

Generalized Stochastic Simulation Algorithm for Artificial Chemistry

Gillespie in artificial chemistry

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Bacterial reaction graphs

```

>AACT3112 -dbfa Chromosome:K08584v2 |Chromosome:130x2551 | gene:130112 | ...
AGGAAAGCGTAGCACTCCGCTTACCCACCGCATCCATCCATTTACDCAKCGATAGCGGTGGG
GGCTTA
>AACT3112 -dbfa Chromosome:K08584v2 |Chromosome:337x2799 | gene:130112 | ...
ATGCGAGTGTGGAGTTGGGATCGGACATGCGATGATGATGAAATGGAGACGCTTCTGCG
GCGGATATTTGGAAAGCAATGCTCAAGGAGAGGCGAGTGGGACAGTCTCTCTGCGGCGCC
GCCAAATCTCCACCAACCTGTGGCGGTGATGGAAACCGTGGACCGGCGAGTCTCT
...
    
```

cDNA fasta file

(a) Step 1: From cDNA fasta file, get gene labels.

The image shows two screenshots from the KEGG database. The left screenshot displays a search result for 'Escherichia coli K-12 MG1615' with a table of gene entries. The right screenshot shows a detailed view of an enzyme entry, 'ENZYME 1.1.1.13', with fields for name, class, reaction, and product.

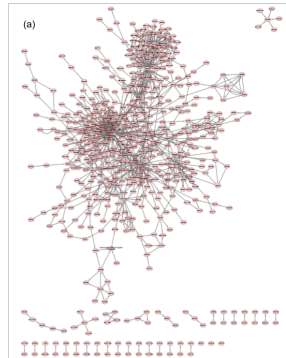
KEGG gene entries

KEGG enzyme entries

(b) Step 2: From KEGG gene entries, get EC (c) Step 3: From KEGG enzyme entries, get substrates and products.



(d) Step 4: Build graph and filter ubiquitous metabolites. Ubiquitous metabolites : H₂O, ATP, ADP, NAD⁺, NADH, NADPH, NADP⁺, CO₂, ammonia, sulfate, thioredoxin, phosphate, PPI, H⁺.



Artificial Chemistry

Motivation

- ▶ (Real) Chemistry is ... difficult
 - Maybe artificial chemistry is easier
 - can model interesting properties (for e.g. space/diffusion)
- ▶ (Real) Life is ... chemistry
 - *Artificial* Life should have ... artificial chemistry (AC)

What is done (usually)

- ▶ chemistry is prescribed : small dimension (small # of reactions)
- ▶ straightforward : chemistry graph is simple
- ▶ (somewhat) unrealistic
 - transition energies are ignored (easier)
 - mass conservation is sloppy
 - $A + B \mapsto C$ and $C \mapsto A$

AC : what is expected

- ① large : should have a huge number of reactions
- ② energy : all reactions are not possible
- ③ mass : reversibility should not be hacked
- ④ open : we don't know all reactions/molecules

Hutton Artificial Chemistry (1)

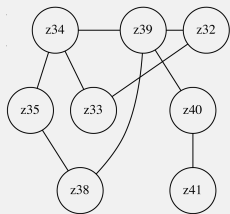
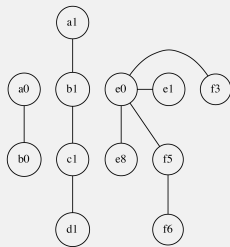
Hutton J.

- ▶ several papers on his AC scheme
 - ▶ only one follow up
 - ▶ straightforward : chemistry is simple
 - ▶ very complex : chemistry reactions network can be extremely complex
 - ▶ however (somewhat) unrealistic
-
- ▶ Hutton, Tim J. Evolvable self-replicating molecules in an artificial chemistry. *Artificial life* 8.4 (2002) : 341-356.
 - ▶ Hutton, Tim J. Evolvable self-reproducing cells in a two-dimensional artificial chemistry. *Artificial life* 13.1 (2007) : 11-30.
 - ▶ Hutton, Tim J. A functional self-reproducing cell in a two-dimensional artificial chemistry. *ALIFE9*. 2004.
 - ▶ Hutton, Tim J. The organic builder : A public experiment in artificial chemistries and self-replication. *Artificial life* 15.1 (2009) : 21-28.

