

CSE 421: Introduction to Algorithms



Dynamic Programming

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Dynamic Programming

- **Dynamic Programming**

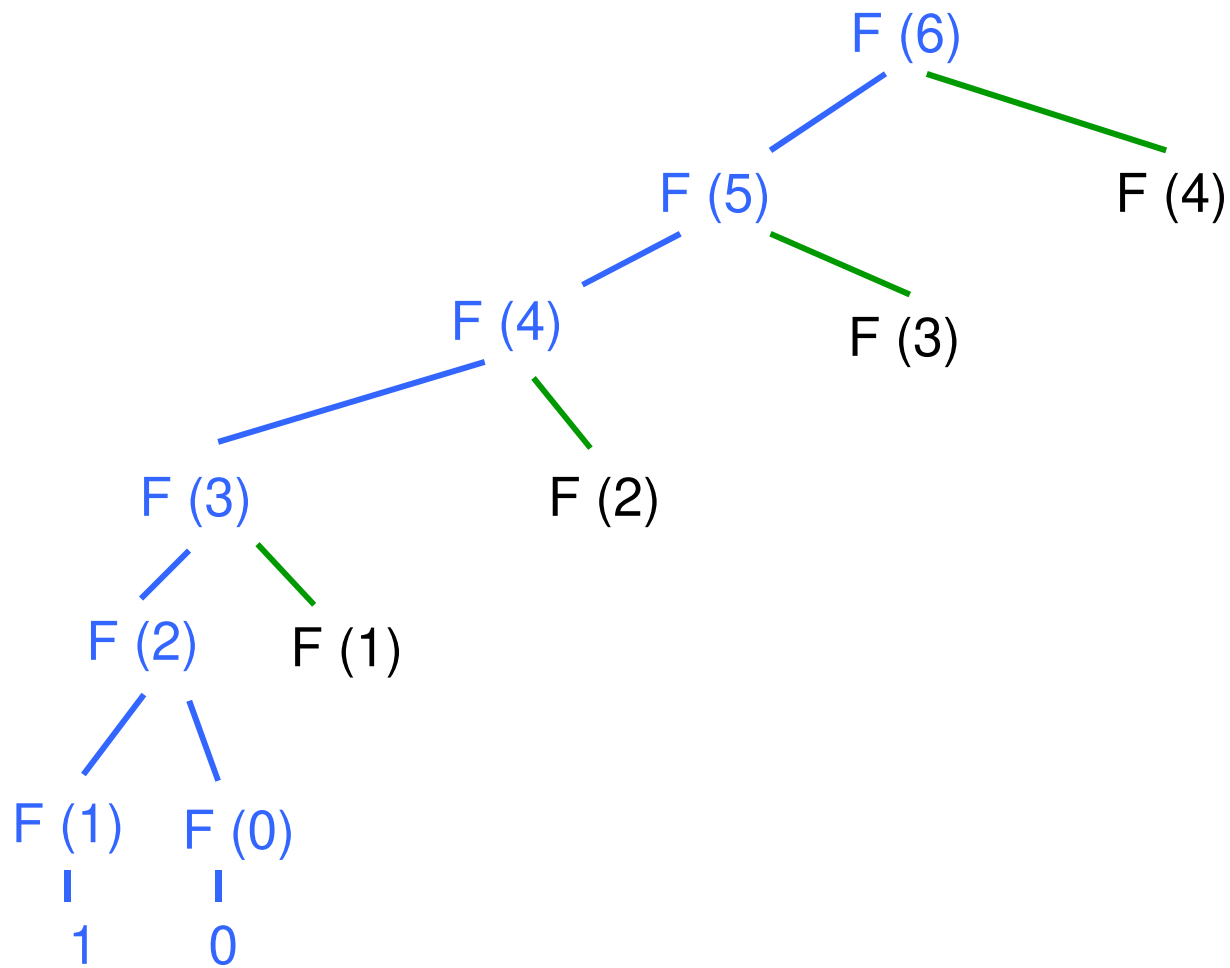
- Give a solution of a problem using smaller sub-problems where the parameters of all the possible sub-problems are determined in advance
- Useful when the same sub-problems show up again and again in the solution



