# Quantum Gravity Simulations with MIXMAX Generator 

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## Introduction to Causal Dynamical Triangulations

## What is Causal Dynamical Triangulations?

Causal Dynamical Triangulations (CDT) is a background independent approach to quantum gravity. It provides a lattice regularization of the formal gravitational path integral via a sum over causal triangulations.

$$
\int \mathrm{D}[g] e^{i S^{E H}[g]} \rightarrow \sum_{\mathcal{T}} e^{-S^{R}[\mathcal{T}]}
$$



## MIXMAX and CDT

For the last three years we have been using the standard version of the MIXMAX pseudorandom number generator with $N=256$, $s=-1, m=0$ (and discarding the first component) in our CDT code.
It shows many advantages:

- Great quality (passes Big Crush from TestU01)
- Very fast (although it's a small part, $<0.5 \%$ CPU time)
- Convenient vector size $(255=3 \times 85)$
- Convenient output size (61-bits, double)


## It just works!

We encountered no issues which could be related to the RNG.

## Regularization via triangulation

- A four-dimensional simplicial manifold is obtained by gluing pairs of four-simplices along their three-faces (tetrahedra).
- The metric is flat inside each 4-simplex.
- Curvature is localized at triangles.
- Global proper-time foliation. Spatial states are three-dimensional geometries. Discretized states are build from equilateral tetrahedra.
- Length of time links $a_{t}$ and space links $a_{s}$ is constant.

Fundamental building blocks


4D


## Regge action

The Einstein-Hilbert action has a natural realization on piecewise linear geometries called Regge action

$$
S^{E}[g]=-\frac{1}{G} \int \mathrm{~d} t \int \mathrm{~d}^{D} \times \sqrt{g}(R-2 \Lambda)
$$



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$$
S^{R}[\mathcal{T}]=-K_{0} N_{0}+K_{4} N_{4}+\Delta\left(N_{14}-6 N_{0}\right)
$$

$N_{0}$ number of vertices
$N_{4}$ number of simplices
$N_{14}$ number of simplices of type $\{1,4\}$
$K_{0} K_{4} \Delta$ bare coupling constants $\left(G, \Lambda, a_{t} / a_{s}\right)$

## Causal Dynamical Triangulations

- The partition function of quantum gravity is defined as a formal integral over all geometries weighted by the Einstein-Hilbert action.

$$
Z=\int \mathrm{D}[g] e^{i S^{E H}[g]}
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- To make sense of the gravitational path integral one uses the standard method of regularization - discretization.
- The path integral is written as a nonperturbative sum over all causal triangulations $\mathcal{T}$ Wick rotation is well defined due to global proper-time foliation.
- Using Monte Carlo techniques we can approximate expectation values of observables.


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Monte Carlo simulations - initial and minimal configuration To start the simulations we need to construct an initial triangulation with the desired topology.

- Minimal triangulation: five tetrahedra.

Toroidal topology $T^{3} \times S^{1}$

- Regular initial triangulation: 1024 four-simplices per slice
- Minimal triangulation: layered and interlaced structure, 15 vertices (not $3^{3}$ ) and 90 tetrahedra.



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Layer $L_{2}, y=2$


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## Monte Carlo simulations - Pachner moves

Random walk over configuration space:

- Ergodicity all possible configurations can be generated by moves
- Fixed topology moves don't change the topology
- Causality moves preserve the foliation
- 4D CDT set of 7 moves

Example of a 2D Monte Carlo move


## Monte Carlo Markov Chain

- We perform a random walk in the phase-space of configurations (space of piecewise linear geometries).
- Each step is one of the 4D CDT moves.
- The weight (acceptance probability) $W(\mathcal{A} \rightarrow \mathcal{B})$ of a move from configuration $\mathcal{A}$ to $\mathcal{B}$ is determined (not uniquely) by the detailed balance condition:

$$
P(\mathcal{A}) W(\mathcal{A} \rightarrow \mathcal{B})=P(\mathcal{B}) W(\mathcal{B} \rightarrow \mathcal{A})
$$

- The Monte Carlo algorithm ensures that we probe the configurations with the probability $P(\mathcal{A})$.
- After sufficiently long time, the configurations are independent.
- All we need, is the probability functional for configurations $P(\mathcal{A})$ up to the normalization (Regge action).


## Dynamical lattice

- The Monte Carlo moves alter the underlying spacetime lattice making it dynamical. The (sub)-simplices appear and disappear modifying the connectivity.
- To effectively deal with such system various techniques are used.

Basic stages of a single move attempt:

- Choose which move type (with adjusted probabilities).
- Choose where to make the move (index).
- Verify move validity (hash table) and calculate its weight (maximal information).
- Check whether to make the move (detailed balance). Reject the move, or accept the move.
- Update the triangulation if move was accepted.