Hutton Artificial Chemistry (2)

Main properties

- ▶ molecules are graphs
- ▶ nodes are domains
- ▶ edges are chemical bonding
- ▶ domains have a fixed type and a changing state
 - described by a pair ($t|s$)
 - usually a letter and a number $a0$, $b1$...
- ▶ any domain can have any numbers of links
- ▶ the chemistry is composed of fully connected subgraphs



Geometry and Physics

- ▶ HAC is $2D$
- ▶ spatially resolved
- ▶ each domain has an id/position
- ▶ links are *springs* (however weird coding)
- ▶ (vaguely) Brownian in viscous medium

Hutton Artificial Chemistry (3)

Reactions

Reactions are of the form :



- ▶ $(t_i|s_j)$ is a domain (within a molecule)
- ▶ $(\cdot|+)$: is either linked \cdot or no (collision) $+$
- ▶ note that type is unchanged
- ▶ modification are only local (other links unmodified)
- ▶ conformation \cdot is performed instantaneously
- ▶ reaction $+$ is performed instantaneously upon collision
- ▶ when conflict : chose at random

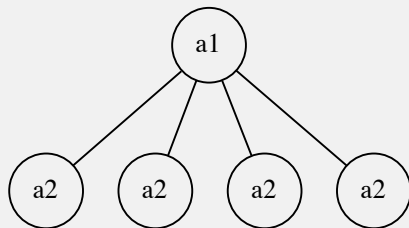
Hutton Artificial Chemistry (4)

Examples

Starting a reactor with several a_0 s and 1 (one) a_1 with



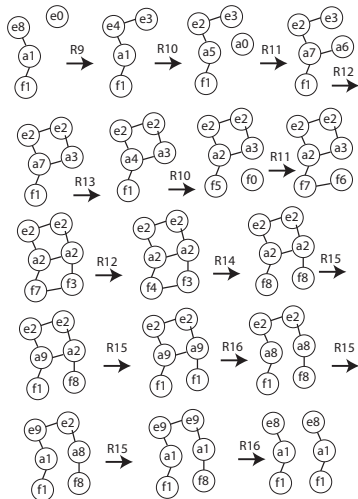
whereas $a_0 + a_1 \mapsto a_2.a_1$ yields



Hutton Artificial Chemistry (5) : replication

Main properties

Using wildcards ...



Pros and Cons

- ▶ Pros :
 - very general
 - 2D and nice
 - mass conservation
- ▶ Cons :
 - 2D and nice
 - absurdly long
 - no reaction rates
 - *intelligently* designed

STAARC : STochastic Atom-based ARtificial Chemistry

- 1 Getting rid of space
- 2 Same data structure (without space)
- 3 reactions with rates
- 4 SSA formalism (Gillespie)
- 5 slightly involved (but not too much)
- 6 github.com/hsoula/staarc

Modifications

- ▶ Reaction have a rate :

$$(t_1|s_1)(\cdot|+)(t_2|s_2) \mapsto (t_1|s_3)(\cdot|+)(t_2|s_4) : \lambda$$

- ▶ For each + reaction :

- We count the number of $n_1 = (t_1|s_1)$ and $n_2 = (t_2|s_2)$
- minus the number of $n_{12} = (t_1|s_1).(t_2|s_2)$ already linked
- the propensity is $a = \lambda(n_1 * n_2 - n_{12})$

- ▶ For each . reaction :

- we count the number of $n_{12} = (t_1|s_1).(t_2|s_2)$ already linked
- the propensity is $a = \lambda n_{12}$

- ▶ then classical Gillespie algorithm

- ▶ once a reaction is selected we apply the modification to a given pair (selected at random uniformly)

First step

compute all propensities a_i Compute the combined rates of all reactions

$$a_0 = \sum_{i=1}^R a_i$$

To compute the time of the next reaction, draw a random number

$$\tau = -\log(rn)/a_0$$

This answers the when

What reaction

To compute the what, we choose randomly the equation weighted by their relative weight i.e

$$\Pr(\text{next reaction is } i) = \frac{a_i}{a_0}$$

in practice, shoot a random number uniform rn in $[0, 1]$ and find r such as :

$$\sum_{i=1}^{r-1} a_i \leq a_0 rn < \sum_{i=1}^r a_i$$

STAARC : STochastic Atom-based ARtificial Chemistry

Properties

- 1 now all reactions have rates : real chemistry
- 2 simulated 3D well mixed medium
- 3 almost all HAC properties conserved
- 4 we can simulate diffusion (reaction rate for collision)
- 5 way faster computation

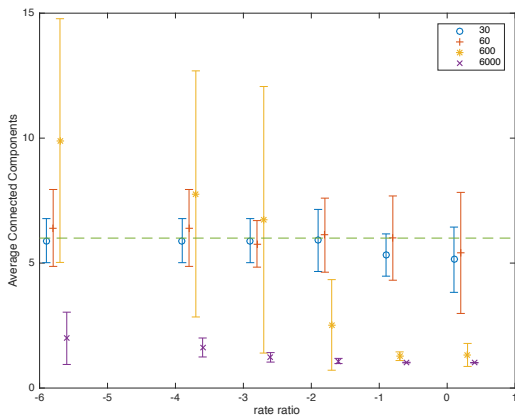
This turn a local resolution to a global one

- ▶ well mixed medium
- ▶ with infinite crowding

not intuitive ...