A simple case: Computing Fibonacci Numbers

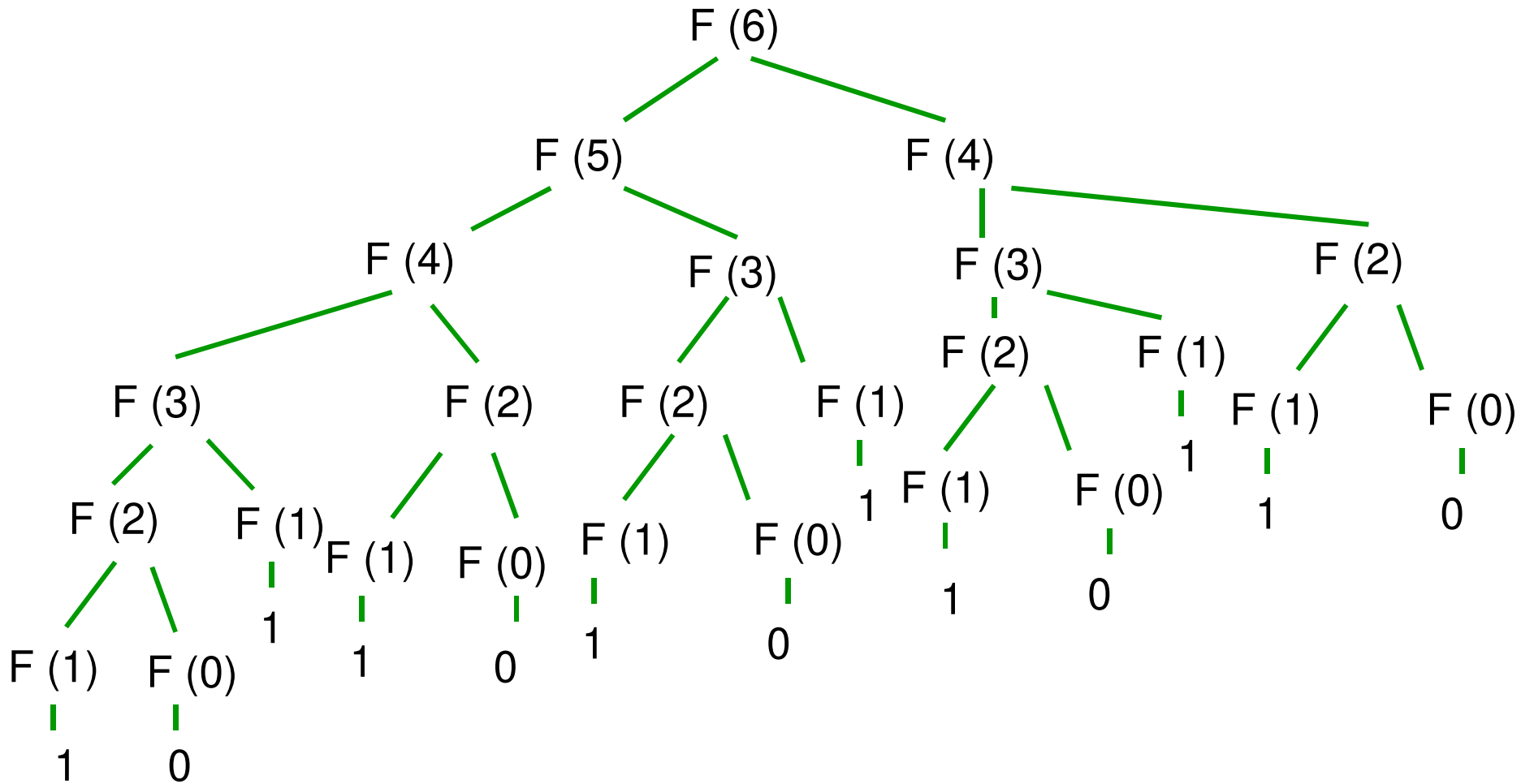
- Recall $F_n = F_{n-1} + F_{n-2}$ and $F_0 = 0$, $F_1 = 1$
- Recursive algorithm:
 - Fibo(**n**)
 - if **n=0** then return(**0**)
 - else if **n=1** then return(**1**)
 - else return(Fibo(**n-1**)+Fibo(**n-2**))

