# **EE360C: Algorithms**

Dynamic Programming (2/4)

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#### **Least Squares**

- Foundational problem in statistic and numerical analysis
- Given *n* points in the plane  $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$ , find a line y = ax + b that minimizes the sum of the squared error:

$$SSE = \sum_{i=1}^{n} (y_i - ax_i - b)^2$$

#### Solution

The solution from calculus tells us that the minimum error is achieved when:

$$a = \frac{n \sum_{i} x_{i} y_{i} - (\sum_{i} x_{i})(\sum_{i} y_{i})}{n \sum_{i} x_{i}^{2} - (\sum_{i} x_{i})^{2}}$$

and

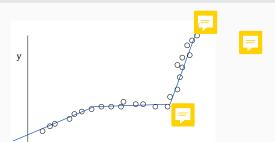
$$b = \frac{\sum_{i} y_{i} - a \sum_{i} x_{i}}{n}$$

#### **Segmented Least Squares**

- Points lie roughly on a sequence of several line segments
- Given *n* points in the plane  $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$  with  $x_1 < x_2 < \dots < x_n$ , find a sequence of lines that minimizes f(x)

#### Question

What is a reasonable choice for f(x) to balance accuracy (goodness of fit) and parsimony (number of lines)



#### The Tradeoff

We have to tradeoff the number of lines for the summed error

Find a sequence of lines that minimizes

- The sum of the sums of the squared errors E in each segment
- The number of lines L

This results in a tradeoff function E + cL for some constant c

# **Dynamic Programming: Multiway Choice**

#### **Notation**

- OPT(j) =minimum cost for points  $p_1, p_2, ..., p_j$
- $e(i,j) = \text{minimum sum of squares for points } p_i, p_{i+1}, \dots, p_j$

# **To Compute** OPT(j)

- Last segment uses points  $p_i, p_{i+1}, \dots, p_j$  for some i
- Cost = e(i, j) + c + OPT(i 1)

# **Segmented Least Squares: Algorithm**

```
INPUT: n, p<sub>1</sub>,...,p<sub>N</sub>, c

Segmented-Least-Squares() {
    M[0] = 0
    for j = 1 to n
        for i = 1 to j
            compute the least square error e<sub>ij</sub> for the segment p<sub>i</sub>,..., p<sub>j</sub>

for j = 1 to n
    M[j] = min 1 s i s j (e<sub>ij</sub> + c + M[i-1])

return M[n]
}
```

### **Running Time**

• Bottleneck: computing e(i,j) for  $O(n^2)$  pairs, O(n) per pair using the previous formula

# Knapsack

# The Knapsack Problem

#### **Knapsack Problem**

- Given n objects and a "knapsack"
- Item *i* weighs  $w_i > 0$  kilograms and has value  $v_i > 0$
- Knapsack has a capacity of W kilograms
- Goal: fill the knapsack so as to maximize the total value

#### Greedy?

Remember the greedy algorithm we explored was not optimal...

# **Knapsack: A False Start**

Definition: OPT(i) is the max profit for the subset of items  $1, \ldots, i$ 

- Case 1: OPT does not select item i
  - *OPT* selects the best of 1, 2, ..., i 1
- Case 2: OPT selects item i
  - accepting item i does not immediately imply that we will have to reject other items
  - without knowing what other items were selected before i, we don't even know if we have enough room for i

#### Conclusion

We need more subproblems!

# Dynamic Programming: Adding a Variable

Definition: OPT(i, w) is the max profit of items 1, ..., i with weight limit w

- Case 1: OPT does not select item i
  - *OPT* selects best of  $\{1, 2, ..., i 1\}$  using weight limit w
- Case 2: OPT selects item i
  - new weight limit w − w<sub>i</sub>
  - *OPT* selects best of  $\{1, 2, ..., i 1\}$  using this new weight limit

$$OPT(i, w) = \begin{cases} 0 & \text{if } i = 0 \\ OPT(i-1, w) & \text{if } w_i > w \\ \max(OPT(i-1, w), v_i + OPT(i-1, w-w_i)) & \text{otherwise} \end{cases}$$

