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# Recursive Monte Carlo for linear ODEs

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# Contents

**Abstract**

This thesis introduces recursive Monte Carlo for solving linear initial value problems with a vision towards partial differential equations. Further proposing 3 new methods: RRMC an unbiased method for linear initial value problems that has similar convergence behavior in step size to traditional methods, coupled splitting an unbiased technique for linear Fredholm integral equations of the second kind that has similar convergence behavior to fix-point methods and finally a generic unbiased path resampling method similar to walk on spheres applied to recursive Monte Carlo of an IVP obtained by semi-discretizing the heat equation.

# 1   Introduction

## 1.1   Related Work

The primary motivating paper for this work is the work by Sawhney et al. (2022) [Saw+22], which introduces the Walk-on-Sphere (WoS) method for solving second-order elliptic PDEs with varying coefficients and Dirichlet boundary conditions. Their techniques have shown high accuracy even in the presence of geometrically complex boundary conditions. We were inspired to apply the underlying mechanics of these Monte Carlo (MC) techniques to Inital Value Problems (IVPs) to explore parallel in time and the possibility of extending their techniques to other types of PDEs.

We made an interactive data map of the literature read mainly in the function of this thesis available at https://huggingface.co/spaces/ISIPINK/zotero_map.

Existing randomized algorithms for solving IVPs in general are either based on MC integration or randomizing parts of deterministic algorithms. Typically motivated by lower smoothness conditions where randomization adds significant convergence speed or solving ODEs where randomization aids in parallelization .

Daun's 2011 [Dau11] proposes a biased algorithm for IVPs that has optimal Information-Based Complexity (IBC) for some smoothness classes. Similarly to us, their method uses control variates.

Ermakov and Smilovitskiy's 2021 [ES21] proposes an unbiased method for a Cauchy problem for large systems of linear ODEs. Similarly to us, their method is based on Volterra integral equations.

# 2   Background

## 2.1   Monte Carlo Integration

In this subsection, we review basic MC theory.

**Notation 2.1.1** (Random Variables)

Random variables (RVs) will be denoted with capital letters, e.g., $X$, $Y$ or $Z$.

MC integration is any method that involves random sampling to estimate an integral.

**Definition 2.1.2** (Uniform Monte Carlo Integration)

We define uniform MC integration of $f : \mathbb{R} \to \mathbb{R}$ over $[0, 1]$ as an estimation of the expected value of $f(U)$, with $U \sim \text{Uniform}(0, 1)$. Combined with the Best Linear Unbiased Estimators (BLUEs), MC Integration in that case, can be summarized in the following formula:

$$\int_0^1 f(s)ds \approx \frac{1}{n} \sum_{j=1}^n f(U_j), \tag{1}$$

```
1   uniform_MC_int(f, n) = sum(f(rand()) for _ in 1:n) / n
```

where $n$ is the amount of samples used and $U_j$ i.i.d. Uniform$(0, 1)$.

Because estimators are random variables (RVs), the cost and error are also random variables. In most cases, obtaining these RVs is difficult to impossible. Directly comparing estimators based on these RVs can be challenging; there is no simple Pareto front. Instead, comparisons can be made using statistics.

Accuracy comparisons between estimators are typically conducted with (root-)mean-square error (RMSE).

**Definition 2.1.3** (Root-Mean-Square Error)

We define the Root-Mean-Square Error (RMSE) of an estimator $\tilde{\theta}$ for $\theta$ as follows:

$$\text{RMSE}(\tilde{\theta}) = \sqrt{E[||\tilde{\theta} - \theta||_2^2]}. \tag{2}$$

Even comparisons based on RMSE can be counterintuitive; consider Stein's paradox, for example. We limit ourselves to simple cases such as 1-dimensional unbiased estimators, making MSE equivalent to variance. Estimating variance is most cases possible and can be used to calculate confidence intervals using Chebyshev's inequality.

Average floating point operations or time per simulation are common cost statistics. It may also be useful to consider 'at risk' (analogous to 'value at risk') in terms of memory or wall time.

If we limit ourselves to (1) with a big sample size and finite variance assumption on error and simulation time, simulations can be computed in parallel, making them well-suited for a GPU implementation. These assumptions are useful for establishing a baseline and when they are close to being optimal, they become highly practical. In this case, there is a linear trade-off between average simulation time and variance which motivate the definition of MC efficiency for comparing estimators.

**Definition 2.1.4** (Monte Carlo Efficiency)

Define MC efficiency of an estimator $F$ as follows:

$$\epsilon[F] = \frac{1}{\text{Var}(F)T(F)}, \tag{3}$$

with $T$ the average simulation time.

**Related Work 2.1.5** (Monte Carlo Efficiency)

For a reference see [Vea97] page 45.

For smooth 1-dimensional integration, the linear trade-off between variance and average simulation time is not even close to optimal see Theorem 2.4.5.

When it comes to comparing better trade-offs, Information-Based Complexity (IBC) is often employed. IBC primarily serves as a qualitative measure and does not necessarily imply the practicality of an algorithm. We will not delve into a rigorous definition of IBC.

**Definition 2.1.6** (Information-Based Complexity)

IBC is a way to describe asymptotically (for increasing accuracy/function calls) the trade-off between the average amount of function calls (information) needed and accuracy.

**Example 2.1.7** (IBC of (1))

In (1) the function calls trades of linearly with variance. For $n$ function calls, the RMSE $= O\left(\frac{1}{\sqrt{n}}\right)$ or equivalently, if we want a RMSE of $\varepsilon$ we would need $O\left(\frac{1}{\varepsilon^2}\right)$ function calls.

## 2.2 Recursive Monte Carlo

In this subsection, we introduce Recursive Monte Carlo (RMC) with the following initial value problem:

$$y_t = y, \quad y(0) = 1. \tag{4}$$

By integrating both sides of (4), we obtain:

$$y(t) = 1 + \int_0^t y(s)ds. \tag{5}$$

(5) represents a recursive integral equation, specifically, a linear Volterra integral equation of the second type.

**Notation 2.2.1** $(U, U_j)$

We will frequently use the uniform distribution, so we will abbreviate it

$$U_j \text{ i.i.d Uniform}(0,1). \tag{6}$$

The subscripts are used to clarify independence between uniforms. However, when there is no risk of confusion, we simply use $U$.

By estimating the recursive integral in (5) using MC, we derive the following estimator:

$$Y(t) = 1 + ty(Ut). \tag{7}$$

If $y$ is well-behaved, then $E[Y(t)] = y(t)$. However, we cannot directly simulate $Y(t)$ without access to $y(s)$ for $s < t$. Nevertheless, we can replace $y$ with an unbiased estimator without affecting $E[Y(t)] = y(t)$, by the law of total expectation

$(E[X] = E[E[X|Z]])$. By replacing $y$ with $Y$ itself, we obtain a recursive expression for $Y$:

$$Y(t) = 1 + tY(Ut). \qquad (8)$$

```
1   Y(t, eps) = t > eps ? 1 + t * Y(rand() * t, eps) : 1
```

(8) is a Recursive Random Variable Equation (RRVE).

**Definition 2.2.2** (Recursive Random Variable Equation (RRVE))
A Recursive Random Variable Equation (RRVE) is an equation that defines a family of random variables in terms of itself.

Simulation of $Y$ with (8), would recurse indefinitely (every $Y$ needs to sample another $Y$). To stop the recursion, approximate $Y(t) \approx 1$ near $t = 0$ introducing minimal bias. Later, we will discuss Russian roulette; see Definition 2.3.2, which can be used as an unbiased stopping mechanism.

## 2.3   Modifying Monte Carlo

In this subsection, we discuss techniques for modifying RRVEs in a way that preserves the expected value of the solution while acquiring more desirable properties. These techniques are only effective when applied by using prior information about the problem or computational costs.

We will frequently interchange RVs with the same expected values. Therefore introduce following notation:

**Notation 2.3.1** ($\cong$)

$$X \cong Y \iff E[X] = E[Y].$$

Russian roulette is a MC technique commonly employed in rendering algorithms. The concept behind Russian roulette is to replace an RV with a less computationally expensive approximation sometimes.

**Definition 2.3.2** (Russian roulette)
We define Russian roulette on $X$ with free parameters $Y_1 \cong Y_2$, $p \in [0, 1]$ and $U$ independent of $Y_1, Y_2, X$ as follows:

$$X \cong \begin{cases} \frac{1}{p}(X - (1-p)Y_1) & \text{if } U < p \\ Y_2 & \text{else} \end{cases}. \qquad (9)$$

**Notation 2.3.3** ($B(p)$)
Often Russian roulette will be used with $Y_1 = Y_2 = 0$. In that case, we use Bernoulli variables to shorten notation.

$$B(p) \sim \text{Bernoulli}(p) = \begin{cases} 1 & \text{if } U < p \\ 0 & \text{else} \end{cases}. \qquad (10)$$

**Example 2.3.4** (Russian roulette)

Consider the estimation of $E[Z]$, with $Z$:

$$Z = U + \frac{f(U)}{1000}. \tag{11}$$

Here, $f : \mathbb{R} \to [0, 1]$ is expensive to compute. Directly estimating $E[Z]$ would involve evaluating $f$ for each sample which contributes little to the accuracy. To address this, we can modify $Z$ to:

$$Z \cong U + B\left(\frac{1}{100}\right)\frac{f(U)}{10}. \tag{12}$$

This requires calling $f$ on average once every 100 samples. This significantly reduces the computational burden while increasing the variance slightly thereby increasing the overall MC efficiency.

**Related Work 2.3.5** (Example 2.3.4)

In Example 2.3.4, it is also possible to estimate the expectations of the 2 terms of $Z$ separately. Given the variances and computational costs of both terms, you can calculate the asymptotically optimal division of samples for each term. However, this is no longer the case with RMC. In [Rat+22], a method is presented to estimate the optimal Russian roulette/splitting factors for rendering.