Call tree - start





Full call tree





Memoization (Caching)

- Remember all values from previous recursive calls
- Before recursive call, test to see if value has already been computed
- **Dynamic Programming**
 - Convert memoized algorithm from a recursive one to an iterative one



Fibonacci

Dynamic Programming Version

- **FiboDP(n):**
 F[0] ← 0
 F[1] ← 1
 for i=2 to n do
 F[i] ← F[i-1] + F[i-2]
 endfor
 return(F[n])



Fibonacci: Space-Saving Dynamic Programming

- **FiboDP(n):**
 prev ← 0
 curr ← 1
 for **i=2** to **n** do
 temp ← **curr**
 curr ← **curr+prev**
 prev ← **temp**
 endfor
 return(**curr**)



Dynamic Programming

- Useful when
 - same recursive sub-problems occur repeatedly
 - Can anticipate the parameters of these recursive calls
 - The solution to whole problem can be figured out with knowing the internal details of how the sub-problems are solved
 - principle of optimality
 - “Optimal solutions to the sub-problems suffice for optimal solution to the whole problem”



Three Steps to Dynamic Programming

- Formulate the answer as a recurrence relation or recursive algorithm
- Show that the number of different values of parameters in the recursive calls is “small”
 - e.g., bounded by a low-degree polynomial
 - Can use memoization
- Specify an order of evaluation for the recurrence so that you already have the partial results ready when you need them.

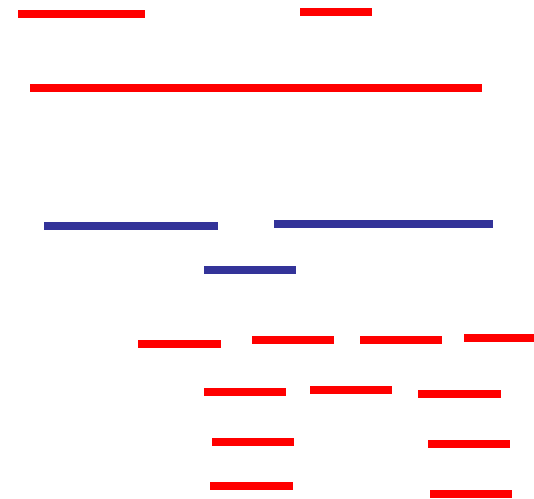


Weighted Interval Scheduling

- Same problem as interval scheduling except that each request i also has an associated **value** or **weight** w_i
 - w_i might be
 - amount of money we get from renting out the resource for that time period
 - amount of time the resource is being used $w_i = f_i - s_i$
- **Goal**: Find compatible subset **S** of requests with maximum total weight

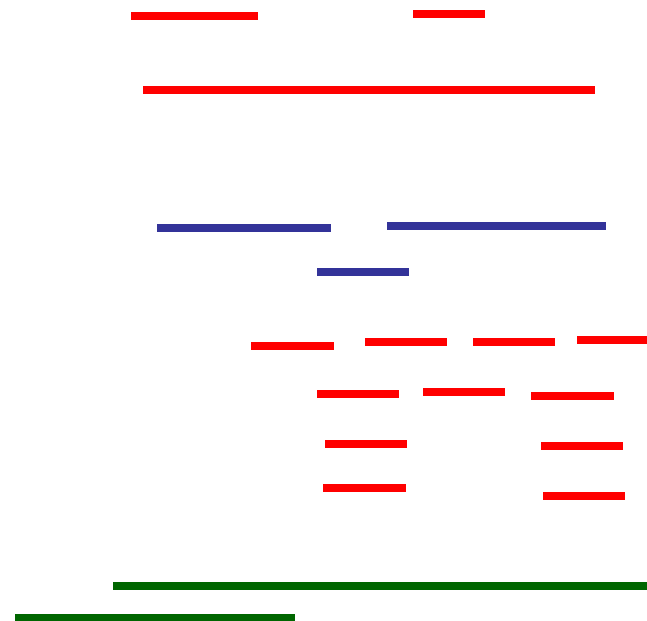
Greedy Algorithms for Weighted Interval Scheduling?