# **Knapsack: Bottom Up**

```
Input: n, w_1, ..., w_N, v_1, ..., v_N
for w = 0 to W
  M[0, w] = 0
for i = 1 to n
   for w = 1 to W
      if (w_i > w)
          M[i, w] = M[i-1, w]
      else
          M[i, w] = \max \{M[i-1, w], v_i + M[i-1, w-w_i]\}
return M[n, W]
```

# **Knapsack: Running Time**

#### **Running Time**

- θ(nW)
- Is this a polynomial-time algorithm?
- Nope. It's "pseudo-polynomial"

**RNA Secondary Structure** 

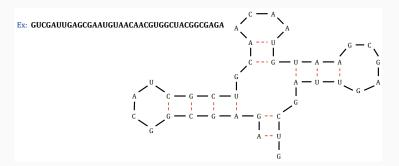
# **RNA Secondary Structure**

#### **RNA**

String  $B = b_1 b_2 \dots b_n$  over the alphabet  $\{A, C, G, U\}$ 

#### **Secondary Structure**

RNA is single stranded, so it tends to loop back and form base pairs with itself. This structure is often essential to the function of the molecule. Legal base pairs are (A, U) or (C, G).



# **RNA Secondary Structure**

#### **Secondary Structure**

A set of pairs  $S = \{(b_i, b_i)\}$  that satisfy:

- Watson and Crick: S is a matching and each pair in S is a Watson-Crick component (matching A to U or C to G)
- No Sharp Turns: The ends of each pair are separated by at least 4 intervening bases. That is, if (b<sub>i</sub>, b<sub>j</sub>) ∈ S, then i < j 4.</li>
- Non-Crossing: If  $(b_i, b_j)$  and  $(b_k, b_l)$  are two pairs in S, the we cannot have i < k < j < l

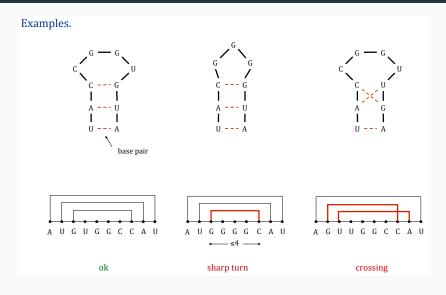
#### **Free Energy**

The usual hypothesis is that an RNA molecule will form the secondary structure that has the optimum total *free energy*, which we approximate by the number of base pairs.

#### Goal

Given an RNA molecule  $B = b_1 b_2 \dots b_n$ , find a secondary structure S that maximizes the number of base pairs.

# **RNA Secondary Structure: Examples**



# **Dynamic Programming Over Intervals**

#### **Notation**

OPT(i,j) is the maximum number of base pairs in a secondary structure of the substring  $b_i b_{i+1} \dots b_j$ 

- Case 1: if  $i \ge j 4$ 
  - OPT(i,j) = 0 by the no sharp turns condition
- Case 2: Base b<sub>i</sub> is not involved in a pair.
  - OPT(i,j) = OPT(i,j-1)
- Case 3: Base  $b_i$  pairs with  $b_t$  for some  $i \le t < j 4$ 
  - The non crossing constraint decouples the resulting subproblems
  - $OPT(i.j) = 1 + \max_t(OPT(i, t-1) + OPT(t+1, j-1))$  such that  $i \le t < j-4$  and  $(b_j, b_t)$  is a Watson and Crick pair

# **Bottom Up Dynamic Programming Over Intervals**

#### Question

What order to solve the subproblems?

#### **Answer**

Do the shortest intervals first.

# **Dynamic Programming Summary**

#### Recipe

- Characterize the structure of the problem
- · Recursively define the value of an optimal solution
- · Compute the value of the optimal solution
- Construct the optimal solution from computed information

# **Dynamic Programming Techniques**

- Binary choice: weighted interval scheduling.
- Multi-way choice segmented least squares.
- Adding a new variable: knapsack
- Dynamic programming over intervals: RNA secondary structure

#### Top-down vs. Bottom-up?

# Questions