## Internals of CDT code: indices and labels

- Each object (point, link, triangle or simplex) is assigned two integer numbers: a label and an index.
- The label is fixed during the object lifetime allowing for its unique identification. Labels are not ordered continuously (removal).
- Indices are ordered continuously. They are split into few types to quickly find a valid move location. Index of a (sub)-simplex may change for various reasons (hidden randomness).


## Indices



Labels

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- The label is fixed durino the obiect lifetime allowino for its unique idf New version ( $>30 \%$ faster) (removal) rewritten code
- Indices ar simple, separate containers for to quickly may chan
- tabelarization
- avoid int $\leftrightarrow$ f.p. conversions
- better hashing
 sub-simplices
types implex
- common field for coordination number and time-likeness


Labels

## Internals of CDT code: data structures

- Simulations are limited by CPU time consumption, while memory usage is not a bottle-neck.
- To enhance the acceptance rate of the moves and gain fast access to necessary data maximal information philosophy is adopted.
- Various information about points, links, triangles and simplices (but not tetrahedra) are stored:

```
struct triangle
{
int p[3]; /* Vertex labels */
int n; /* Coordination number */
int s; /* Simplex which owns this triangle */
int i; /* Index */
int h; /* Hash code of triangle */
int next; /* Next triangle in hash table list */
}
```


## Internals of CDT code: hash tables

- To perform a fast check if a link or triangle with given vertices exists, a hash table is used.
- Verification of tetrahedron existence is more complicated and demands making a loop around a triangle.
Given labels of vertices $\mathrm{p}[0]<\mathrm{p}[1]<\mathrm{p}[2]$ we calculate the hash code,

$$
h=(p[0] \ll 8) \text { xor }(p[1] \ll 4) \text { xor } p[2]
$$

Triangles with the same hash code are stored in a singly-linked list.


## Numerical setup

- Monte Carlo algorithm probes the space of configurations with the probability $P[\mathcal{T}]=\frac{1}{Z} e^{-S[\mathcal{T}]}$.
- To calculate the expectation value of an observable, the path integral is approximated by a sum over a finite set of Monte Carlo configurations.

$$
\begin{aligned}
\langle\mathcal{O}[g]\rangle & =\frac{1}{Z} \int \mathcal{D}[g] \mathcal{O}[g] e^{-S[g]} \\
& \downarrow \\
\langle\mathcal{O}[\mathcal{T}]\rangle & =\frac{1}{Z} \sum_{\mathcal{T}} \mathcal{O}[\mathcal{T}] e^{-S[\mathcal{T}]} \\
& \downarrow \\
\langle\mathcal{O}[\mathcal{T}]\rangle & \approx \frac{1}{K} \sum_{i=1}^{K} \mathcal{O}\left[\mathcal{T}^{(i)}\right]
\end{aligned}
$$

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## What can we measure?

From the computational point of view, the basic observables are the parameters which appear in the bare action:
$N_{0}$ number of vertices
$N_{41}$ number of simplices of type $\{4,1\}$
$N_{32}$ number of simplices of type $\{3,2\}$
The number of all remaining sub-simplices can be expressed as a linear combination of above parameters.
The simplest observable providing information about spacetime geometry is spatial volume $n_{t}$ defined as a number of tetrahedra building a three-dimensional slice $t=1 \ldots T$.
(Proper definition of e.g. curvature is non-trivial.)

## Phase structure



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Second order phase transition line $K_{0}$

## De Sitter phase - background geometry

- In phase C the time translation symmetry is spontaneously broken and the three-volume profile $n_{t}$ is bell-shaped.
- The average volume $\left\langle n_{t}\right\rangle$ is with high accuracy given by formula

$$
\left\langle n_{t}\right\rangle=H \cos ^{3}\left(\frac{t}{W}\right)
$$

a classical vacuum solution.


## De Sitter phase - properties

- Phase C, the so called de Sitter phase, is physically most interesting.
- A four-dimensional background geometry emerges dynamically. The background geometry corresponds to Euclidean de Sitter space ( $S^{4}$ ), a classical vacuum solution.
- It is also possible to study quantum fluctuations around it.



## Autocorrelation: MIXMAX vs Ran3

The MC moves introduce only small local changes and the configurations may be correlated for a long time.

$$
\text { د } 0.20 \text { ( } 0 \text { ( }
$$

## Autocorrelation time: MIXMAX vs Ran3

Often in the vicinity of phase transition, a phenomenon of critical slowing down is observed.
It takes longer for the system to thermalize, amplitude of fluctuations increases and the autocorrelation time grows.




## The new bifurcation phase




- The spatial volume profile $\left\langle n_{t}\right\rangle$ is similar as in phase $C$. But the agreement with $\cos ^{3}(t)$ is broken.
- The transfer matrix bifurcates into two branches. At some volume the kinetic term splits into a sum of two shifted Gausses.
- Every second slice singular vertices of very high order appear. New order parameter: $\max _{v} o(v)$.
- Not captured by global properties of the triangulation $\left(N_{0}, N_{32}\right)$.


