

Stochastic simulation of reaction networks

Week 5

Basic Outline: 3 functions

```
gilldriver
```

```
[t,dimer]=mygill(tfin,rtab,x,c)
```

```
h=update(x)
```

[within driver]

Gilldriver

What does the driver do?

Gilldriver

What does the driver do?

- Set parameters (these are given in notes)
- Set `rtab` (hard-code)
- Run Gillespie (`mygill`) `nr` times for first 6 reactions
 - Plot mean and STD
- Run Gillespie (`mygill`) `nr` times for all 8 reactions
 - Plot mean and STD

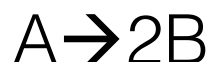
[within driver]

Reactant indices
always in least to
greatest order for
those in *that reaction*

Building rtab

- General idea: alternate between index and #
{[reactant index #gain/lose reactant index #gain/lose]
[reactant index #gain/lose for however many reactants]
... [indices and gain/loss for last reaction]}

For example, take reactions:

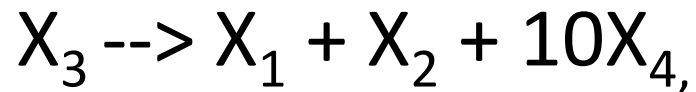


A has index 1;
B has index 2;
C has index 3

rtab={ [1 -1 2 -1 3 1] (lose an A, lose a B, gain a C)
[1 -1 2 2] (lose an A, gain 2 Bs)

Practice

Regarding the rtab convention established in the notes, if I wish to encode the reaction



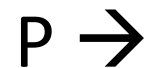
then the associated row of rtab should be:

- A. [1 2 3 4 1 1 -1 10]
- B. [1 1 2 1 3 -1 4 10]
- C. [1 -1 2 -2 3 1 4 -10]

Your turn

Work with the person next to you to build **rtab** by hand for the reactions for this project.

<u>Indices</u>	
R:	1
Pr:	2
RPr:	3
P:	4
D:	5
I:	6



Remember at first we only want to run Gillespie on these 6 reactions. Then run again for the *whole* ruletab!

How can we represent the first 6 rows of ruletab for our first call to mygill?

After setting values and building ruletab, run mygill

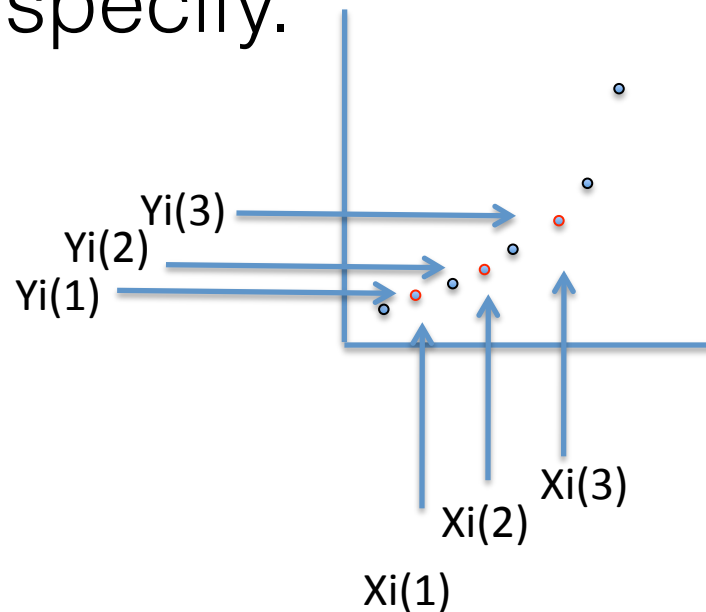
- Run mygill nr times (value specified earlier in code)
- `[t,dimer]=mygill(tfin,rtab,x,c)`
- What is x? What is c? Where do we find those?
- Interpolate. But before addressing this, a word on *time*:
 - we want to plot dimer vs. t for all times from 0 to tfin (100), at an increment of tinc (.1).
 - Build tvec as specified in notes
 - Each of the *nr* times you run mygill, interpolate *t* and *x* with *tvec* (see next slide).

Hint: find STD
using std(x) in
MATLAB

Using interp1

Use: $YI = \text{INTERP1}(X,Y,XI)$

Basic interpolation: you have $y(x)$: vector of y values for given vector of x values. You want to find vector of y_i values for vector of x_i values that you specify.



From notes (modify using
interp):
for $j=1:nr$
[t,x] = mygill(tfin,rtab,x0,c);
X(j,:) = x;
end
avg = mean(X,1);
--find STD

$$h = \text{update}(x)$$

- h is a vector, each of whose elements represents the number of ways the corresponding reaction can occur.

Remember that x contains the quantities of each reactant.

$R + Pr \rightarrow RPr$ $h(1) = \#$ ways this reaction can occur

$RPr \rightarrow R + Pr$ $h(2) = \#$ ways this reaction can occur

$RPr \rightarrow 10P + R + Pr$...

$P + P \rightarrow D$

$D \rightarrow P + P$

$P \rightarrow$

$Pr + D \rightarrow I$

$I \rightarrow Pr + D$ $h(8) = \#$ ways this reaction can occur

Write down $h(1)$ through $h(8)$ as functions of elements in x .

`[t, dimer]=mygill(tfin,rtab,x,k)`

“1. The user is unsure of how to choose maxiter and would prefer to provide the final time, tfin.”

→ Substitute **while** for **for** (how?)

-first call update `h=update(x)`

-Cryptic notes: The use of update is relatively straightforward, e.g., `a = c. * h`; With this a in hand you may now use cumsum and find to find the next reaction index (in just a couple of lines - and with NO if clauses). With this index in hand you may visit the proper element of **rtab**.

-record your dimer count before cycling through while loop again. (Which element of x is dimer?)

Within the while loop

see [lotkaedit.m](#)

- Update counter
- Call update function
- Reaction probability: $a_m = h_m c_m$
- “We shall have occasion to call on:”
 $a_0 = a_1 + a_2 + \dots + a_M$ use MATLAB’s sum function
- Generate random number r_1
- Set $t(i) = t(i-1) + T$ (tau in lotka) (solve 8.1 for T); 8.1: $r_1 = \exp(-a_0 T)$
- Generate random number r_2
- use cumsum and find and subsequently access elements of rtab to update x (quantities of each reactant present)
- Store dimer count, $x(5)$.