**Example 2.3.6** (Russian roulette on (8))

To address the issue of indefinite recursion in (8), Russian roulette can be employed by approximating the value of $Y$ near $t = 0$ with 1 sometimes. Specifically, we replace the coefficient $t$ in front of the recursive term with $B(t)$ when $t < 1$. The modified recursive expression for $Y(t)$ for $t < 1$ becomes:

$$y(t) \cong Y(t) = 1 + B(t)Y(Ut) \tag{13}$$

```
1  Y(t) = rand() < t ? 1 + Y(rand() * t) : 1 # correct for t<1
```

Interestingly, $\forall t \leq 1 : Y(t)$ is the number of recursion calls to sample $Y(t)$ such that the average number of recursion calls to sample $Y(t)$ equals $e^t$.

Splitting is a technique that increases samples of important terms, similar to how Russian roulette decreases the samples of unimportant terms.

**Definition 2.3.7** (splitting)

Splitting $X$ refers to utilizing multiple $X_j \cong X$ (not necessarily independent) to reduce variance by taking their average:

$$X \cong \frac{1}{N}\sum_{j=1}^{N} X_j. \tag{14}$$

Splitting the recursive term in an RRVE can result in additive branching recursion, necessitating cautious management of terminating the branches promptly to prevent exponential growth in computational complexity. To accomplish this, termination

strategies that have been previously discussed can be employed. Subsequently, we will explore the utilization of coupled recursion as a technique to mitigate additive branching recursion in RRVEs (see Example 3.1.7).

**Example 2.3.8** (splitting on (13))

We can "split" the recursive term of (13) into two parts as follows:

$$y(t) \cong Y(t) = 1 + \frac{B(t)}{2}(Y_1(U_1 t) + Y_2(U_1 t)) \tag{15}$$

```
1  function Y(t) # correct for t<1
2      u = rand()
3      rand() < t ? 1 + (Y(u * t) + Y(u * t)) / 2 : 1
4  end
```

$$y(t) \cong Y(t) = 1 + \frac{B(t)}{2}(Y_1(U_1 t) + Y_2(U_2 t)) \tag{16}$$

```
1  Y(t) = rand() < t ? 1 + (Y(rand() * t) + Y(rand() * t)) / 2 : 1
```

where $Y_1(t)$ and $Y_2(t)$ are i.i.d. with $Y(t)$.

**Definition 2.3.9** (control variates)

Define control variating $f(U)$ with $\tilde{f}$ an approximation of $f$ as:

$$f(U) \cong f(U_1) - \tilde{f}(U_1) + E[\tilde{f}(U)] \tag{17}$$
$$= (f - \tilde{f})(U_1) + E[\tilde{f}(U)]. \tag{18}$$

Note that control variating requires the evaluation of $E[\tilde{f}(U)]$. When this is estimated instead, we refer to it as 2-level MC and recursively applying 2-level is multilevel MC.

**Example 2.3.10** (control variate on (8))

To create a control variate for (8) that effectively reduces variance, we employ the approximation $y(t) \approx \tilde{y} = 1 + t$ and define the modified recursive term as follows:

$$y(t) \cong Y(t) = 1 + t(Y(U_1 t) - \tilde{y}(U_1 t) + E[\tilde{y}(Ut)]) \tag{19}$$
$$= 1 + t\left(E[1 + Ut]\right) + t(Y(U_1 t) - 1 - U_1 t) \tag{20}$$
$$= 1 + t + \frac{t^2}{2} + t(Y(U_1 t) - 1 - U_1 t). \tag{21}$$

```
1  function Y(t) # correct for t<1
2      u = rand()
3      1 + t + t^2 / 2 + (rand() < t ? Y(u * t) - 1 - u * t : 0)
4  end
```

Note that while we could cancel out the constant term of the control variate, doing so would have a negative impact on the Russian roulette implemented.

**Related Work 2.3.11** (MC modification)

For further reference on Russian roulette, splitting and control variates see [Vea97].

## 2.4  Monte Carlo Trapezoidal Rule

In this subsection, we introduce an MC trapezoidal rule that exhibits similar convergence behavior to the methods discussed later. The MC trapezoidal rule is essentially a regular Monte Carlo method enhanced with control variates based on the trapezoidal rule.

**Definition 2.4.1** (MC trapezoidal rule)
We define the MC trapezoidal rule for approximating the integral of function $f$ over the interval $[x, x + \Delta x]$ with a Russian roulette rate $l$ and $\tilde{f}$ represents the linear approximation of $f$ corresponding to the trapezoidal rule as follows:

$$\int_x^{x+\Delta x} f(s) ds \tag{22}$$

$$= \int_x^{x+\Delta x} \tilde{f}(s) ds + \int_x^{x+\Delta x} f(s) - \tilde{f}(s) ds \tag{23}$$

$$= \Delta x \frac{f(x) + f(x + \Delta x)}{2} + E\left[ f(S) - \tilde{f}(S) \right] \tag{24}$$

$$\cong \Delta x \frac{f(x) + f(x + \Delta x)}{2}$$
$$+ \Delta x l B \left( \frac{1}{l} \right) \left( f(S) - f(x) - \frac{S - x}{\Delta x} (f(x + \Delta x) - f(x)) \right), \tag{25}$$

where $S \sim \text{Uniform}(x, x + \Delta x)$.

**Lemma 2.4.2** (RMSE MC Trapezoidal Rule)
The MC trapezoidal rule for a twice differentiable function has

$$\text{RMSE} = O\left( \Delta x^3 \right). \tag{26}$$

*Proof.* Start from (25). The MSE is the variance so we can ignore addition by constants.

$$\text{MSE} = \text{Var}\left( \Delta x l B \left( \frac{1}{l} \right) \left( f(S) - f(x) - \frac{S - x}{\Delta x} (f(x + \Delta x) - f(x)) \right) \right) \tag{27}$$

We substitute $S = \Delta x U + x$ and apply Taylor's theorem finishing the proof:

$$\text{MSE} = \text{Var}\left( \Delta x l B \left( \frac{1}{l} \right) (f(\Delta x U + x) - f(x) - U(f(x + \Delta x) - f(x))) \right) \tag{28}$$

$$= \text{Var}\left( \Delta x l B \left( \frac{1}{l} \right) \left( U \Delta x f'(x) + \frac{U^2 \Delta x^2}{2} f''(Z_1) - U \left( \Delta x f'(x) + \Delta x^2 f''(z_2) \right) \right) \right) \tag{29}$$

$$= \text{Var}\left( \Delta x l B \left( \frac{1}{l} \right) \left( \frac{U^2 \Delta x^2}{2} f''(Z_1) - \frac{U \Delta x^2}{2} f''(z_2) \right) \right) \tag{30}$$

$$= \Delta x^6 \text{Var}\left( l B \left( \frac{1}{l} \right) \left( \frac{U^2}{2} f''(Z_1) - \frac{U}{2} f''(z_2) \right) \right), \tag{31}$$

for some $Z_1 \in [x, S], z_2 \in [x, x + \Delta x]$. The variance term is bounded because the variance of a bounded RV is bounded. Note that the proof does not rely on Russian roulette ($l = 1$). $\qquad\square$

**Related Work 2.4.3** (proof of Lemma 2.4.2)

A more generalizable proof for other types of control variates can be constructed by applying the 'separation of the main part' technique, as shown in Lemma 4 of [HM93].

**Definition 2.4.4** (composite (MC) trapezoidal rule)

Define the MC trapezoidal rule for approximating the integral of function $f$ over the interval $[0, 1]$ with a uniform grid with $n$ intervals as follows:

$$\int_0^1 f(s)ds \approx \Delta x \sum_{j=0}^{n-1} \frac{f(x_j) + f(x_j + \Delta x)}{2}. \tag{32}$$

```
1  trapezium(f, n) = sum((f(j / n) + f((j + 1) / n)) / 2 for j in 0:n-1) / n
```

Define the corresponding composite MC trapezoidal with a Russian roulette rate $l$ as follows:

$$\int_0^1 f(s)ds \cong \Delta x \sum_{j=0}^{n-1} \frac{f(x_j) + f(x_j + \Delta x)}{2}$$
$$+ lB\left(\frac{1}{l}\right)\left(f(S_j) - f(x_j) - \frac{S_j - x_j}{\Delta x}(f(x_j + \Delta x) - f(x_j))\right), \tag{33}$$

```
1  function MCtrapezium(f, n, l=100)
2      sol = 0
3      for j in 0:n-1
4          if rand() * l < 1
5              x, xx = j / n, (j + 1) / n
6              S = x + rand() * (xx - x) # \sim Uniform(x, xx)
7              sol += l * (f(S) - f(x) - n * (S - x) * (f(xx) - f(x))) / n
8          end
9      end
10     return trapezium(f, n) + sol
11 end
```

where $S_j \sim \text{Uniform}(x, x + \Delta x)$.

Figure 1 suggests that the order of convergence of RMSE of the composite MC trapezoidal rule is better by 0.5 than the normal composite trapezoidal rule. The MC trapezoidal rule has on average $\frac{1}{l}$ more function calls than the normal trapezoidal rule. For the composite rule with $n$ intervals, there are $\text{Binomial}(n, \frac{1}{l})$ additional function calls (repeated Bernoulli experiments).

