15110 Principles of Computing, Carnegie Mellon University

UNIT 10A Discrete Simulati

Last Time

- How to generate pseudo-random numbers
- Using randomness in interesting applications
- Monte Carlo simulations: running many experiments with random inputs to approximate an answer to a question when an analytical solution is impossible or to hard to obtain

Waking Up

- Understanding systems
- Data Visualization
 - <u>http://www.ted.com/talks/hans rosling shows the best stats you v</u> <u>e ever seen?language=en</u>
 - http://www.ted.com/talks/deb_roy_the_birth_of_a_word?language em_____en___(4:15, 9:30)
 http://www.carbonmap.org/
- Simulations

Understanding Systems

- Data Visualization and Simulations are different
- We try to visualize the results of simulations to make it easy to see/understand the systems

because generally what we try to see / understand or predict is complicated because of the nature of systems.

Systems

- Collection of tracks and railway cars \rightarrow railroad system
- Collection of HW and SW \rightarrow computer System
- Collection of teachers, students \rightarrow school system

Dynamic, Interactive, Complicated

How Can we Study a System?

Experiment with the **actual system**

Experiment with a model of the system

Physical model

May not exist, may be unsafe to work with, expensive to build and modify, some change too slowly over time

- Mathematical model
 - Analytical solution: Equations or systems may be too complex for closed-form or analytical solution
 - Simulation: The imitative representation of the functioning of one system or process by means of the functioning of another, for example a computer program.

Classification due Law and Kelton Simulation, Modeling and Analysis

Computer simulation is a process of making a computer behave like a cow, an airplane, a battlefield, a social system, a terrorist, a HIV virus, a growing tree, a manufacturing plant, a mechanical system, an electric circuit, a stock market, a galaxy, a molecule, or any other thing. This is done with a specific purpose, mainly in order to carry out some "what if" experiments over the computer model instead of the real system.

> Modeling and Simulation, S. Raczynski

Uses of Simulation

- Performance optimization, safety engineering, testing of new technologies.
- Gaining a better understanding of natural and human systems, and making predictions.
- Providing lifelike experiences in training, education, games.

Large Scale Simulations

- Computing power of today enables large scale simulations. For example,
 - Department of Defense: Battle simulations
 - National Center for Atmospheric Research : 1,000 years of climactic changes <u>http://www.youtube.com/watch?v=d8sHvhLvfBo</u>
 - Blue Brain Project at EPFL to reverse engineer the human brain <u>http://www.youtube.com/watch?v=ySgmZOTkQA8</u>

Advantages of Using Simulation

- If we use simulation we can
 - Control sources of variation
 - Choose the scale of time
 - Stop and review
 - Replicate results more easily





Models

- A model is an abstraction of the real system. It represents the system and the rules that govern the behavior of the system.
- The model represents the system itself, whereas the simulation represents the operation of the system over time.

Modeling Concerns

Abstraction: In building models a major issue is to achieve a certain level of accuracy while keeping the complexity manageable

- Identify factors that are the most relevant to the functioning of the system.
- How important is time evolution? (Static vs. dynamic models)
- How important is it to capture continuous behavior over time? (Discrete vs. continuous models)
 - Discrete models: essential variables are enumerable such as integers
 Continuous models: essential variables range over non-enumerable
 - sets such as real numbers
- Do parts of the system exhibit random behavior? (Deterministic vs. stochastic models)

Computational Science

- Computational sciences use computational models (special kind of mathematical models) as the basis of obtaining scientific knowledge.
- Unifies
 - Modeling, algorithms, simulations
 - Computing environment developed to solve science, engineering, medicine, and humanities problems
- Helps explain and predict phenomena using a mechanistic view

Simulation Models are Descriptive

- They tell us how a system works under given conditions but not how to set the conditions to make the system work best
- Simulation does not "optimize" but it helps us in finding an optimal set of parameter settings.

DISCRETE SIMULATION: A Simple Example

Discrete Time and Discrete Events

- Real time vs. model time
 - In simulating the movements of a galaxy one hour simulation may cover billions of years
- In discrete simulation we assume time changes in discrete steps (ticks) and the states of simulated entities change instantaneously

Experimenting with Models

- NetLogo is a programmable modeling environment for simulating natural and social phenomena using discrete simulation.
 - You can learn more about it and download it for free from <u>https://ccl.northwestern.edu/netlogo/</u>.
- It also comes with the Models Library, a large collection of pre-written simulations that can be used and modified.
 - In this lecture we will do an example based on Wilensky, U. (1998). NetLogo Virus model. http://ccl.northwestem.edu/netlogo/models/Virus. Center for Connected Learning and Computer-Based Modeling. Northwestem University, Evansion, IL.