- No criterion seems to work
 - Earliest start time s_i
 - Doesn't work
 - Shortest request time $f_i - s_i$
 - Doesn't work
 - Fewest conflicts
 - Doesn't work



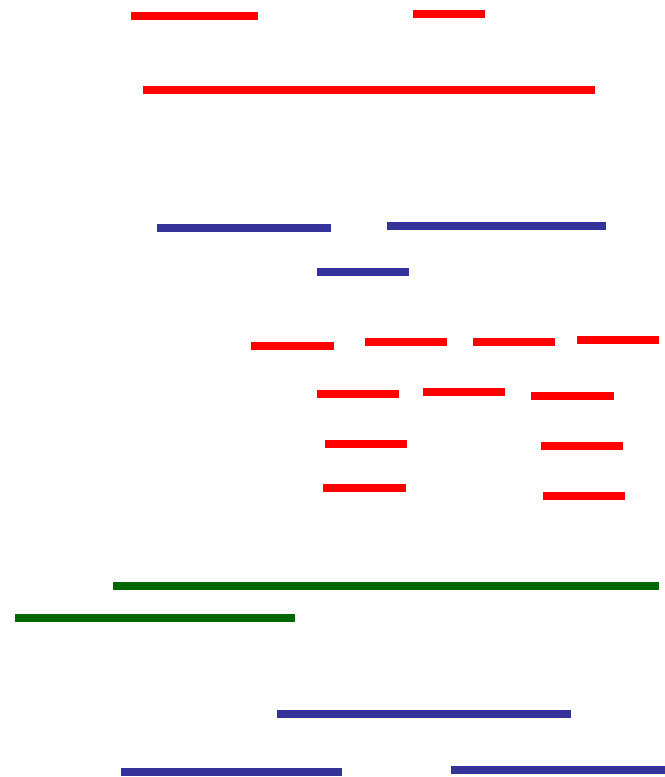
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 - Earliest finish time f_i
 - Doesn't work



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 - Shortest request time $f_i - s_i$
 - Doesn't work
 - Fewest conflicts
 - Doesn't work
 - Earliest finish time f_i
 - Doesn't work
 - Largest weight w_i
 - Doesn't work





Towards Dynamic Programming: Step 1 – A Recursive Algorithm

- Suppose that like ordinary interval scheduling we have first sorted the requests by finish time f_i so $f_1 \leq f_2 \leq \dots \leq f_n$
- Say request i comes **before** request j if $i < j$

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- For any request j let $p(j)$ be
 - the largest-numbered request before j that is compatible with j
 - or 0 if no such request exists



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- Suppose that like ordinary interval scheduling we have first sorted the requests by finish time f_i so $f_1 \leq f_2 \leq \dots \leq f_n$
- Say request i comes **before** request j if $i < j$
- For any request j let $p(j)$ be
 - the largest-numbered request before j that is compatible with j
 - or 0 if no such request exists
- Therefore $\{1, \dots, p(j)\}$ is precisely the set of requests before j that are compatible with j



Towards Dynamic Programming: Step 1 – A Recursive Algorithm

- Two cases depending on whether an optimal solution O includes request n
 - If it **does** include request n then all other requests in O must be contained in $\{1, \dots, p(n)\}$



Towards Dynamic Programming: Step 1 – A Recursive Algorithm

- Two cases depending on whether an optimal solution O includes request n
 - If it **does** include request n then all other requests in O must be contained in $\{1, \dots, p(n)\}$
 - Not only that!
 - Any set of requests in $\{1, \dots, p(n)\}$ will be compatible with request n
 - So in this case the optimal solution O must contain an optimal solution for $\{1, \dots, p(n)\}$
 - **“Principle of Optimality”**



Towards Dynamic Programming: Step 1 – A Recursive Algorithm

- All subproblems involve requests $\{1, \dots, i\}$ for some i
- For $i=1, \dots, n$ let $\mathbf{OPT}(i)$ be the **weight** of the optimal solution to the problem $\{1, \dots, i\}$
- The two cases give
$$\mathbf{OPT}(n) = \max[w_n + \mathbf{OPT}(p(n)), \mathbf{OPT}(n-1)]$$



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- The two cases give
$$\mathbf{OPT}(n) = \max[w_n + \mathbf{OPT}(p(n)), \mathbf{OPT}(n-1)]$$
- Also
 - $n \in \mathbf{O}$ iff $w_n + \mathbf{OPT}(p(n)) > \mathbf{OPT}(n-1)$