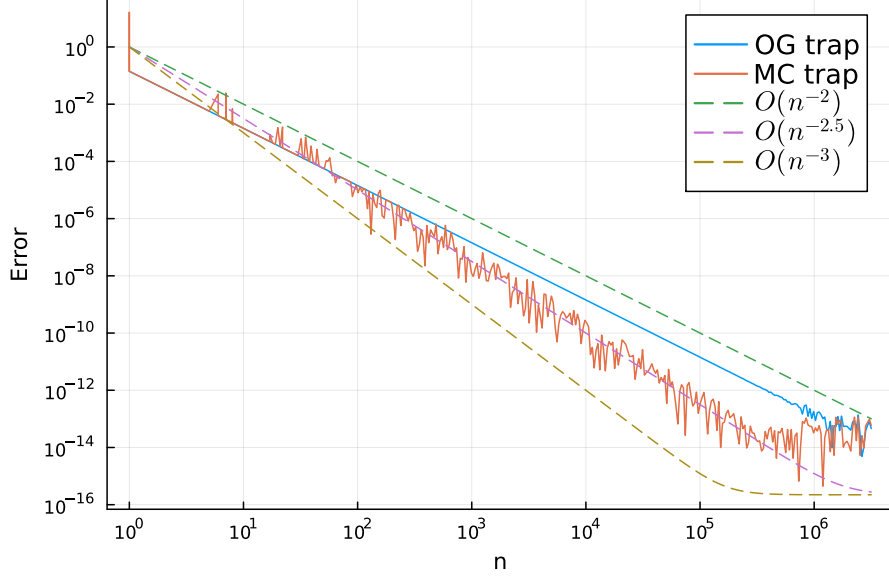**Theorem 2.4.5** (RMSE Composite Trapezoidal MC Rule)

*Figure 1: Log-log plot of the error of (33) for $\int_0^1 e^s ds$ with $l = 100$. At floating point accuracy, the convergence ceases.*

The composite trapezoidal MC rule with $n$ intervals for a twice differentiable function has

$$\text{RMSE} = O\left(\frac{1}{n^{2.5}}\right). \tag{34}$$

The proof uses Lemma 2.4.2, and is similar to the proof of the normal trapezoidal rule. The main difference is the accumulation of 'local truncation' errors into 'global truncation' error. Normally there is a loss of one order but the MC trapezoidal loses only a half order because the accumulation happens in variance instead of bias.

$$\sqrt{\text{Var}\left(\sum_{j=1}^n \Delta x^3 U_j^2\right)} = \Delta x^3 \sqrt{\sum_{j=1}^n \text{Var}(U_j^2)} \tag{35}$$

$$= \Delta x^3 \sqrt{n\text{Var}(U^2)} \tag{36}$$

$$= O(\Delta x^{2.5}). \tag{37}$$

Note that the meaning of a bound on the error, which behaves as $O(\Delta x^2)$, and a bound on the RMSE, also behaving as $O(\Delta x^2)$, is different. A bound on the error implies a bound on the RMSE, but not vice versa.

**Related Work 2.4.6** (Monte Carlo Trapezoidal Rule)

Due to an argument like Stein's paradox, it is always possible to bias the composite MC trapezoidal rule to achieve lower RMSE. The optimal IBC for the deterministic, random, and quantum integration are known for some smoothness classes; see [HN01] for details.

## 2.5 Unbiased Non-Linearity

In this subsection, we present techniques for handling polynomial non-linearity. The main idea behind this is using independent samples $y^2 \cong Y_1 Y_2$ with $Y_1$ independent of $Y_2$ and $Y_1 \cong Y_2 \cong y$.

**Example 2.5.1** $(y_t = y^2)$

Consider the following ODE:

$$y_t = y^2, \quad y(1) = -1. \tag{38}$$

The solution to this equation is given by $y(t) = -\frac{1}{t}$. By integrating both sides of (38), we obtain the following integral equation:

$$y(t) = -1 + \int_1^t y(s)y(s)ds. \tag{39}$$

To estimate the recursive integral in (38), we use $Y_1, Y_2$ i.i.d. $Y$ in following RRVE:

$$y(t) \cong Y(t) = -1 + (t-1)Y_1(S_1)Y_2(S_1), \tag{40}$$

```
1  function Y(t) # Y(u)^2 != Y(u) * Y(u) !!!
2      t > 2 && error("doesn't support t > 2")
3      S1 = rand() * (t - 1) + 1
4      rand() < t - 1 ? -1 + Y(S1) * Y(S1) : -1
5  end # Y(t) = 0 or Y(t) = -1
```

where $S_1 \sim \text{Uniform}(1, t)$.

**Example 2.5.2** $(e^{E[X]})$

$e^{\int x(s)ds}$ is a common expression encountered when studying ODEs. In this example, we demonstrate how you can generate unbiased estimates of $e^{E[X]}$ with simulations of $X$. The Taylor series of $e^x$ is:

$$e^{E[X]} = \sum_{n=0}^{\infty} \frac{E^n[X]}{n!} \tag{41}$$

$$= 1 + \frac{1}{1}E[X]\left(1 + \frac{1}{2}E[X]\left(1 + \frac{1}{3}E[X]\left(1 + \dots\right)\right)\right). \tag{42}$$

Change the fractions of (42) to Bernoulli processes and replace all $X$ with independent $X_j \cong X$.

$$e^{E[X]} = E\left[1 + B\left(\frac{1}{1}\right)E[X_1]\left(1 + B\left(\frac{1}{2}\right)E[X_2]\left(1 + B\left(\frac{1}{3}\right)E[X_3]\left(1 + \dots\right)\right)\right)\right] \tag{43}$$

$$\cong 1 + B\left(\frac{1}{1}\right)X_1\left(1 + B\left(\frac{1}{2}\right)X_2\left(1 + B\left(\frac{1}{3}\right)X_3\left(1 + \dots\right)\right)\right). \tag{44}$$

```
1  num_B(i) = rand() * i < 1 ? num_B(i + 1) : i - 1
2  res(X, n) = (n != 0) ? 1 + X() * res(X, n - 1) : 1
3  expE(X) = res(X, num_B(0))
```

Sampling (44) requires a finite amount of samples from $X_j$'s with probability 1.

**Related Work 2.5.3** (Example 2.5.2)

NVIDIA has a great paper on optimizing Example 2.5.2 [Ket+21].

## 2.6 Recursion

In this subsection, we discuss recursion-related techniques.

**Technique 2.6.1** (coupled recursion)

The idea behind coupled recursion is sharing recursion calls of multiple RRVEs for simulation. This does make them dependent.

**Example 2.6.2** (coupled recursion)

Consider calculating the sensitivity of following ODE to a parameter $a$:

$$y_t = ay, y(0) = 1 \Rightarrow \tag{45}$$

$$\partial_a y_t = y + a\partial_a y \tag{46}$$

Turn (45) and (46) into RRVEs. To emphasize that they are coupled and should recurse together we write them in a matrix equation:

$$\begin{bmatrix} y(t) \\ \partial_a y(t) \end{bmatrix} \cong \begin{bmatrix} Y(t) \\ \partial_a Y(t) \end{bmatrix} = X(t) = \begin{bmatrix} 1 \\ 0 \end{bmatrix} + t \begin{bmatrix} a & 0 \\ 1 & a \end{bmatrix} X(Ut). \tag{47}$$

```
1  (q = [1, 0]; A(a) = [a 0; 1 a])
2  X(t, a) = (rand() < t) ? q + A(a) * X(rand() * t, a) : q
```

Observe how this eliminates the additive branching recursion present in (46).

**Technique 2.6.3** (recursion in recursion)

Later we will use recursion in recursion which can get involved. Recursion in recursion is like proving an induction step of an induction proof with induction.

**Related Work 2.6.4** (recursion in recursion)

A beautiful example of recursion in recursion is the next flight variant of WoS in [Saw+22].

Most programming languages do support recursion, but it often comes with certain limitations such as maximum recursion depth and potential performance issues. Non-branching recursion can be avoided by directly traversing the computational path in the reverse order. Poisson time processes can be sampled in the reverse order because of equal forward paths and backward paths distributions.

**Definition 2.6.5** (main Poisson)

Consider the following linear IVP with $\forall s > 0 \in \mathbb{R} : f(s) \in \mathbb{R}^n, A(s) \in \mathbb{R}^{n \times n}$:

$$y' = Ay + f \Leftrightarrow \tag{48}$$

$$y' + \sigma y = (A + \sigma I)y + f \Leftrightarrow \tag{49}$$

$$e^{-\sigma t}(e^{\sigma t}y)' = (A + \sigma I)y + f \Leftrightarrow \tag{50}$$

$$y(t) = e^{-\sigma t}y(0) + \int_0^t e^{(s-t)\sigma} \left( (A(s) + \sigma I)y(s) + f(s) \right) ds. \tag{51}$$

With $\sigma > 0 \in \mathbb{R}$. Do following substitution: $e^{(s-t)\sigma} = \tau$ equivalent to importance sampling the exponential term:

$$y(t) = \int_0^{e^{-\sigma t}} y(0) d\tau + \int_{e^{-\sigma t}}^1 \left( \frac{A(s)}{\sigma} + I \right) y(s) + \frac{f(s)}{\sigma} d\tau. \tag{52}$$

Sampling $\tau$ uniformly is equivalent to sampling a Poisson process.

**Julia Code 2.6.6** (implementation of 2.6.5)

(52) can be turned into a RRVE by sampling $\tau \sim U$, no Russian roulette is needed for termination because in the recursion the first integral get sampled with probability 1. No recursion is used for the implementation as we can sample the Poisson process in the reverse order.

```
1  function Y(t, sig, A::Function, f::Function, y0)
2      (s = -log(rand()) / sig; sol = y0)
3      while s < t
4          sol += (A(s) * sol .+ f(s)) ./ sig
5          s -= log(rand()) / sig
6      end
7      sol
8  end
```

**Related Work 2.6.7** (main Poisson)

A very similar estimator is used to solve ODEs derived from the 1-dimensional telegraph equations in [AR16]. Unlike 4.0.3 the recursion process they use can be reversed because the paths consist of a Poisson process and reflections.

Tail recursion is commonly used in rendering to implement non-branching recursion.

**Technique 2.6.8** (non-branching tail recursion)

Tail recursion involves reordering all operations so that almost no operation needs to happen after the recursion call. This allows us to return the answer without retracing all steps when we reach the last recursion call.

