# Algorithmic Approaches for Biological Data, Lecture #21

Katherine St. John

City University of New York American Museum of Natural History

25 April 2016



Guest Lecturer: Dr. Eric Ford

 Searching Graphs: Breadth-First & Depth-First Searches



Guest Lecturer: Dr. Eric Ford

- Searching Graphs: Breadth-First & Depth-First Searches
- Hill Climbing



Guest Lecturer: Dr. Eric Ford

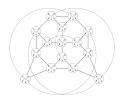
- Searching Graphs: Breadth-First & Depth-First Searches
- Hill Climbing
- Optimality Criteria & Complexity



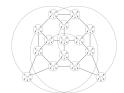
Guest Lecturer: Dr. Eric Ford

- Searching Graphs: Breadth-First & Depth-First Searches
- Hill Climbing
- Optimality Criteria & Complexity
- Parsimony Example: scoring trees in linear time

• Two common strategies:

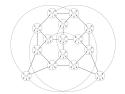


Bastert et al., 2002



Bastert et al., 2002

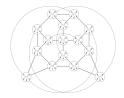
- Two common strategies:
  - ▶ Breadth First Search (BFS): visit all the neighbors, then visit all the neighbors' neighbors, etc.



Bastert et al., 2002

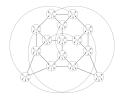
#### • Two common strategies:

- Breadth First Search (BFS): visit all the neighbors, then visit all the neighbors' neighbors, etc.
- ▶ Depth First Search (DFS): for each neighbor, visit its' neighbors, and continue as far down as possible, and repeat.



Bastert et al., 2002

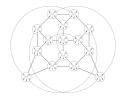
- Two common strategies:
  - Breadth First Search (BFS): visit all the neighbors, then visit all the neighbors' neighbors, etc.
  - Depth First Search (DFS): for each neighbor, visit its' neighbors, and continue as far down as possible, and repeat.
- Bookkeeping is important:



Bastert et al., 2002

#### • Two common strategies:

- Breadth First Search (BFS): visit all the neighbors, then visit all the neighbors' neighbors, etc.
- Depth First Search (DFS): for each neighbor, visit its' neighbors, and continue as far down as possible, and repeat.
- Bookkeeping is important:
  - Keep a "To Do" list (priority queue) of nodes still to visit.

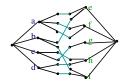


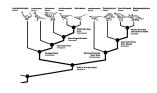
Bastert et al., 2002

#### • Two common strategies:

- Breadth First Search (BFS): visit all the neighbors, then visit all the neighbors' neighbors, etc.
- Depth First Search (DFS): for each neighbor, visit its' neighbors, and continue as far down as possible, and repeat.
- Bookkeeping is important:
  - Keep a "To Do" list (priority queue) of nodes still to visit.
  - Mark nodes as you visit them, so, you know not to visit again.

#### In Pairs: Searching Graphs

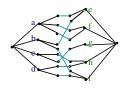


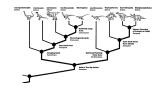


	а	Ь	С	d	е
а	0	1	1	0	0
b	0	0	1	1	0
С	1	0	0	1	0
d	0	0	1	0	1
е	0	0	0	0	0

- For above, choose start and end points to be the worst for breath first search (i.e. 'hide' the endpoint so that it takes the most number of steps to reach it).
- Por above, choose start and end points to be the worst for depth first search
- What is a graph on 6 vertices with the worst start/end pair for breath first search?
- What about for depth first search?

## Discussion: Searching Graphs

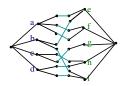


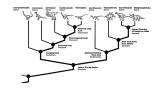


	a	Ь	С	d	е
а	0	1	1	0	0
b	0	0	1	1	0
С	1	0	0	1	0
d	0	0	1	0	1
е	0	0	0	0	0

• What makes a search difficult for BFS?

## Discussion: Searching Graphs

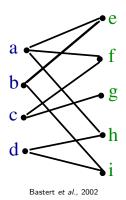


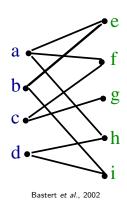


	a	Ь	С	d	е
а	0	1	1	0	0
b	0	0	1	1	0
С	1	0	0	1	0
d	0	0	1	0	1
е	0	0	0	0	0

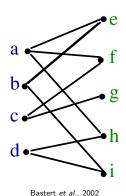
- What makes a search difficult for BFS?
- What makes a search difficult for DFS?