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## Toroidal topology - phase transitions

For the toroidal topology, the $C_{b i f} / C_{d S}$ transitions reveals a hysteresis and very long thermalization time. Reason?

- Strong finite size effects
- Phase transition order depends on topology?


How to improve the simulations?

## Motivation

- Limited single-core performance (gates, frequency, voltage, power)
- Cheap access to many cores

Naive parallelization is used to multiply the statistic, but ...

- ... a thermalized configuration is needed.
- Thermalization stage is a bottleneck
- Critical slowing down near phase transition
- Need to speed-up a single Monte Carlo Markov chain $\Rightarrow$ parallelization
- Large minimal triangulation $\Rightarrow$ strong finite size effects $\Rightarrow$ large triangulations
- Simple parallelization does not work ...


## Simple parallelization

A simple parallel algorithm, where many threads modify the same triangulation, has a very large overhead

- Detect collisions
- Thread synchronization: withdraw a move
- Move weight depends on global parameters - threads are coupled



## Parallel rejection

(by Jack Laiho, Phys. Rev. D 96 (2017) 064015)

- In some regions, the acceptance rate of MC moves is very low (critical slowing down)
- In such case a parallel rejection algorithm can be applied
- Multiple threads execute a series of attempted moves
- When one of them succeeds, rest of the threads are stopped and the triangulation is updated
- No collision detection is necessary
- To guarantee compatibility with the scalar version, youngest move has to be chosen (wait)
- Each attempted move is identified by three random numbers:
- which move
- where to make the move
- whether to make the move (detailed balance)
- A block of random numbers is precomputed and split between threads. Synchronization with the RNG (separate thread).


## Parallel rejection - diagram

|  | Thread |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  | 1 | 2 | 3 | 4 |  |
|  | 1 | 2 | 3 |  | Enumerate |
|  | 5 | 6 | 7 | 8 | set of a few |
|  | 9 |  | 11 | 12 | random number |
|  | 13 | 10 | 15 | 16 |  |
| . | 17 | 14 | - | - |  |
| $\square$ |  |  |  |  | Attempted |
| $\stackrel{\square}{0}$ |  |  |  |  | MC move |
| $\bigcirc$ |  |  |  |  |  |
|  | 41 |  |  |  |  |
|  |  | 38 |  |  |  |
|  |  |  |  |  | Accepted move |
|  |  |  |  |  |  |
|  |  |  | 59 |  |  |
|  |  |  |  |  |  |
|  |  |  |  |  |  |
|  |  |  |  |  |  |

## Parallel rejection - issues

- For three threads we gained $100 \%$ speed up for very small acceptance rate $\frac{1}{400}$
- Different move types significantly vary in acceptance rate
- Fixed number of random numbers (3) consumed per attempted move
- Replicability, compatibility with scalar version
- Thread synchronization is expensive
- Low level libraries, single barrier from pthreads
- Atomic operations (counter), busy waiting


## Replica exchange

- In this method we run a number of processes (replicas) with slightly different values of coupling constants.
- Periodically, the configurations are exchanged between processes with a certain probability (detailed balance).
A CDT configuration is a heavy object (triangulation + additional data). Instead, the sets of a few coupling constants (denoted as $\beta$ ) are exchanged.



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## Replica exchange

- Exchange probability fulfills the detailed balance condition,

$$
\begin{aligned}
P(A) & =P_{1}\left(\mathcal{T}_{1}\right) P_{2}\left(\mathcal{T}_{2}\right), \quad P(B)=P_{2}\left(\mathcal{T}_{1}\right) P_{1}\left(\mathcal{T}_{2}\right), \\
W(A \rightarrow B) & =\min \left\{1, \frac{P_{2}\left(\mathcal{T}_{1}\right) P_{1}\left(\mathcal{T}_{2}\right)}{P_{1}\left(\mathcal{T}_{1}\right) P_{2}\left(\mathcal{T}_{2}\right)}\right\}
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- Implementation: Managing process and Linux message queues to notify working processes when and with whom exchange



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## Autocorrelation time

- History of one configuration (Markov chain) contributes to many simulations and reduces the autocorrelation time.
- Exchange rate has to be higher than the autocorrelation time
- For 31 replicas a significant decrease of autocorrelation time is observed. A point near A/C transition (toroidal topology):

Auto-correlation of $\mathrm{N}_{32}$


## Replica exchange - hysteresis

Monte Carlo history of the relevant order parameter $\operatorname{Max} o(p)$ for single value of coupling constants. Visible jump and exchanges.
$\Delta=0.370$


## Replica exchange - hysteresis

Average value of the order parameter as a function of coupling constant. Comparison: replica exchange vs single threaded.


Thank You!