The non-branching recursion presented in the RRVEs can be implemented using tail recursion due to the associativity of all operations $((xy)z = x(yz))$ involved.

**Julia Code 2.6.9** (tail recursion for 2.6.5)

We collect the sum in the variable sol and the matrix multiplications in W. Notice that the computation is going backwards in time.

```
1  using LinearAlgebra
2  function Y(t, sig, A::Function, f::Function, y0)
3      (sol = zero(y0); W = I; s = t + log(rand()) ./ sig)
4      while 0 < s
5          sol += W * f(s) ./ sig
6          W += W * A(s) ./ sig
7          s += log(rand()) ./ sig
8      end
9      sol + W * y0
10 end
```

Tail recursion loses the intermediate values of the recursive calls and may increase computational costs because of the reordering of operations. The computational cost of matrix multiplications are sensitive to reordering operations.

**Related Work 2.6.10** (non-branching tail recursion)

[VSJ21] proposes an efficient unbiased backpropagation algorithm for rendering exploiting properties of non-branching tail recursion by inverting local Jacobian matrices which is only feasible for small matrices. To access intermediate values in tail recursion they use path replay.

The expensive matrix multiplications disappear when solving for $v^T y(t)$.

**Julia Code 2.6.11** (adjoint tail recursion for 2.6.5)

We collect the sum in the variable sol and instead of matrix multiplications in W matrix-vector multiplications in $v$.

```
function Yadj(t, sig, A::Function, f::Function, y0, v) # unbiased v'*y
    (sol = 0; s = t + log(rand()) ./ sig)
    while 0 < s
        sol += dot(v, f(s)) ./ sig
        v += v * A(s) ./ sig
        s += log(rand()) ./ sig
    end
    sol + dot(v, y0)
end
```

We are interested in methods that can keep $v$ sparse. When $(v)_j = \delta_{jn}$ we obtain an estimator for $(y(t))_n$.

# 3 Ordinary Differential Equations

## 3.1 Green's Functions

In this subsection, we discuss how to transform ODEs into integral equations and subsequently solve them. Our main tool for this is Green's functions.

A Green's function is a type of kernel function used for solving linear problems with linear conditions. In this context, Green's functions are analogous to homogeneous and particular solutions to a specific problem, which are combined through integration to solve more general problems.

**Related Work 3.1.1** (Green's function)

Our notion of the Green's function is similar to that presented in [HGM01].

**Notation 3.1.2** ($H$)

We denote the Heaviside step function with:

$$H(x) = \begin{cases} 0 & \text{if } x < 0 \\ 1 & \text{else} \end{cases}. \tag{53}$$

**Notation 3.1.3** ($\delta$)

We denote the Dirac delta function as $\delta(x)$.

To clarify Green's functions, let us look at the following example.

**Example 3.1.4** ($y_t = y$ average condition)

Consider equation:

$$y_t = y, \tag{54}$$

with following condition making the solution unique:

$$\int_0^1 y(s)ds = e - 1. \tag{55}$$

The solution to this equation remains the same: $y(t) = e^t$. We define the corresponding source Green's function $G(t, x)$ for $y_t$ and this type of condition as follows:

$$G_t = \delta(x - t), \quad \int_0^1 G(s, x)ds = 0. \tag{56}$$

Solving this equation yields:

$$G(t, x) = H(t - x) + x - 1. \tag{57}$$

We define the corresponding boundary Green's function $G(t, x)$ for $y_t$ and this type of condition as follows:

$$P_t = 0, \quad \int_0^1 P(s)ds = e - 1. \tag{58}$$

Solving this equation yields:

$$P(t) = e - 1. \tag{59}$$

The Green's functions are constructed to form the following integral equation for (54):

$$y(t) = P(t) + \int_0^1 G(t, s)y(s)ds. \tag{60}$$

Converting (60) into an RRVE using RMC, we obtain:

$$y(t) \cong Y(t) = e - 1 + 2B\left(\frac{1}{2}\right)Y(S)(H(t - S) + S - 1), \tag{61}$$

where $S \sim U$. We plot realizations of (61) in Figure 2.

(60) is a Fredholm integral equation of the second kind.

**Definition 3.1.5** (Fredholm equation of the second kind)

A Fredholm equation of the second kind for $\varphi$ is of the following form:

$$\varphi(t) = f(t) + \lambda \int_a^b K(t, s)\varphi(s)ds. \tag{62}$$

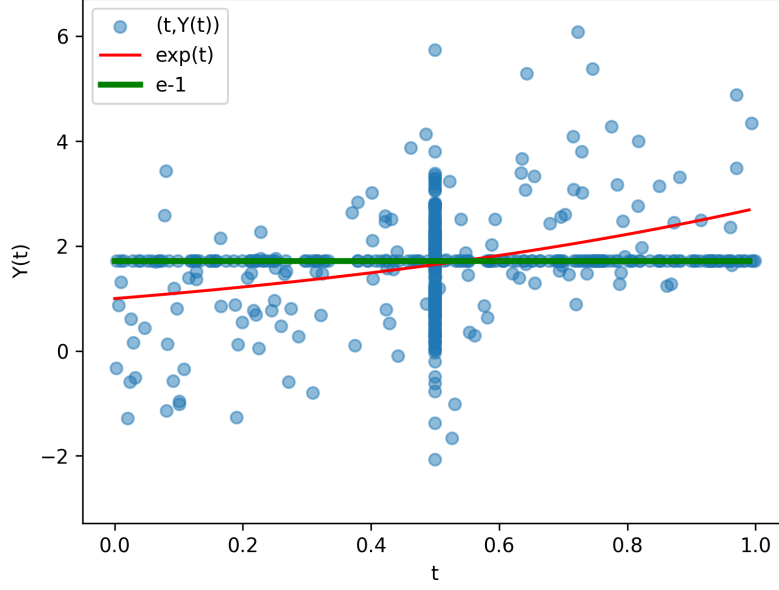Here, $K(t, s)$ represents a kernel, and $f(t)$ is a given function.

*Figure 2: Recursive calls $(t, Y(t))$ of ([61](#)) when calling $Y(0.5)$ 300 times. Points accumulate on the Green's line due to the Russian roulette, and at $t = 0.5$ because of the first recursion call.*

If both $K$ and $f$ satisfy certain regularity conditions, then for sufficiently small $\lambda$, it is relatively straightforward to establish the existence and uniqueness of solutions using a fixed-point argument.

**Example 3.1.6** (Dirichlet $y_{tt} = y$)

Consider following BVP:

$$y_{tt} = y. \tag{63}$$

With boundary conditions $y(b_0), y(b_1)$. The Green's functions corresponding to $y_{tt}$ and Dirichlet conditions are:

$$P(t|x) = \begin{cases} \frac{b_1 - t}{b_1 - b_0} & \text{if } x = b_0 \\ \frac{t - b_0}{b_1 - b_0} & \text{if } x = b_1 \end{cases}, \tag{64}$$

$$G(t, s) = \begin{cases} -\frac{(b_1 - t)(s - b_0)}{b_1 - b_0} & \text{if } s < t \\ -\frac{(b_1 - s)(t - b_0)}{b_1 - b_0} & \text{if } t < s \end{cases}. \tag{65}$$

Directly from these Green's functions, we obtain the following integral equation and RRVE:

$$y(t) = P(t|b_0)y(b_0) + P(t|b_1)y(b_1) + \int_{b_0}^{b_1} G(t, s)y(s)ds, \tag{66}$$

$$Y(t) = P(t|b_0)y(b_0) + P(t|b_1)y(b_1) + lB\left(\frac{1}{l}\right)(b_1 - b_0)G(t, S)y(S), \tag{67}$$

with the Russian roulette rate $l \in \mathbb{R}$ and $S \sim \text{Uniform}(b_0, b_1)$. In Figure [3](#) and [4](#) we test convergence of ([67](#)).
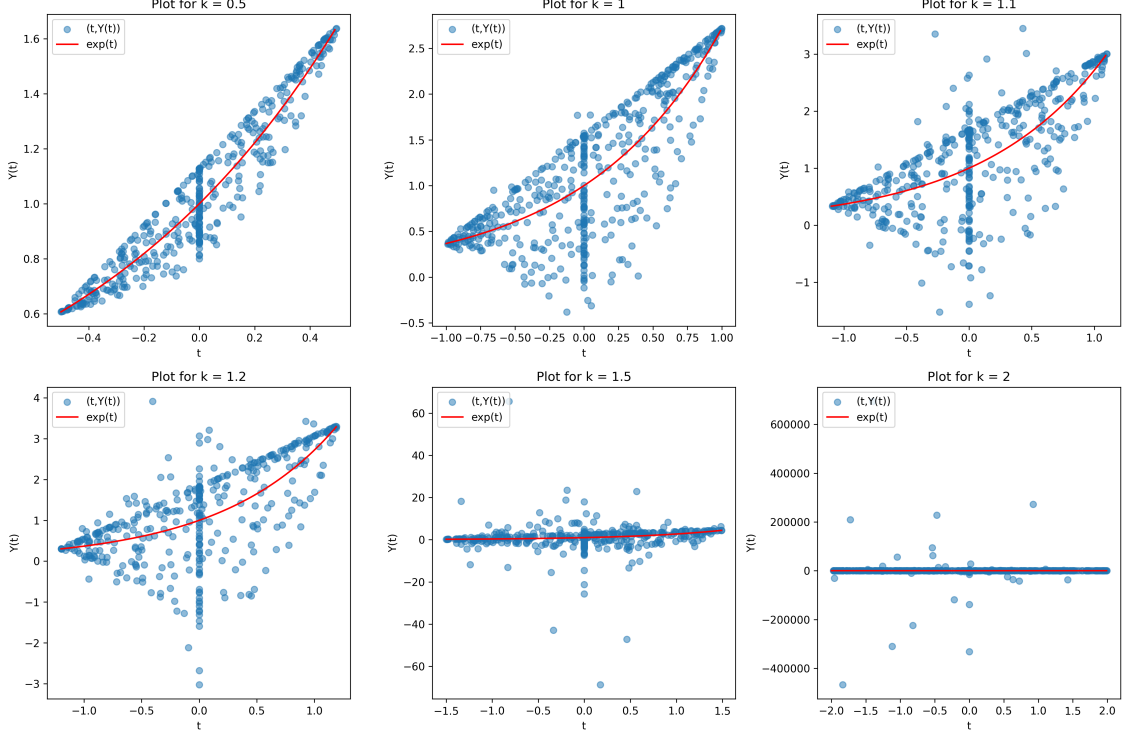
*Figure 3: Recursive calls $(t, Y(t))$ of (67) when calling $Y(0)$ 75 times, with $l = 1.2$ and boundary conditions $y(-k) = e^{-k}$ and $y(k) = e^k$.*

We tested coupled splitting on Example 3.1.6. Coupled splitting removes the additive branching of splitting by coupling (reusing) samples.