Towards Dynamic Programming: Step 1 – A Recursive Algorithm

- Sort requests and compute array $p[i]$ for each $i=1, \dots, n$

ComputeOpt(n)

if $n=0$ then return(0)

else

$u \leftarrow$ ComputeOpt($p[n]$)

$v \leftarrow$ ComputeOpt($n-1$)

 if $w_n + u > v$ then return($w_n + u$)

 else return(v)

endif



Towards Dynamic Programming: Step 2 – Small # of parameters

- **ComputeOpt(n)** can take exponential time in the worst case
 - 2^n calls if $p(i)=i-1$ for every i



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- There are only n possible parameters to **ComputeOpt**
- Store these answers in an array **OPT[n]** and only recompute when necessary
 - **Memoization**



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- There are only n possible parameters to **ComputeOpt**
- Store these answers in an array **OPT[n]** and only recompute when necessary
 - **Memoization**
- Initialize **OPT[i]=0** for $i=1, \dots, n$



Dynamic Programming: Step 2 – Memoization

ComputeOpt(**n**)

if **n=0** then return(**0**)

else

u ← MComputeOpt(**p[n]**)

v ← MComputeOpt(**n-1**)

if **w_n+u > v** then

return(**w_n+u**)

else return(**v**)

endif

MComputeOpt(**n**)

if **OPT[n]=0** then

v ← ComputeOpt(**n**)

OPT[n] ← **v**

return(**v**)

else

return(**OPT[n]**)

endif



Dynamic Programming Step 3: Iterative Solution

- The recursive calls for parameter **n** have parameter values **i** that are $< n$

IterativeComputeOpt(**n**)

array **OPT**[0..**n**]

OPT[0] ← 0

for **i**=1 to **n**

if $w_i + \mathbf{OPT}[p[i]] > \mathbf{OPT}[i-1]$ then

$\mathbf{OPT}[i] \leftarrow w_i + \mathbf{OPT}[p[i]]$

else

$\mathbf{OPT}[i] \leftarrow \mathbf{OPT}[i-1]$

endif

endfor

Producing the Solution

IterativeComputeOptSolution(**n**)

array **OPT**[0..**n**], **Used**[1..**n**]

OPT[0] ← 0

for **i**=1 to **n**

if $w_i + \mathbf{OPT}[p[i]] > \mathbf{OPT}[i-1]$ then

$\mathbf{OPT}[i] \leftarrow w_i + \mathbf{OPT}[p[i]]$

Used[**i**] ← 1

else

$\mathbf{OPT}[i] \leftarrow \mathbf{OPT}[i-1]$

Used[**i**] ← 0

endif

endfor

i ← **n**

S ← ∅

while **i** > 0 do

if **Used**[**i**] = 1 then

$\mathbf{S} \leftarrow \mathbf{S} \cup \{i\}$

i ← **p**[**i**]

else

i ← **i** - 1

endif

endwhile



Example

	1	2	3	4	5	6	7	8	9
s_i	4	2	6	8	11	15	11	12	18
f_i	7	9	10	13	14	17	18	19	20
w_i	3	7	4	5	3	2	7	7	2
$p[i]$									
OPT[i]									
Used[i]									



Example

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w_i	3	7	4	5	3	2	7	7	2
$p[i]$	0	0	0	1	3	5	3	3	7
OPT[i]	3	7	7	8	10	12	14	14	16
Used[i]	1	1	0	1	1	1	1	0	1



Example

	1	2	3	4	5	6	7	8	9
s_i	4	2	6	8	11	15	11	12	18
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$p[i]$	0	0	0	1	3	5	3	3	7
OPT[i]	3	7	7	8	10	12	14	14	16
Used[i]	1	1	0	1	1	1	1	0	1

