReplicatesArrayRQMC, except that an iterator on a randomized version of p2 is used in place of the noise stream to randomize p at the different steps of the chain. The dimension of p2 must equal the number of uniforms required to randomize p, and its number of points must equal the number of steps in the chain. One point of p2 is used to randomize **p** at each step. The stream **noise** is used to randomize **p2** between the independent replications.

Similar to simulReplicatesArrayRQMC, but also gives the variance improvement factor with respect to MC, assuming that varMC gives the variance per run for MC.

Similar to simulReplicatesArrayRQMC2, but also gives the variance improvement factor with respect to MC, assuming that varMC gives the variance per run for MC.

ArrayOfComparableChainsStop

Similar to ArrayOfComparableChains, except that each chain stops whenever it reaches its stopping time. Simulation and sorting (at each step) is continued only for the chains that have not yet reached their stopping time. This is pursued until either all chains have stopped or the maximum number of steps has been reached.

package umontreal.iro.lecuyer.markovchain;

public class ArrayOfComparableChainsStop extends ArrayOfComparableChains

Constructor

public ArrayOfComparableChainsStop (MarkovChainComparableStop baseChain)

Methods

public double simulArrayRQMC (PointSet p, int numSteps, RandomStream noise)

The simulation of each chain stops whenever it reaches its stopping time or numSteps steps. To simulate each copy of the chain until the stopping time is reached, it suffices to set numSteps = Integer.MAX_VALUE.

ArrayOfDoubleChains

Similar to ArrayOfComparableChains, except that instead of working with n clones of a MarkovChain, we use a *single* MarkovChainDouble object for all the chains. The states of the chains are maintained in an array of real numbers (double) and the nextStepDouble method is used to advance each chain by one step. The performance measure is assumed to be additive over all steps of all copies of the chain. The sum is cumulated in a *single* accumulator for all copies of the chain, updated at each step of each copy.

package umontreal.iro.lecuyer.markovchain;

public class ArrayOfDoubleChains extends ArrayOfComparableChains

Constructor

public ArrayOfDoubleChains (MarkovChainDouble baseChain)

Creates a virtual array for the chain **baseChain**. The method **makeCopies** must be called to make the copies.

Methods

public void makeCopies (int n)

Creates the vector of states for **n** copies of the base chain.

ArrayOfDoubleChainsStop

Similar to ArrayOfDoubleChains, except that each chain stops whenever it reaches its stopping time. When simulating the array of chains, any chain that has reached its stopping time is put in the state ∞ . Simulation and sorting (at each step) is continued only for the chains that have not yet reached their stopping time. This is pursued until either all chains have stopped or the maximum number of steps has been reached.

package umontreal.iro.lecuyer.markovchain;

public class ArrayOfDoubleChainsStop extends ArrayOfDoubleChains

Constructor

public ArrayOfDoubleChainsStop (MarkovChainDoubleStop baseChain)

Methods

Here, the simulation of a chain stops whenever it reaches its stopping time or numSteps steps. To simulate each copy of the chain until the stopping time is reached, it suffices to set numSteps = Integer.MAX_VALUE.

${\bf Left Scrambled Sobol Sequence}$

A Sobol sequence randomized by a left matrix scramble followed by a digital random shift.

package umontreal.iro.lecuyer.markovchain;

public class LeftScrambledSobolSequence extends SobolSequence

Constructor

```
public LeftScrambledSobolSequence (int k, int w, int dim)
```

Same as SobolSequence(k, w, dim), except that its randomize method will do a left matrix scramble followed by a random digital shift.

LeftScrambledFaureSequence

A Faure sequence randomized by a left matrix scramble followed by a digital random shift.

package umontreal.iro.lecuyer.markovchain;

public class LeftScrambledFaureSequence extends FaureSequence

Constructor

public LeftScrambledFaureSequence (int b, int k, int r, int w, int dim)
Same as FaureSequence(b, k, r, w, dim), except that its randomize method will do a
left matrix scramble followed by a random digital shift.

References

- C. Lécot and B. Tuffin. Quasi-Monte Carlo methods for estimating transient measures of discrete time Markov chains. In H. Niederreiter, editor, *Monte Carlo and Quasi-Monte Carlo Methods 2002*, pages 329–343, Berlin, 2004. Springer-Verlag.
- [2] P. L'Ecuyer, C. Lécot, and B. Tuffin. Randomized quasi-Monte Carlo simulation of Markov chains with an ordered state space. In H. Niederreiter and D. Talay, editors, *Monte Carlo and Quasi-Monte Carlo Methods 2004*, 2005. To appear.