**Example 3.1.7** (coupled splitting on Example 3.1.6)

In addition to normal splitting (see definition 2.3.7), we can also split the domain in (66) as follows:

$$y(t_1) = P(t_1|b_0)y(b_0) + P(t_1|b_1)y(b_1) \tag{68}$$

$$+ \int_{b_0}^{\frac{b_0+b_1}{2}} G(t_1, s_1)y(s_1)ds_1 + \int_{\frac{b_0+b_1}{2}}^{b_1} G(t_1, s_2)y(s_2)ds_2 \tag{69}$$

$$y(t_2) = P(t_2|b_0)y(b_0) + P(t_2|b_1)y(b_1) \tag{70}$$

$$+ \int_{b_0}^{\frac{b_0+b_1}{2}} G(t_2, s_1)y(s_1)ds_1 + \int_{\frac{b_0+b_1}{2}}^{b_1} G(t_2, s_2)y(s_2)ds_2. \tag{71}$$

By coupling, we can eliminate the additive branching recursion in the RRVEs corresponding to (68) and (70). In other words, to estimate $y(t_1)$ and $y(t_2)$ we need to estimate $y(s_1)$ and $y(s_2)$ instead of doing it twice we reuse the same estimate. This results in the following RRVE:

$$X(t_1, t_2) = \begin{bmatrix} P(t_1|b_0) & P(t_1|b_1) \\ P(t_2|b_0) & P(t_2|b_1) \end{bmatrix} \begin{bmatrix} y(b_0) \\ y(b_1) \end{bmatrix} + W \begin{bmatrix} G(t_1, S_1) & G(t_1, S_2) \\ G(t_2, S_1) & G(t_2, S_2) \end{bmatrix} X(S_1, S_2), \tag{72}$$
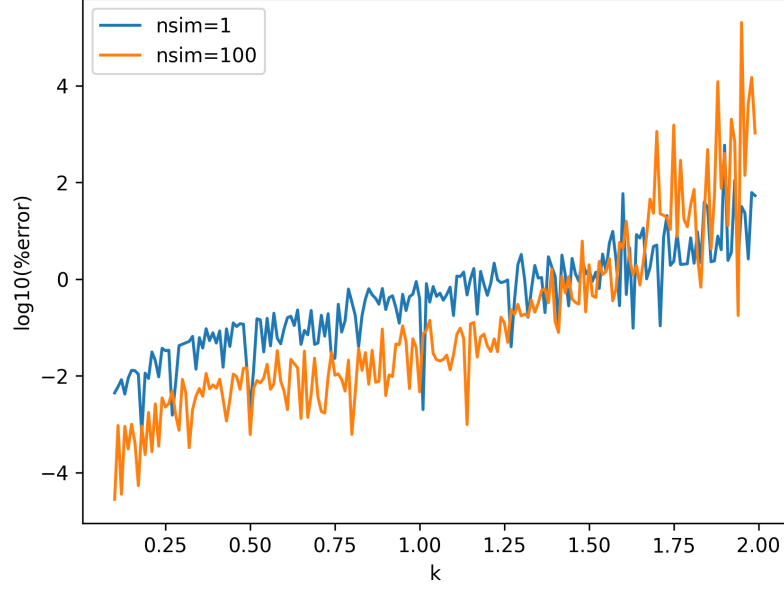
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*Figure 4: The logarithmic percentage error of $Y(0)$ for (67), with $l = 1.2$ and boundary conditions $y(-k) = e^{-k}$ and $y(k) = e^k$, displays an exponential increase until approximately $k = 1.5$, beyond which additional simulations fail to reduce the error, indicating that the variance does not exist.*

```
1   Pb0(t, b0, b1) = (b1 - t) / (b1 - b0)
2   Pb1(t, b0, b1) = (t - b0) / (b1 - b0)
3   G(t, s, b0, b1) = (t < s) ?
4                     -(b1 - s) * (t - b0) / (b1 - b0) :
5                     -(b1 - t) * (s - b0) / (b1 - b0)
6   function X(T::Array, y0, y1, b0, b1, l)
7       PP = hcat([Pb0(t, b0, b1) for t in T], [Pb1(t, b0, b1) for t in T])
8       sol = PP * [y0, y1]
9       if rand() * l < 1
10          (u = rand(); n = length(T))
11          SS = [b0 + (u + j) * (b1 - b0) / n for j in 0:n-1]
12          GG = reshape([G(t, S, b0, b1) for t in T, S in SS], n, n)
13          sol += l * GG * X(SS, y0, y1, b0, b1) .* (b1 - b0) ./ n
14      end
15      sol
16  end
```

with $W$ some weighting matrix, $S_1$ and $S_2$ can be chosen in various ways. (72) is unbiased in the following way: $X(t_1, t_2) \cong [y(t_1) \quad y(t_2)]^T$.

**Related Work 3.1.8** (coupled splitting)

Coupled splitting is partly inspired by the approach of [SM09], which reduces variance by using larger submatrices in unbiased sparsification of matrices. Reusing samples for WoS is discussed in both [Mil+23] and [BP23].

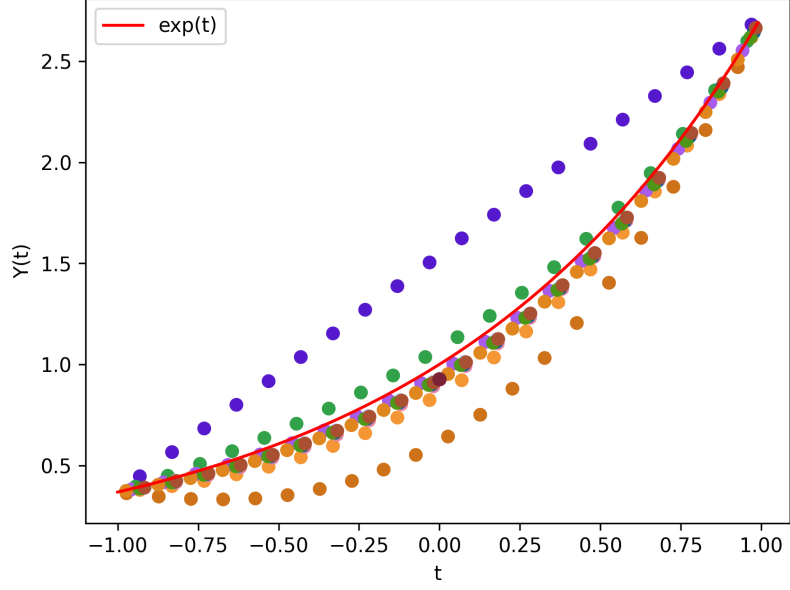Figure 5 resembles fixed-point iterations, leading us to hypothesize that the conver-

*Figure 5: Recursive calls of ([72](#)) when calling $X(0)$ once. We chose $X$ to have $20$ points and colored each call. The $S_j$ are coupled such that they are equally spaced. The initial conditions for this call are $y(-1) = e^{-1}$ and $y(1) = e^1$, with Russian roulette rate $l = 1.2$.*

gence speed is very similar to fix-points methods until the accuracy of the stochastic approximation of the operator is reached (the approximate operator bottleneck). The approximation of the operator can be improved by increasing the coupled splitting amount when approaching the bottleneck. Alternatively when reaching the bottleneck it is possible to rely on splitting to converge.

**Related Work 3.1.9** (convergence coupled splitting)

See [GH21] for a discussion on the convergence of recursive stochastic algorithms.

**Related Work 3.1.10** (IBC integral equations)

Optimal IBC is known for integral equations see [Hei98] for the solution at 1 point and the global solution.

## 3.2  Initial Value Problems

Classic IVP solvers rely on shrinking the time steps for convergence. In this subsection, we introduce Recursion in Recursion MC (RRMC) for IVPs which tries to emulate this behavior.

**Example 3.2.1** (RRMC $y_t = y$)

We demonstrate RRMC for IVPs with

$$y_t = y, \quad y(0) = 1. \tag{73}$$

Imagine we have a time-stepping scheme $(t_n), \forall n : t_{n-1} < t_n$ then the following integral equations hold:

$$y(t) = y(t_n) + \int_{t_n}^{t} y(s)ds, \quad t > t_n. \tag{74}$$

Turn these in the following class of RRVEs:

$$y(t) \cong Y_j(t) = y(t_j) + (t - t_j)Y_j((t - t_j)U + t_j), \quad t > t_j. \tag{75}$$

A problem with these RRVEs is that we do not know $y(t_j)$. Instead, we can replace it with an unbiased estimate $y_j$ which we keep fixed in the inner recursion:

$$y(t) \cong Y_j(t) = y_j + (t - t_j)Y_j((t - t_j)U + t_j), \quad t > t_j \tag{76}$$

$$y(t_j) \cong y_j = \begin{cases} Y_{j-1}(t_j) & \text{if } j \neq 0 \\ y(t_0) & \text{if } j = 0 \end{cases}. \tag{77}$$

```
1  function Y_in(t, tn, yn, h)
2      S = tn + rand() * (t - tn)   # \sim Uniform(T,t)
3      return rand() < (t - tn) / h ? yn + h * Y_in(S, tn, yn, h) : yn
4  end
5  function Y_out(tn, h) # h is out step size
6      TT = tn > h ? tn - h : 0
7      return tn > 0 ? Y_in(tn, TT, Y_out(TT, h), h) : 1
8  end
```

We refer to (76) as the inner recursion and (77) as the outer recursion of the recursion in recursion. The measured RMSE for estimating $y(t)$ is of the order $O(h^{1.5})$, where $h$ represents the step size. For the implementation we used a scaled version of the time process from Example 2.3.6 for the inner recursion, such that the average number of total inner recursion calls is $en$, where $n$ represents the total number of outer recursion calls.