• How would you implement BFS?

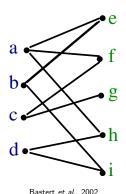




- How would you implement BFS?
  - ▶ Inputs: A graph G, a start and an end.

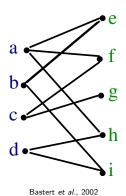


- How would you implement BFS?
  - ▶ Inputs: A graph G, a start and an end.
  - Algorithm: For each neighbor, visit its' neighbors, and continue as far down as possible, and repeat.



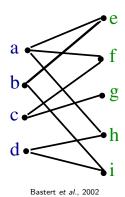
• How would you implement BFS?

- ▶ Inputs: A graph G, a start and an end.
- Algorithm: For each neighbor, visit its' neighbors, and continue as far down as possible, and repeat.
- Bookkeeping is important:

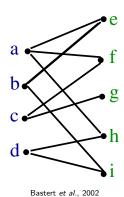


• How would you implement BFS?

- ▶ Inputs: A graph G, a start and an end.
- Algorithm: For each neighbor, visit its' neighbors, and continue as far down as possible, and repeat.
- Bookkeeping is important:
  - Keep a "To Do" list to visit and a dictionary of those already visited.



- How would you implement BFS?
  - ▶ Inputs: A graph G, a start and an end.
  - Algorithm: For each neighbor, visit its' neighbors, and continue as far down as possible, and repeat.
- Bookkeeping is important:
  - Keep a "To Do" list to visit and a dictionary of those already visited.
     toDo = [start]
    - $visited = {}$
  - Mark nodes as you visit them and add neighbors to the To Do list:



• How would you implement BFS?

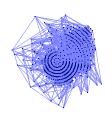
- ▶ Inputs: A graph G, a start and an end.
- Algorithm: For each neighbor, visit its' neighbors, and continue as far down as possible, and repeat.
- Bookkeeping is important:
  - Keep a "To Do" list to visit and a dictionary of those already visited. toDo = [start]

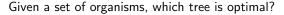
 $visited = {}$ 

Mark nodes as you visit them and add neighbors to the To Do list:

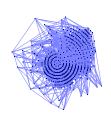
```
while len(toDo) > 0:
    nextNode = toDo.pop(0)
    visited[nextNode] = 1
    For n unvisited neighbor of nextNode,
        toDo.append(n)
```

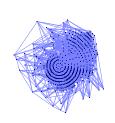
4 D > 4 A > 4 B > 4 B > B 9 9 9



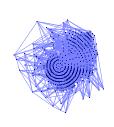


Two standard criteria for optimality:

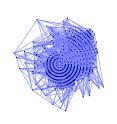




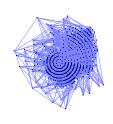
- Two standard criteria for optimality:
  - Maximum Parsimony: find tree with fewest changes



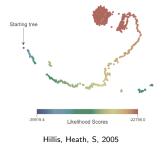
- Two standard criteria for optimality:
  - Maximum Parsimony: find tree with fewest changes
  - Maximum Likelihood: find most likely tree (with respect to a model of evolution)



- Two standard criteria for optimality:
  - Maximum Parsimony: find tree with fewest changes. (NP-hard, Foulds & Graham, 1982).
  - Maximum Likelihood: find most likely tree (with respect to a model of evolution) (NP-hard, Roch, 2008).



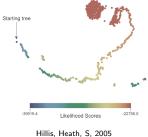
- Two standard criteria for optimality:
  - Maximum Parsimony: find tree with fewest changes. (NP-hard, Foulds & Graham, 1982).
  - Maximum Likelihood: find most likely tree (with respect to a model of evolution) (NP-hard, Roch, 2008).
- Later this lecture, we will define (and write pseudocode) for the maximum parsimony criteria.