Replication : molecules size

Simple experiment : the replication according to the initial number of particles ($n \in \{30, 60, 600, 6000\}$) and λ ratio (between collision reaction + and conformation reaction .)



Properties

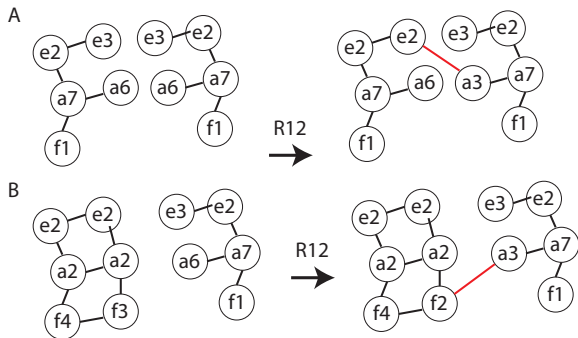
- ▶ Not very resilient - original sequence quickly lost
- ▶ Scales very badly

Not that intelligently designed :)

Race conditions

- ▶ Race condition – diffusion vs conformation
- ▶ Race condition – concurrent replications

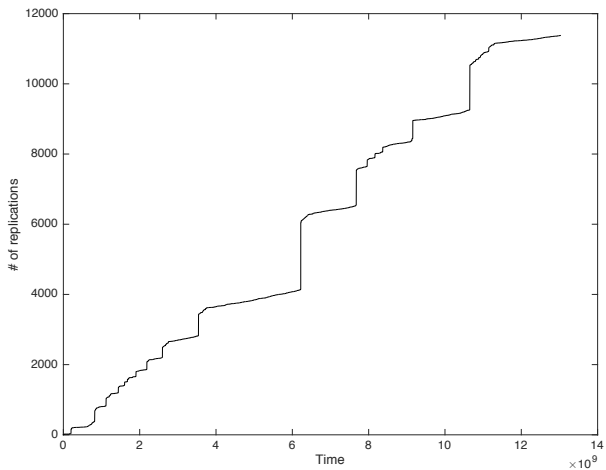
STAARC : replication



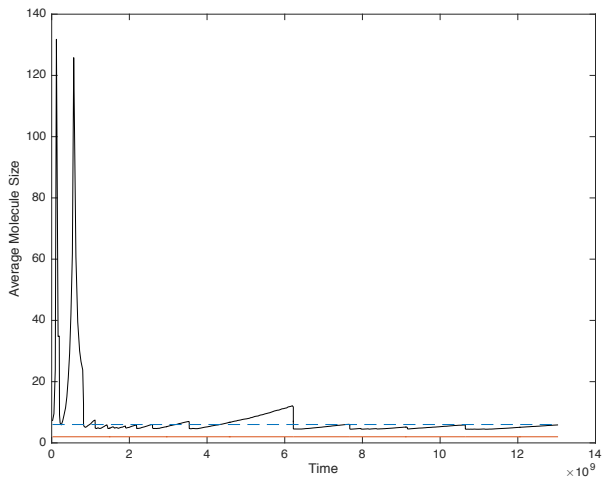
Long replication experiment

- ▶ Starting with size 6 replication seed
- ▶ Production and degradation
 - $(x|0) \rightarrow \emptyset$
 - $\emptyset \rightarrow (x|0)$
- ▶ Let the simulation for 450,000 reactions
- ▶ Compute average size and number of divisions

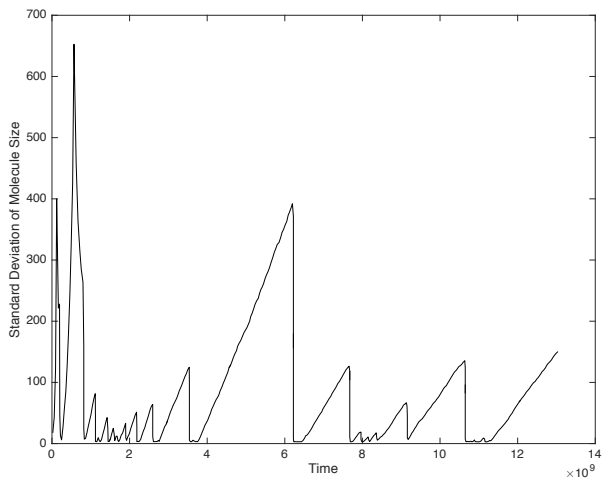
Long replication : replication event



Long replication : average size



Long replication : standard deviation



Long replication experiment

- ▶ Replication is extremely stable
- ▶ Division occurs in bumps
- ▶ The limit size is ... 6
- ▶ Big compounds created transiently