**Conjecture 3.2.2** (local RMSE RMC)
Consider a general linear IVP:

$$y_t(t) = A(t)y(t) + g(t), y(t_0), \tag{78}$$

with $A$ a matrix and $g$ a vector function, each once differentiable with the corresponding RRVE:

$$y(t) \cong Y(t) = y(t_0) + hB\left(\frac{t - t_0}{h}\right)A(S)Y(S) + g(S), \tag{79}$$

where $S \sim \text{Uniform}(t_0, t)$. Then, the following relation holds:

$$E[||y(t_1) - Y(t_1)||^2] = O(h^2), \tag{80}$$

with $t_1 = t_0 + h$.

To achieve a higher order of convergence RRMC can be combined with control variates.

**Example 3.2.3** (CV RRMC $y_t = y$)

Let us control variate Example 3.2.1.

$$y(t) = y(t_n) + \int_{t_n}^{t} y(s)ds, \quad t > t_n. \tag{81}$$

We build a control variate with a lower-order approximation of the integrand:

$$y(s) = y(t_n) + (s - t_n)y_t(t_n) + O((s - t_n)^2) \tag{82}$$

$$\approx y(t_n) + (s - t_n)f(y(t_n), t_n) \tag{83}$$

$$\approx y(t_n) + (s - t_n)\left(\frac{y(t_n) - y(t_{n-1})}{t_n - t_{n-1}}\right) \tag{84}$$

$$\approx y(t_n)(1 + s - t_n). \tag{85}$$

Using (85) as a control variate for the integral:

$$y(t) = y(t_n) + \int_{t_n}^{t} y(s)ds \tag{86}$$

$$= y(t_n) + \int_{t_n}^{t} y(s) - y(t_n)(1 + s - t_n) + y(t_n)(1 + s - t_n)ds \tag{87}$$

$$= y(t_n)\left(1 + (1 - t_n)(t - t_n) + \frac{t^2 - t_n^2}{2}\right) + \int_{t_n}^{t} y(s) - y(t_n)(1 + s - t_n)ds. \tag{88}$$

Figure 6 displays the error of realizations of an RRVE constructed from (88) for various step sizes.
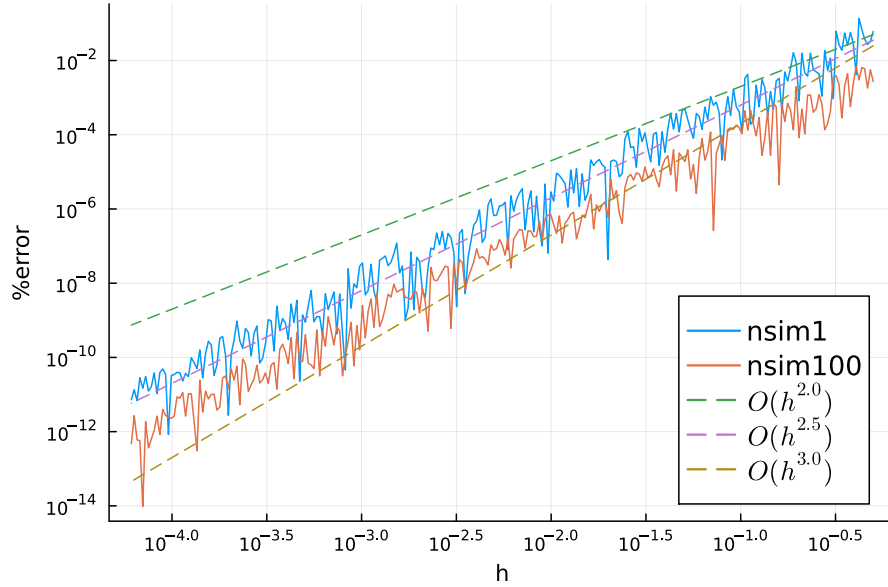


*Figure 6: Log-log plot of the error for Example 3.2.3 at $Y(10)$.*

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**Related Work 3.2.4** (CV RRMC)

[Dau11] similarly uses control variates to achieve a higher order of convergence.

Similar to explicit solvers, RRMC performs poorly on stiff problems. We attempted to make RRMC more like implicit solvers but this proved difficult. Instead, we believe that an approach more like exponential integrators is more viable.

# 4 Heat Equation

In this section we discuss how to solve ODEs derived from the semi-discretized heat equation with Dirichlet boundary conditions in 1 point and acceleration with recursive first-passage sampling combined with presampled path stitching.

**Definition 4.0.1** (1D heat equation Dirichlet)

We define the 1D heat equation for $u(x,t)$ with $(x,t) \in \Omega = [0,1]^2$ with Dirichlet boundary conditions the following way:

$$\frac{\partial u}{\partial t} = \frac{\partial^2 u}{\partial x^2} \Leftrightarrow \tag{89}$$

$$u_t(x,t) = u_{xx}(x,t), \quad \forall (x,t) \in \Omega = [0,1]^2. \tag{90}$$

Given $\{u(x,0), u(1,t), u(0,t) | \forall x, t \in [0,1]\}$.

**Definition 4.0.2** (semi-discretization 1D heat equation Dirichlet)

We define the semi-discretization 1D heat equation for $\tilde{u}$ as space discretized heat equation using the central difference scheme:

$$\tilde{u}_t(j\Delta x, t) = \frac{\tilde{u}((j+1)\Delta x, t) - 2\tilde{u}(j\Delta x, t) + \tilde{u}((j-1)\Delta x, t)}{\Delta x^2} \tag{91}$$

$\forall (j,t) \in \{0, 1, \ldots, N\} \times [0,1]$.
Given $\{\tilde{u}(j\Delta x, 0), \tilde{u}(1,t), \tilde{u}(0,t) | \forall j \in \{0, 1, \ldots, N\}, \forall t \in [0,1]\}$.

It is well-known that the solution to the semi-discretization 1D heat equation converges to the solution of the 1D heat equation with corresponding boundary conditions as $\Delta x \to 0$.
We want a point estimator for the solution to the semi-discretization 1D heat equation. We obtain this by using (52) with $\sigma = \frac{2}{\Delta x^2}$ and at the interior point $j$ this is:

$$\tilde{u}(j\Delta x, t) = \int_0^{e^{-\frac{2t}{\Delta x^2}}} \tilde{u}(j\Delta x, 0) d\tau + \int_{e^{-\frac{2t}{\Delta x^2}}}^1 \frac{\tilde{u}((j+1)\Delta x, s) + \tilde{u}((j-1)\Delta x, s)}{2} d\tau. \tag{92}$$

**Julia Code 4.0.3** (implementation of estimator of (92))

(92) can be turned into a estimator $Y(j\Delta x, t, \Delta x) \cong \tilde{u}(j\Delta x, t)$ by sampling $\tau \sim U$, sampling one of the $\tilde{u}$ of the second integral to avoid branching recursion and recursing when $\tilde{u}$ is unknown.

```julia
function Ytail(x, t, dx, u_bound::Function)
    while true
        xold = x
        t += dx^2 * log(rand()) / 2
        x += rand(Bool) ? dx : -dx
        return t - eps() <= 0 ? u_bound(xold, 0) :
               x - eps() <= 0 ? u_bound(0, t) :
               x + eps() >= 1 ? u_bound(1, t) : continue
    end
end
```

$Y(j\Delta x, t, \Delta x)$ is just the boundary condition for the first passage point from the time space domain for a Poisson time process and a random walk in space. In this implementation as $\Delta x \to 0$ the cost of calculating the first passage time grows $\approx O(\Delta x^{-2})$ and the process converges to Brownian motion. Note that the estimator needs only 1 function call from the boundary conditions.

**Example 4.0.4** (4.0.2 + source + coefficients)

Example 4.0.2 can be extended with a source term and 0-order coefficients:

$$u_t = u_{xx} + a(x,t)u + f(x,t). \tag{93}$$

Again by using (52) with $\sigma = \frac{2}{\Delta x^2} + a_0$ looking at an interior point we obtain:

$$u = \int_0^{e^{-t\sigma}} u_0 d\tau + \int_{e^{-t\sigma}}^1 \sigma^{-1} \left( \frac{u_+ + u_-}{\Delta x^2} + (a(x,s) + a_0)u + f \right) d\tau. \tag{94}$$

To remove branching recursion we sample one of the $u$'s, we base the sampling probability based on the magnitude of coefficients, let $a_m$ be approximately the magnitude of $(a(x,s) + a_0)$. We chose to sample the $(a(x,s) + a_0)u$ term and $f$ with probability $p_{\text{source}} = \frac{a_m}{a_m + \frac{2}{\Delta x^2}}$. As $\Delta x \to 0$ the $\frac{u_+ + u_-}{\Delta x^2}$ term is the main contribution to the second integral therefore being sampled almost always. In total for 1 fully recursed sample, the $f(x,s)$ and the $(a(x,s) + a_0)u$ term only is sampled a few times and this does not scale with $\Delta x$. We sampled $f$ together with $(a(x,s) + a_0)u$ for simplicity.