• Goal: Find the tree with the optimal score



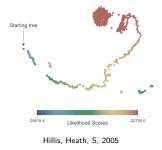
- Goal: Find the tree with the optimal score
- Local search techniques prevail:



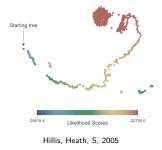
- Hills, Heath, 5, 2005
- Goal: Find the tree with the optimal score
- Local search techniques prevail:
  - ▶ Begin with a tree



- Goal: Find the tree with the optimal score
- Local search techniques prevail:
  - Begin with a tree
  - ► Choose the next tree from its neighbor (e.g. best scoring)

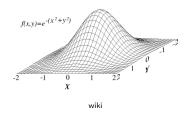


- Goal: Find the tree with the optimal score
- Local search techniques prevail:
  - Begin with a tree
  - Choose the next tree from its neighbor (e.g. best scoring)
  - Repeat



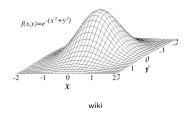
- Goal: Find the tree with the optimal score
- Local search techniques prevail:
  - Begin with a tree
  - Choose the next tree from its neighbor (e.g. best scoring)
  - Repeat
- Many variations on the theme: branch-and-bound, MCMC, genetic algorithms,...

# Hill Climbing



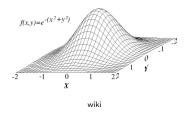
 Local Search is often Hill Climbing: at every step, you move 'up hill'.

# Hill Climbing



- Local Search is often Hill Climbing: at every step, you move 'up hill'.
- Works well if there is a single hill.

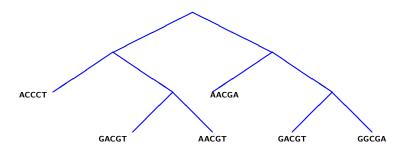
# Hill Climbing



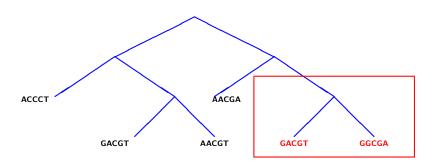
- Local Search is often Hill Climbing: at every step, you move 'up hill'.
- Works well if there is a single hill.
- If there are multiple hills, could get stuck.

• Find the tree that can explain the observed sequences with a minimal number of substitutions.

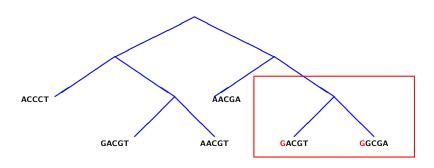
- Find the tree that can explain the observed sequences with a minimal number of substitutions.
- Given sequences for leaves and a tree, first measure "minimal number of substitutions"



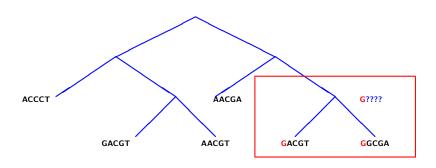
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



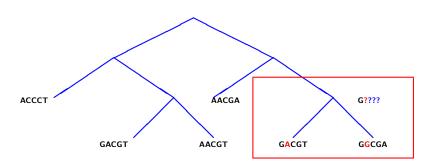
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



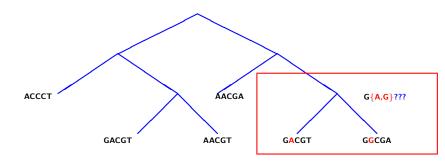
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



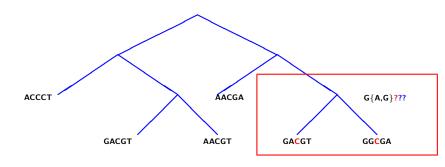
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



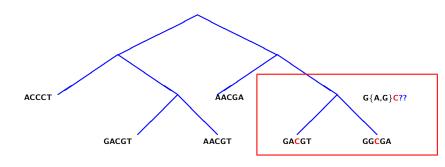
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



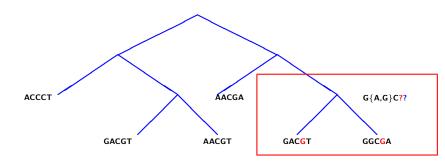
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



