# Lab #15

### **Physics 91SI Spring 2012**

**Objective:** In this lab, you will use Cython and C to massively speed up an implementation of John Conway's *The Game of Life* in Python

As usual, log on to corn and clone over the starter repository:

hg clone /afs/ir.stanford.edu/class/physics91SI/src/lab15 lab15

The repository contains several source code files, as well as a file called setup.py . As demonstrated in lecture, this file can be used to automatihcally generate an executable python module from the present Cython and C files. You should call the script like this:

```
python setup.py build_ext --inplace
```

This will take most current version of your C and Cython files and compile them into a Python module that you can then import just like any other Python library.

Remember that you have to use this command every time you modify the C (.c) or Cython (.pyx) files in the folder, otherwise you won't have the newest version as an executable.

Also, remember to hg commit often to save your changes, and <u>submit your code</u> at the end of class.

#### **Part 0:** The Game of Life

In this part of the lab, you should briefly familiarize yourself with the algorithms and the Big-Oh scaling of *The Game of Life*. Don't spend more than 10 minutes on this part.

The Game of Life, or simply Life, provides a toy simulation of population dynamics. The program consists of a "board" (a 2D grid) of cells. In our version, each cell is either alive or dead, signified by the respective point in the grid containing either a 0 or a 1. In the code, the board is a 2D array of long's. The game evolves in discrete time steps. Every step, the following rules are applied:

- Any cell that is *dead* and has exactly 3 neighbors is set to *alive* (reproduction)
- Any cell that is *alive* and has 2 or 3 neighbors survives (stays alive)
- Any cell that is alive and has more than 3 or less than 2 neighbors dies (overpopulation and underpopulation, respectively)

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Take a look at the file life.py. Familiarize yourself with the algorithm and convince yourself that it is doing the right thing. What is the scaling of the algorithm in N?

The life() function starts the actual simulation. Read the comments to figure out what arguments it takes. Be aware that the "c" and "cython" versions of the algorithm are not implemented yet. It will be your job to implement them later.

You can try out the simulation as follows: open up an iPython session and type at the command line:

```
from life import life
```

life(40,0.2,plotting=True,version="python")

You should be able to watch the simulation. Now use the %timeit magic command to time the execution:

```
%timeit life(40,0.2,plotting=False,version="python")
```

Note that we have **set plotting to False**. This is because the plotting over the ssh X-server takes a long time and outweighs the actual execution time. Try it out for different N and make

sure the scaling is what you expected it to be.

## Part 1: Life in Cython

In this part, you'll rewrite the update() function in cython and observe the speedup that comes with this simple change in the code. In order to keep the old python version, you will write this new function in the file life\_cy.pyx. All function calls and definitions have already been written; all you have to do is fill in the body for update\_cy() and getNeighbors(). Make use of all of the tricks that you learned about in lecture to maximize the performance of those two functions. They won't look very different from the old python versions, but you should be able to get at least a factor of 25 speedup with some simple modifications. Use the %timeit command in an iPython session with plotting set to False and compare the performance when version is set to "python" to when it is set to "cython". What speedup do you get? You should also run the simulation with plotting switched on in "cython" mode to observe whether the program is actually working correctly.

### Part 2: Life in C

Now it is time to maximize the performance of the *Life* simulation. Notice that the cython file life\_cy.pyx is set up to call a pure C function, called update\_pure\_c() through a wrapper function, update\_c(). It is now your job to go to the file life.c and implement the update\_pure\_c() and getNeighbor() functions in pure C. Once you have written the two functions, first run the simulation in "c" mode without %timeit with plotting set to True. Does it behave correctly? If so, go ahead and time the execution without plotting for all three versions. By writing straightforward C code, you should be able to get a relative speedup of more than 400 over the pure Python version.