**Julia Code 4.0.5** (implementation of estimator of Example 4.0.4)

Because of the high chance of sampling the $\frac{u_+ + u_-}{\Delta x^2}$ it is efficient to sample the inter-arrival until not sampling it which is geometrically distributed.

```julia
using Distributions
function Yvar(x, t, dx, a0, am, u_bound::Function, f::Function, a::Function)
    siginv = 1 / (2 / dx^2 + a0)
    (p_source = am / (am + 2 / dx^2); p_avg = 1 - p_source)
    (geom = Geometric(p_source); expon = Exponential(siginv))
    (sol = 0; w = 1)
    sterm_counter = rand(geom)
    while true
        t -= rand(expon)
        t <= eps() && return sol + w * u_bound(x, 0)
        if sterm_counter != 0
            x += rand(Bool) ? dx : -dx
            w *= 2 * siginv / (dx^2 * p_avg)
```

```
14              sterm_counter -= 1
15              return x - eps() <= 0 ? w * u_bound(0, t) + sol :
16                      x + eps() >= 1 ? w * u_bound(1, t) + sol : continue
17          else # this is only done O(am tavg) times in an estimator
18              sterm_counter = rand(geom)
19              sol += w * f(x, t) * siginv / p_source
20              w *= (a(x, t) + a0) * siginv / p_source
21          end
22      end
23  end
```

Computing the estimator requires knowing more then the first passage point unlike
Example 4.0.2 but this is limited to a few points of the time space process and this
amount does not scale with $\Delta x$.

**Technique 4.0.6** (presampling the process of 4.0.5)

As $\Delta x \to 0$ the bottleneck is sampling the process that moves through space and
time. This process is independent of the boundary conditions, $a(x,t)$ and $f(x,t)$.
So it is possible to precompute samples of the process and reuse them for different
boundary conditions, $a(x,t)$ and $f(x,t)$. Reusing paths makes the solutions corre-
lated. To compute the estimator we need to know where we sampled the source
term to update *sol* and $w$, how many steps we took in between to update $w$ for
the multiplications in not source points (line 13 of 4.0.5) and where termination
happens.

```
1   using Distributions
2   function genPath(x, t, dx, a0, am)
3       (siginv = 1 / (2 / dx^2 + a0); p_source = am / (am + 2 / dx^2))
4       (geom = Geometric(p_source); expon = Exponential(siginv))
5       sterm_counter = rand(geom)
6       spoints = [(x, t, sterm_counter)] #maybe other data struct
7       while true
8           t -= rand(expon)
9           t <= eps() && return (spoints, (x, t, sterm_counter))
10          if sterm_counter != 0
11              x += rand(Bool) ? dx : -dx
12              sterm_counter -= 1
13              return x - eps() <= 0 ? (spoints, (x, t, sterm_counter)) :
14                      x + eps() >= 1 ? (spoints, (x, t, sterm_counter)) :
15                          continue
16          else # this is only done O(am tavg) times in an estimator
17              sterm_counter = rand(geom)
18              push!(spoints, (x, t, sterm_counter))
19          end
20      end
21  end
```

Paths for smaller $\tilde{a}_m$'s can be obtained by thinning the Bernoulli process, by rejecting
source points with probability $\frac{\tilde{a}_m}{a_m}$. As $\Delta x \to 0$ *genPath* converges, the dependence
on $a_0$ also disappears. For the heat equation *genPath* presampled paths only depend
on geometry and the starting point.

**Julia Code 4.0.7** (4.0.5 using precomputed paths)

We implement 4.0.5 with a precomputed path from 4.0.6 using tail recursion. Be-

*Figure 7: 20 samples of genPath($x = 0.5, t = 0.15, dx = \{0.1, 0.001\}, a_0 = 0, a_m = 30$). The source term is sampled at the intermediate points. Note that the amount of intermediate points does not differ much between $\Delta x = 0.1$ and $\Delta x = 0.001$. Points with $t < 0$ do not cause any bias, the method is unbiased for the semi-discretized heat equation.*

cause we know the path beforehand a forward implementation is straightforward. Note that $(x, t)$ is never explicitly used only for figuring out how we exited the domain. The whole calculation is linear in $a, f$ evaluated in the source points and $u_{\text{bound}}$ evaluated in the exit points so there are various ways to optimize it for example using unbiased estimates instead of exact evaluations.

```
1   function YvarPath(path, dx, a0, am, u_bound, f, a)
2       spoints, exit = path
3       (siginv = 1 / (2 / dx^2 + a0); p_source = am / (am + 2 / dx^2))
4       w_multiplier = 2 * siginv / (dx^2 * (1 - p_source))
5       (sol = 0; w = w_multiplier^spoints[1][3])
6       for (x, t, sterm_counter) in spoints[2:end]
7           sol += w * f(x, t) * siginv / p_source
8           w *= (a(x, t) + a0) * siginv / p_source
9           w *= w_multiplier^sterm_counter
10      end
11      (x, t, sterm_counter) = exit # small w can produce NaNs
12      w /= sterm_counter > 0 && abs(w) > eps() ? w_multiplier^sterm_counter : 1
13      t >= eps() ? sol + w * u_bound(x, t) : sol + w * u_bound(x, 0)
14  end
```

## 4.1 First Passage Sampling

In this subsection we discuss recursive first passage sampling to efficiently first passage sample complicated geometries.

**Definition 4.1.1** (first passage time)

Define the first passage time for a process $X_t$ for a set of valid states $V$ as

$$\text{FPt}(X_t, V) = \inf\{t > 0 | (X_t, t) \notin V\}. \tag{95}$$

Note that the first passage time is a RV itself.

25

**Definition 4.1.2** (first passage)

Define the first passage for a process $X_t$ for a set of valid states $V$ as

$$\mathrm{FP}(X_t, V) = (X_\tau, \tau), \tau = \mathrm{FPt}(X_t, V). \tag{96}$$

**Lemma 4.1.3**

When a process has more valid states the first passage time gets larger i.e.

$$V_1 \subset V_2 \Rightarrow \mathrm{FPt}(X_t(\omega), V_1) \leq \mathrm{FPt}(X_t(\omega), V_2). \tag{97}$$

The $\omega$ is to indicate the same realization of $X_t$.

**Theorem 4.1.4** (Green's functions and first passage distribution)

The density of first passages of Code 4.0.3 is the corresponding Dirichlet boundary Green's function for the semi-discretized heat equation.

*Proof.* Follows directly from the definition. Let $P(x, t | x_0, t_0)$ denote the boundary Green's function for the semi-discretized heat equation and $X_t$ the corresponding space time process.

$$P(x, t | x_0, t_0) = E[Y(x_0, t_0)] \tag{98}$$
$$= E[\tilde{u}(X_\tau, \tau) | X_{t_0} = x_0] \tag{99}$$
$$= E[\delta_{((X_\tau, \tau) = (x, t))} | X_{t_0} = x_0] \tag{100}$$
$$= P((X_\tau, \tau) = (x, t) | X_{t_0} = x_0). \tag{101}$$

$\square$

Another way to obtain an unbiased estimator for the semi-discretized heat equation is using unbiased estimates of the boundary Green's boundary function. Because of:

$$\tilde{u}(x_0, t_0) = \int_{\partial\Omega} \tilde{u}(x, t) P(x, t | x_0, t_0) d(x, t) \tag{102}$$

$$= \int_{\partial\Omega} \tilde{u}(x, t) dP((X_\tau, \tau) = (x, t) | X_{t_0} = x_0). \tag{103}$$

**Related Work 4.1.5** (recursively estimating Green's functions)

In [Qi+22] unbiased estimators for the boundary Green's function are constructed by recursively sampling known Green's functions.

Similar to recursively sampling simpler Green's function we recursively sample first passages.

**Technique 4.1.6** (recursive first passage sampling)

Recursive first passage sampling involves sampling an initial, simpler first passage, the base that includes fewer valid states. Using this sampled first passage as a starting point, we then perform the same sampling process until the sampled first passage is almost invalid.

**Example 4.1.7** (Euler first passage sampling)

In this example, we approximately sample the first passage of Brownian motion for a parabolic barrier by simulating Brownian motion with the Euler scheme. We plotted this in Figure 8. The accuracy of the sampled first passage are $O(\Delta t)$ and the cost to sample is $O(\Delta t^{-1})$.

```
1  in_triangle(time, pos) = (1 - time > abs(pos))
2  function sample_euler_triangle(dt=0.001)
3      (time = 0; pos = 0)
4      while in_triangle(time, pos)
5          time += dt
6          pos += sqrt(dt) * randn()
7      end
8      return (time, pos)
9  end
```
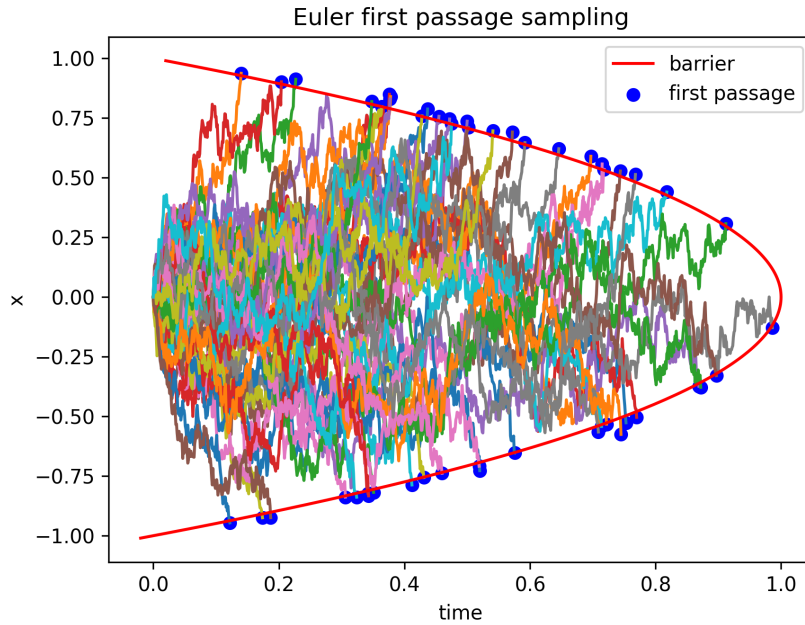


Figure 8: 50 realizations of Euler first passage sampling with step size 0.001.