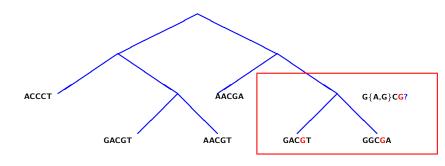
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



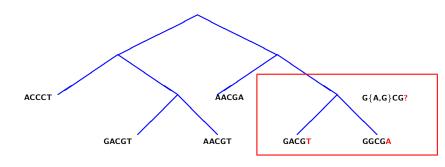
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



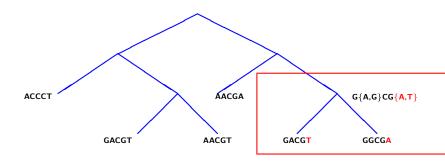
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



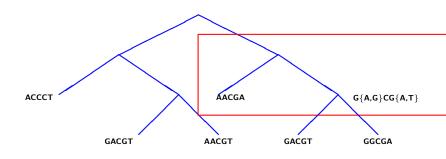
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



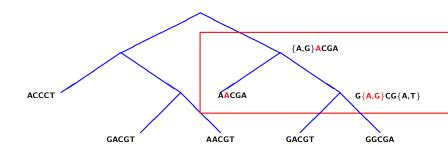
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



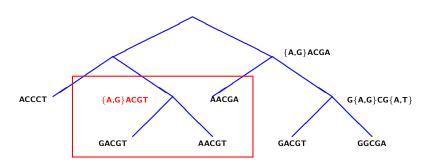
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



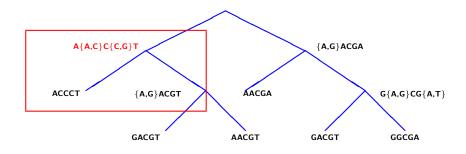
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



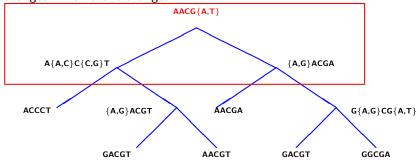
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



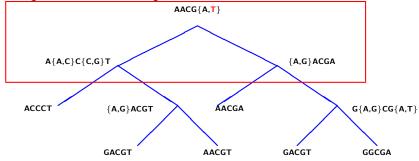
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



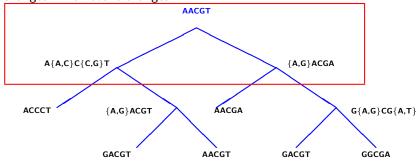
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



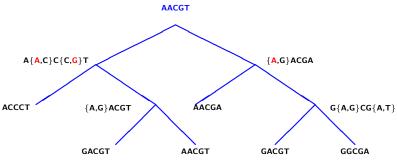
 Given sequences for leaves and a tree, first measure "minimal number of substitutions."

Label the internal nodes with sequences that have minimal number of

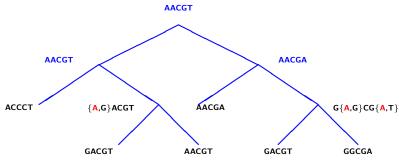
changes. Then count changes.



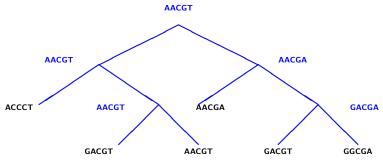
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



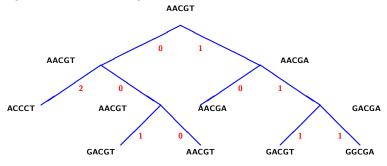
- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



- Given sequences for leaves and a tree, first measure "minimal number of substitutions."
- Label the internal nodes with sequences that have minimal number of changes. Then count changes.



Total change, called the parsimony score is 7.

• Given sequences for leaves, find tree with minimal parsimony score:

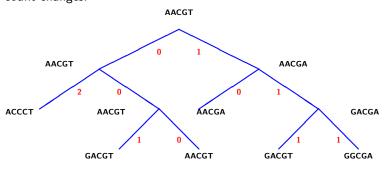
ACCCT AACGA

GACGT AACGT GACGT GGCGA

(Can you find a tree with a score better than 7?)

