CSCI 2270 Data Structures and Algorithms 2/24/2014 finish longest common subsequence

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Office hours: ECCS 128 or ECCS 112

Wed 9:30am-11:30am

Thurs 10:00am-11:00am

Administrivia

- First midterm exam is in grading
- We'll discuss it today in class
- Read Recursion (chapter 4); concentrate on factorial and binary search sections for now
- This week's lab is on LCS alignment
- If you are BORED and you want to go for extra credit, figure out how to extend this to backtrack and display the matching for the 2 input strings, and show it to me next week.

Finished alignment

| - | - | В | Α | N | Α | N | Α |
|---|---|---|---|---|---|---|---|
| - | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| N | 0 | 0 | 0 | 1 | 1 | 1 | 1 |
| Α | 0 | 0 | 1 | 1 | 2 | 2 | 2 |
| N | 0 | 0 | 1 | 2 | 2 | 3 | 3 |
| Α | 0 | 0 | 1 | 2 | 3 | 3 | 4 |

Backtracking to get the match is harder:

| В | Α | N | Α | N | Α |
|---|---|---|---|---|---|
| - | _ | N | Α | N | Α |

Deleting the array

```
- - B A N A N A
- 0 0 0 0 0 0 0 0
N 0 0 1 1 1 1 1
A 0 0 1 1 2 2 2
N 0 0 1 2 3 3 4
```

Deleting the array

```
В
                                N
                                                N
                                        Α
                                0
N
                        0
N
                                        2
                                                3
                                                        3
Α
        0
                0
                                        3
                                                3
```

More than one alignment

| - | - | В | 0 | N | 0 | В | 0 |
|---|---|---|---|---|---|---|---|
| - | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| В | 0 | 1 | 1 | 1 | 1 | 1 | 1 |
| 0 | 0 | 1 | 2 | 2 | 2 | 2 | 2 |
| В | 0 | 1 | 2 | 2 | 2 | 3 | 3 |
| 0 | 0 | 1 | 2 | 2 | 3 | 3 | 4 |

Backtracing the match is harder (multiple matches can happen)

| | В | 0 | N | 0 | В | 0 |
|----|---|---|---|---|---|---|
| | В | - | - | 0 | В | O |
| or | | | | | | |
| | В | 0 | N | 0 | В | 0 |
| | В | 0 | - | - | В | 0 |

Longest common subsequence

Real sequence data is hundreds to millions of letters long...

These 2D tables can get really big!

What if we had to align 3 strings? Oh, the humanity.