**Example 4.1.8** (recursive first passage sampling)

In this example, we sample the first passage of Brownian motion from a parabolic barrier with recursive first passage sampling. For the simpler first passage sampler, we exploit the self-affinity of Brownian motion ($\frac{W_{ct}}{\sqrt{c}} \sim W_t$) by scaling and translating samples from a triangular barrier so its valid states are contained in the parabola generated by the Euler scheme. The precomputed samples are created by 4.1.7. Assuming that the error of the base sampler is insignificant. The accuracy of the sampled first passage are $O(\varepsilon)$ and the cost to sample is $O(\ln(\varepsilon^{-1}))$. Note that resampling with replacement produces dependent samples and introduces additional error. For MC applications dependence is not an issue and when the additional error is significant resampling without replacement can be used.

```
1  function triangle_scale_in_para(time, pos)
2      xx = sqrt(1 - time) - abs(pos)
3      tt = abs(1 - abs(pos)^2 - time)
4      return sqrt(tt) < xx ? sqrt(tt) : xx
5  end
6  function sample_recursive_para(accuracy=0.01, scale_mul=0.9)
7      (time = 0.0; pos = 0.0)
8      scale = triangle_scale_in_para(time, pos)
9      while scale > accuracy
10         scale *= scale_mul
11         dtime, dpos = triangle_sample[rand(1:length(triangle_sample))]
12         dtime, dpos = scale^2 * dtime, scale * dpos
13         (time += dtime; pos += dpos)
14         scale = triangle_scale_in_para(time, pos)
15     end
16     return (time, pos)
17 end
```

When the domain is convex, the maximum scaling of the triangular barrier that fits
is easier to obtain. To dampen barrier overstepping of 4.1.7 we use a smaller scaling
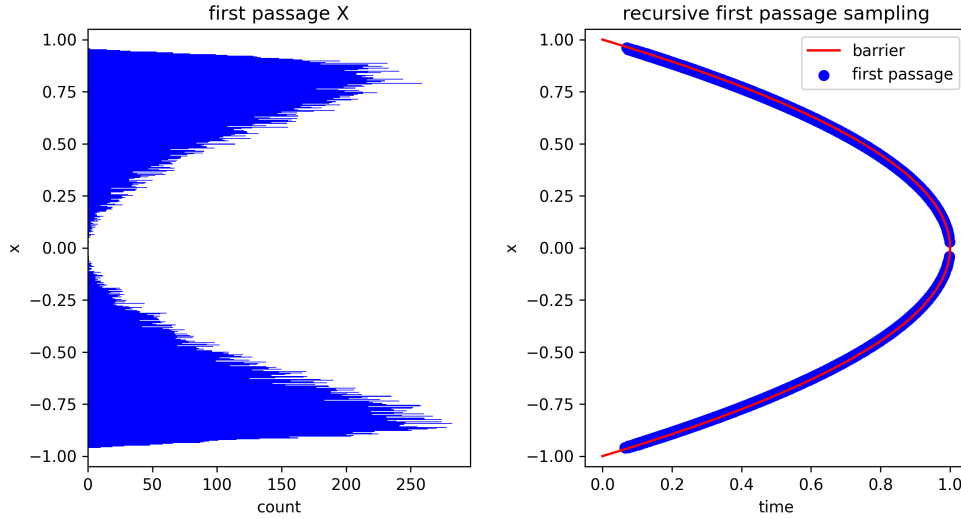than the maximum.



*Figure 9: 50000 of realizations of recursive first passage sampling produced by
(4.1.8). The precomputed sample of 5000 first passages of a triangular barrier uses
the Euler scheme with step size 0.001.*

**Related Work 4.1.9** (recursive first passage sampling)

The original walk on spheres is a recursive first passage algorithm. Recursive first
passage sampling for Brownian motion is discussed in [HT16] and by transformation
also first passage problems for the Ornstein-Uhlenbeck process. A more efficient
alternative to resampling from an Euler scheme is to use tabulated inverse cumulative
probability functions, as demonstrated in [HGM01].

**Technique 4.1.10** (Brownian motion path stitching)

Sampling approximate Brownian motion paths also can be framed as a first passage

problem. Instead of sampling first passage points we sample first passage paths. This requires a simple first passage paths sampler and the paths sampled this way are "stitched" together.

**Example 4.1.11** (path stitching parabola)

Example 4.1.8 but resampling full Euler scheme generated paths. The advantages over generating paths directly from the Euler scheme are that computations for all small steps from the base sampler are avoided and the full path can be represented by its subpaths and their scalings requiring less memory than a fully stored path. The only downsides are correlation between paths and inhomogeneous time steps.
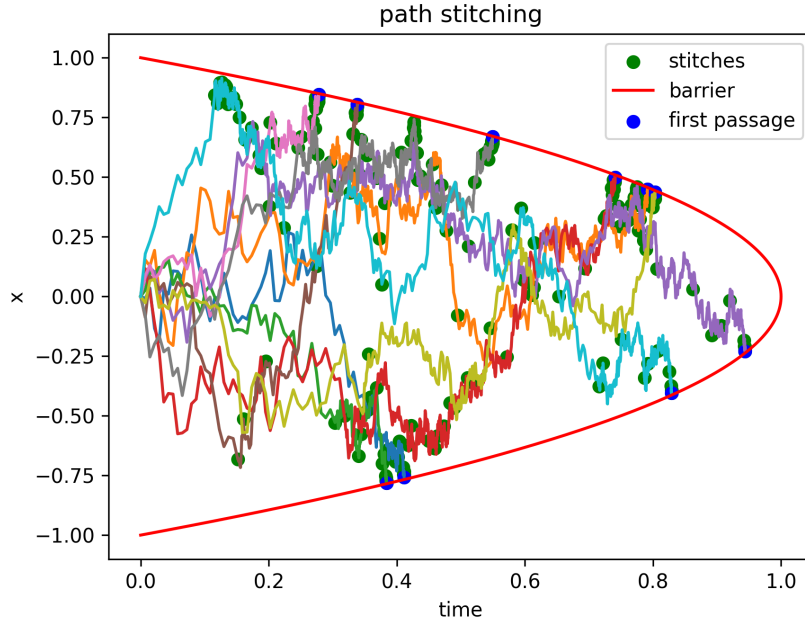


*Figure 10:* 10 *paths build with path stitching build out of resampled Euler scheme generated paths with step size* 0.01*.*

**Related Work 4.1.12** (path stitching)

Path stitching appears frequently in rendering and also in [Das+15] and [JL12].

# 5 Limitations and Future Work

We believe that understanding and optimizing unbiased and deterministic linear ODE solvers is the key to developing better randomized ODE/PDE solvers. Randomized ODE/PDE solvers are useful for cases with little structure where the advantage of IBC is significant or where the linear trade-off between cost and variance is close to optimal.

Besides the time process, other areas of development could include the cheaper construction of control variates, different types of control variates, adaptive schemes, freezing less important terms in the inner recursion or Russian rouletting them into

reasonable approximations, error estimates based on variance in the inner recursion, etc.

To handle stiff problems we weigh towards exponential integrators type of methods. The biggest obstacle to implementing them similar to diagonal RRMC is getting unbiased estimates of $e^{A(t-s)}y(s)$ type of expressions. In the case of a big matrix $A$ unbiased sparsifaction will probably be useful see [SM09]. Initially, we came up with Example 2.5.2, but some time later, we found the paper from NVIDIA [Ket+21] that optimizes that example. Closely related to this is directly estimating the Magnus expansion, where expressions like $e^{\int_0^{\Delta t} A(s)ds}y(0)$ are needed. In this case, [Ket+21] doesn't utilize the smoothness of $A(s)$, which is necessary for optimal IBC.

One of the elements lacking in our findings is rigor. We believe that RMC is an informal approach to an unbiased estimate of the Von Neumann series to the corresponding integral equation. [ES21] presents Theorems 1 and 2 to show that their estimates have finite variance and provide an expression for it. Before becoming aware of [ES21], we previously derived a similar expression (with errors) by employing the law of total variance, similar to (16) in [Rat+22].

We believe that proving the optimality of IBC in Example 3.2.3 is feasible but tedious. [Dau11] presented a proof for optimal IBC for their algorithm. The proof we have in mind is using a lower bound on IBC from integration and proving it is attained.

Optimal IBC isn't everything. Being optimal in IBC doesn't necessarily mean it's optimized. An algorithm that uses 1000 times more function calls may still have the same IBC. Additionally, the computational goal might not align well with the IBC framework. We admire [Bec+22], which employs deep learning to extend beyond the IBC framework. The emphasis here is on performing multiple rapid solves (inferences) while allowing for an expensive precomputation (training). IBC also doesn't take into account the parallel nature of computations. Given infinite parallel resources, it would be reasonable to cease reducing variance by decreasing the step size and instead opt for splitting the final estimator. The only necessary communication in splitting is averaging the final estimator. We hypothesize that for RMC, the wall time at risk increases logarithmically with the size of the splitting.

## Abstract

Deze scriptie onderzoekt recursieve Monte Carlo voor het oplossen van lineaire gewone differentiaalvergelijkingen met het oog op partiële differentiaalvergelijkingen. De voorgestelde algoritmes maken gebruik van de geschikte combinatie van Monte Carlo technieken. Deze Monte Carlo technieken worden geïntroduceerd met voorbeelden en code.

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