Random Chemical World

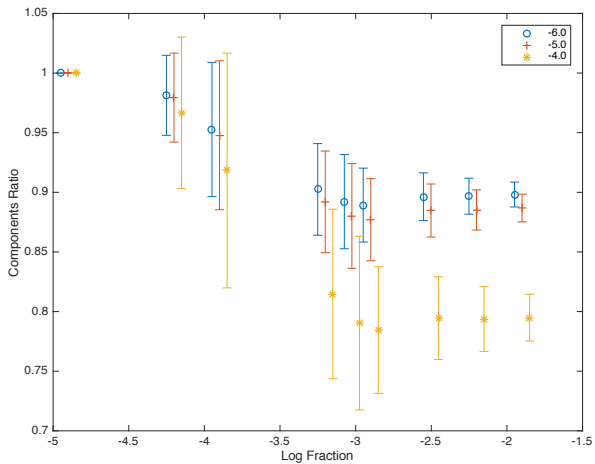
Random generation of reaction

- ▶ available types : $\{a, b, c\}$
- ▶ maximum number of state is 5.
- ▶ compute all the possible reactions
 - with no production nor degradation.
 - keeping only a fraction $p \in [0, 1]$ randomly
- ▶ starting $N = 10,000$ particles $(t|s)$ with $t \in \{a, b, c\}$ and $0 \leq s \leq 4$

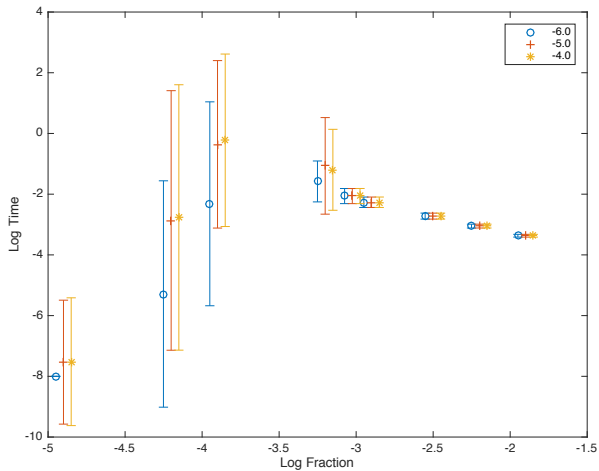
Experiment

- ▶ Maximum of 2,000 reactions.
- ▶ Compute the time needed
- ▶ The ratio of number of molecules

Random Chemical World



Random Chemical World



Phase transition

- ▶ Huge variability in the middle
- ▶ in molecules created
- ▶ in the time to the end of reactions
- ▶ this middle is for a low number of reactions

Drawbacks

- ▶ Results not conclusive
- ▶ Better random selection of reactions
- ▶ Common elements

STAARC

This repository Search

Pull requests Issues Gist

hsoula / staarc

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STAARC: STochastic Atom-based Artificial Chemistry — Edit

7 commits 1 branch 0 releases 1 contributor

Branch: master New pull request

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hsoula update: readme updated quote Latest commit 4404cae on Feb 26

ifemix	add: assembly info / packages	4 months ago
reactor	add: assembly info / packages	4 months ago
replicator	add: assembly info / packages	4 months ago
test	test/	4 months ago
.gitignore	add: ignore file	4 months ago
LICENSE	Initial commit	5 months ago
README.md	update: readme updated quote	4 months ago
staarc.sln	test/	4 months ago
staarc.userprefs	test/	4 months ago

README.md

STAARC: STochastic Atom-based Artificial

github.com/hsoula/staarc

Code

- ▶ Open-Endedness procedural enzyme
- ▶ Tri-molecular reactions
- ▶ Graph-based random chemistry
- ▶ 'Metabolic network'-like graph

Projects

- ▶ Explore Random Chemistry
- ▶ Use it in an evolutionnary set-up
- ▶ Artificial 'bacterial world'

thanks

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Soula HA, Generalized Stochastic simulation algorithm for Artificial Chemistry Proceedings of ALIFE XV, 2016 Cancun, Mexico