Discrete Simulation of Disease Spread

- We are going to use a dynamic, discrete, stochastic simulation model
 - $\hfill\square$ We want to capture how the disease spreads over time
 - We model time discretely as a sequence of days, and use discrete variables to capture the health state of each person
 There is randomness in how the virus spreads
- Simulate the system execution as a sequence of discrete events that change the state of the system instantaneously at each time step

Example: Flu Virus Simulation

Goal: Develop a simple simulation that shows graphically how disease spreads through a population.

Modeling the Spread of Flu Virus

- Every person is either healthy, infected, contagious or immune. We assume that "infected" means infected but not contagious.
- Each day, a healthy person comes in contact with 4 random people. If any of those random people is contagious, then the healthy person becomes infected.
- It takes one day for the infected person to become contagious.
- After a person has been contagious for 4 days, then the person is non-contagious and cannot spread the virus nor can the person get the virus again due to immunity.



Displaying the Population

adjacent does NOT mean that they are physically close.

200 pixels

7

Graphical Simulation			
Simulation captures the evolution of the health state of the population over time. It evolves in discrete steps: change occurs instantaneously as a new day begins.			
	• • • •		











Неа	lth	Stat	es

0	white	healthy	HEALTHY = 0
1	pink	infected	INFECTED = 1
2	red	contagious (day 1)	DAY1 = 2
3	red	contagious (day 2)	DAY2 = 3
4	red	contagious (day 3)	$\mathbf{DAY3} = 4$
5	red	contagious (day 4)	DAY4 = 5
6	purple	immune (non-contagious)	$\mathbf{IMMUNE} = 6$

The health state of the population will be represented using a 20 by 20 matrix where each entry has one of the values above.

Running the Code

import tkinter
from tkinter import Canvas
from random import randrange
from time import sleep

Constants for health states of an individual HEALTHY = 0 INFECTED = 1 DAY1 = 2 DAY2 = 3 DAY3 = 4 DAY4 = 5 IMMUNE = 6

Checking Health State

#returns True if person (matrix[r][c]) is healthy
def healthy(matrix, r, c):
 return matrix[r][c] == HEALTHY

#returns True if person (matrix[r][c]) is infected def infected(matrix, r, c): return matrix[r][c] == INFECTED

#returns True if person (matrix[r][c]) is contagious def contagious(matrix, r, c): return matrix[r][c] >= DAY1 and matrix[r][c] <= DAY4</pre>

#returns True if person (matrix[r][c]) is immune def immune(matrix, r, c): return matrix[r][c] == IMMUNE

Jpdating the matrix

<pre>def simNextDay(data): nextDayData = [] # create new matrix and initialize for i in range(20): nextDayData.append([0] * 20) for i in range(20): # create next day for j in range(20): if immune(data, i, j): nextDayData[i][j] = IMMUNE elif infected(data, i, j) or contagious(data, i, j): nextDayData[i][j] = data[i][j] + 1 elif healthy(data, i, j):</pre>
nextDayData[i][j] = meetPeople(data, i, j) return newMatrix

def meetPeople(currMatrix, row, col):
 for counter in range(4): # repeat 4 times
 if contagious(currMatrix, randrange(20), randrange(20)):
 return INFECTED
 return currMatrix[row][col]

Displaying the matrix

Testing display

```
def test_display():
    window = tkinter.Tk()
    # create a canvas of size 200 X 200
    c = Canvas(window,width=200,height=200)
    c.pack()
    matrix = []
    # create a randomly filled matrix
    for i in range(20):
        row = []
        for j in range(20):
            row.append(randrange(7))
        matrix.append(row)
    # display the matrix using your display function
    display(matrix,c)
```

def simulateFlu(numOfDays): window = tkinter.Tk() # create a canvas of size 200 X 200 c = Canvas(window,width=200,height=200) c.pack() # initialize matrix a to all healthy individu

initialize matrix a to all healthy individuals
population = []
for i in range(20):
 population.append([0] * 20)

infect one random person
population[randrange(20)][randrange(20)] = INFECTED
display(population, c)
sleep(0.3) # Wait in order to show 1st infection

run the simulation for required num of days for day in range(0, num0fDays): c.delete(tkinter.ALL) population = simNextDay(population) display(population,c) sleep(0.3) # Wait in order to show change c.update() #Force changes to display -update screen

What if Our Model Changes?

If a healthy person contacts a contagious person, she gets sick 40% of the time.

def meetPeople(currMatrix, row, col):

for counter in range(4): # repeat 4 times
 if contagious(currMatrix, randrange(20), randrange(20))

and randrange(100) < 40 : return INFECTED

return currMatrix[row][col]

What if Our Model Changes?(cont'd)

- The current model does not capture neighbor relationship. The adjacency of 2 cells does not indicate that they are neighbors.
- What if we used to grid to capture neighbor relationship and assumed that a healthy person gets infected if they have at least one contagious neighbor.

Neighbors

<pre>cell = matrix[i][j] north = matrix[i-1][j]</pre>	NO !
<pre>if i == 0: north = None</pre>	YES!
else: north = matrix[i-1][j]	

Next Time

Continuous simulation