Quiz Section Week 5 April 26, 2016

Review

Programming: Matrices, files, more on functions, organizing programs

Topics (not guaranteed to be comprehensive!)

• Alignments

- Reasons to align sequences
- Needleman-Wunsch algorithm
- Smith-Waterman algorithm
- Effects of parameter variation (including gap penalties)
- Testing for statistical significance of an alignment
- Phylogenetic trees
 - Rooted and unrooted topologies
 - Defining the best tree with UPGMA and Neighbor Joining
 - Concept of parsimony
 - Fitch algorithm: quantifying how parsimonious a tree is, assigning internal states
 - Finding the most parsimonious tree: Hill climbing w/ Nearest-Neighbor interchanges
 - Bootstrapping to quantify confidence in tree partitions
- Clustering
 - Defining a clustering problem
 - Hierarchical clustering
 - Impact of using single/complete/average linkage
 - K-means: Objective and algorithm
- General computation and programming
 - What is an algorithm
 - What is a search heuristic
 - Conceptual definitions of variable and function
 - Algorithm complexity with O(n) notation
 - Data types and converting between them
 - Program flow and control with conditional statements and loops

Phylogenetic trees

UPGMA/Neighbor Joining

- Define the best tree: based on distance between leaves
- Find the best tree using: polynomial time algorithm to construct the best tree from a distance matrix

Parsimony approach

- Define the best tree: Minimum # of mutations required to traverse tree
- Find the best tree: by enumerating all trees (exhaustive search), or by heuristic approach like Nearest-Neighbor Interchange Hill-Climbing

Tree topologies

Are these the same tree?



Tree topologies

Are these the same tree?



How about these?



For N leaves

```
# of unrooted topologies = 3*5*7*...*(2N-5)
# of branches = 2N-3
```

E.g. an unrooted tree with 6 nodes



How many different topologies?

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How many different topologies? 3*5*7 = **105**

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The root could be placed on any branch

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Nearest Neighbor Interchange trees



For each internal branch

generate two variant trees that swap the relationships of the four outside branches

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Nearest Neighbor Interchange trees

relationships of the four outside branches





Nearest Neighbor interchange keeps you from stepping too far in hillclimbing

Clades A and B are Tree **Negative** very closely related parsimony Score 1 -12 2 -4 3 -5 4 -76 5 -52 B 6 -30

Random designation of neighbors



Tree position on surface

NNI designation



NN Practice: Draw both interchanges from swapping this branch



NN Practice: Draw both interchanges from swapping this branch





Fitch algorithm practice



Fitch algorithm practice: bottom-up



Fitch algorithm practice: top-down



Hierarchical clustering with complete linkage example



Programming note: 2D matrices in Python

- List of lists!
- Each row is a different list

| | Α | В | С | D |
|---|---|---|---|---|
| Α | 0 | 1 | 2 | 4 |
| B | 1 | 0 | 2 | 5 |
| С | 2 | 2 | 0 | 5 |
| D | 4 | 5 | 5 | 0 |

matrix = [[0, 1, 2, 4], [1, 0, 2, 5], ...]
print matrix[0][1]
1
print matrix[0]
[0, 1, 2, 4]

Reminder: "Big O" notation for complexity What is the time complexity in O() to compute the sum of a list?

x is a list of length N
sum = 0
for v in x:
 sum = sum + v
print 'The sum is:', sum

Directly proportional to # of items in list! O(N)

How about the time complexity in O() to compute the sum of an NxN matrix?

```
# x is a list of N lists
# each list has N elements
sum = 0
for row in x: # Do this N times
    for v in row: # N times again
        sum = sum + v
print 'The sum is:', sum
# The answer is O(N^2)
```

Given a list of 2D points, compute their center

```
points = [ [1,2], [3,4], [5,6], [7,8] ]
# center point is ( mean_x, mean_y )
mean_x = 0.0
mean_y = 0.0
for i in range(0,len(points)):
    mean_x += points[i][0]
    mean_y += points[i][1]
```

```
center = ( mean_x/len(points), mean_y/len(points) )
print center
(4.0, 5.0)
```

Reading data from a file in Python

```
fin = open('qs5.txt', 'r') # 'r' stands for
'read'
all lines = []
for line in fin: # In a for loop, fin acts
like a list of strings
    print line
    all lines.append(line)
fin.close() # Lets the computer know it can
free up resources used to read the file
print all lines
```

Alternative file-reading structure

my_open_file = open(sys.argv[1])
s1 = my_open_file.readline().strip()
s2 = my_open_file.readline().strip()

Note: if in a different directory, have to supply **file path, e.g.**: python myScript.py /Users/cecilia/genome373/dataFile.txt

Writing data to a file

fout = open('output.txt', 'w') # `w'
stands for `write)
fout.write('Hello! How')
fout.write(' are you?\nI'm fine.') #
`\n' starts a
new line
fout.close()

Useful function: Split a string into its constituent words

```
s = 'Wherefore art thou Romeo?'
words = s.split() # Returns a list of substrings
print words
```

```
['Wherefore', 'art', 'thou', 'Romeo?']
```

```
# split() can use any arbitrary string to split by
words = s.split('r')
print words
['Whe', 'efo', 'e a', 't thou Romeo?']
```