#### In Pairs

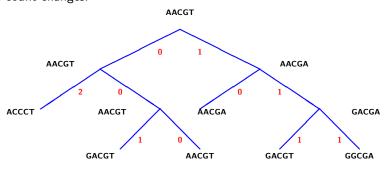
Label the internal nodes with sequences that have minimal number of changes. Then count changes.



Find a better scoring tree.

#### In Pairs

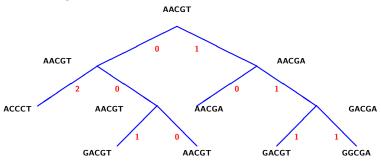
Label the internal nodes with sequences that have minimal number of changes. Then count changes.



- Find a better scoring tree.
- 2 Find the best scoring tree.

#### In Pairs

Label the internal nodes with sequences that have minimal number of changes. Then count changes.



- Find a better scoring tree.
- Find the best scoring tree.
- If all the leaves (tips) were labelled by "AAAAA", what is the best scoring tree?

• How do you code this?



- How do you code this?
  - ▶ Inputs: A tree and sequences on the leaves.



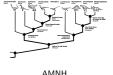
- How do you code this?
  - ▶ Inputs: A tree and sequences on the leaves.
  - Output: The parsimony score of the tree (with respect to the leaf labels).



- How do you code this?
  - Inputs: A tree and sequences on the leaves.
  - Output: The parsimony score of the tree (with respect to the leaf labels).
- What data structures do you need?

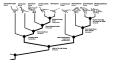


- How do you code this?
  - ▶ Inputs: A tree and sequences on the leaves.
  - Output: The parsimony score of the tree (with respect to the leaf labels).
- What data structures do you need?
  - Tree structure

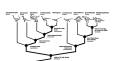


AMNE

- How do you code this?
  - ▶ Inputs: A tree and sequences on the leaves.
  - Output: The parsimony score of the tree (with respect to the leaf labels).
- What data structures do you need?
  - Tree structure
  - Count of the number changes



AMNH



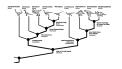
AMNH

- How do you code this?
  - ▶ Inputs: A tree and sequences on the leaves.
  - Output: The parsimony score of the tree (with respect to the leaf labels).
- What data structures do you need?
  - Tree structure
  - Count of the number changes
- Algorithm:
  - First pass: Starting at the leaves, label the internal leaves (with possible multiple labels).



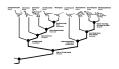
AMNH

- How do you code this?
  - ▶ Inputs: A tree and sequences on the leaves.
  - Output: The parsimony score of the tree (with respect to the leaf labels).
- What data structures do you need?
  - Tree structure
  - Count of the number changes
- Algorithm:
  - First pass: Starting at the leaves, label the internal leaves (with possible multiple labels).
  - ► Second pass: Starting at the root, choose a labeling, then work towards the leaves minimizing the conflicts.

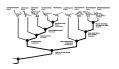


AMNH

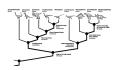
- How do you code this?
  - ▶ Inputs: A tree and sequences on the leaves.
  - Output: The parsimony score of the tree (with respect to the leaf labels).
- What data structures do you need?
  - Tree structure
  - Count of the number changes
- Algorithm:
  - First pass: Starting at the leaves, label the internal leaves (with possible multiple labels).
  - Second pass: Starting at the root, choose a labeling, then work towards the leaves minimizing the conflicts.
- If time, write out pseudocode...



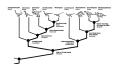
 Searching Graphs: Breadth-First & Depth-First Searches



- Searching Graphs: Breadth-First & Depth-First Searches
- Hill Climbing



- Searching Graphs: Breadth-First & Depth-First Searches
- Hill Climbing
- Optimality Criteria: Parsimony Example



- Searching Graphs: Breadth-First & Depth-First Searches
- Hill Climbing
- Optimality Criteria: Parsimony Example
- Email lab reports to kstjohn@amnh.org



- Searching Graphs: Breadth-First & Depth-First Searches
- Hill Climbing
- Optimality Criteria: Parsimony Example
- Email lab reports to kstjohn@amnh.org
- Challenges available at rosalind.info