$S = \{9, 7, 2\}$

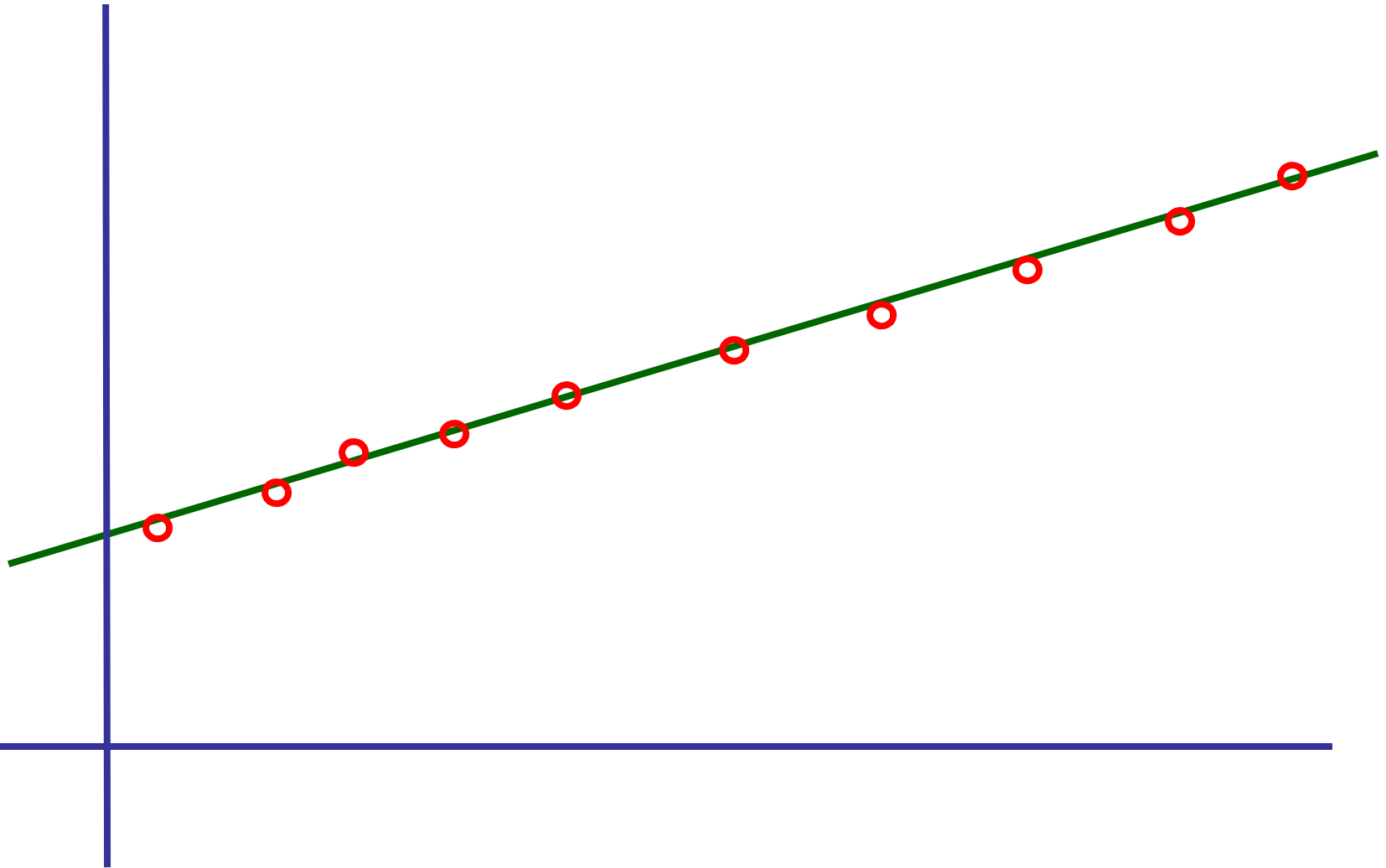


Segmented Least Squares

■ Least Squares

- Given a set \mathbf{P} of n points in the plane $\mathbf{p}_1=(x_1,y_1), \dots, \mathbf{p}_n=(x_n,y_n)$ with $x_1 < \dots < x_n$ determine a line \mathbf{L} given by $y=ax+b$ that optimizes the total ‘squared error’
 - $\text{Error}(\mathbf{L}, \mathbf{P}) = \sum_i (y - ax_i - b)^2$
- A classic problem in statistics
- Optimal solution is known (see text)
 - Call this $\text{line}(\mathbf{P})$ and its error $\text{error}(\mathbf{P})$

Least Squares

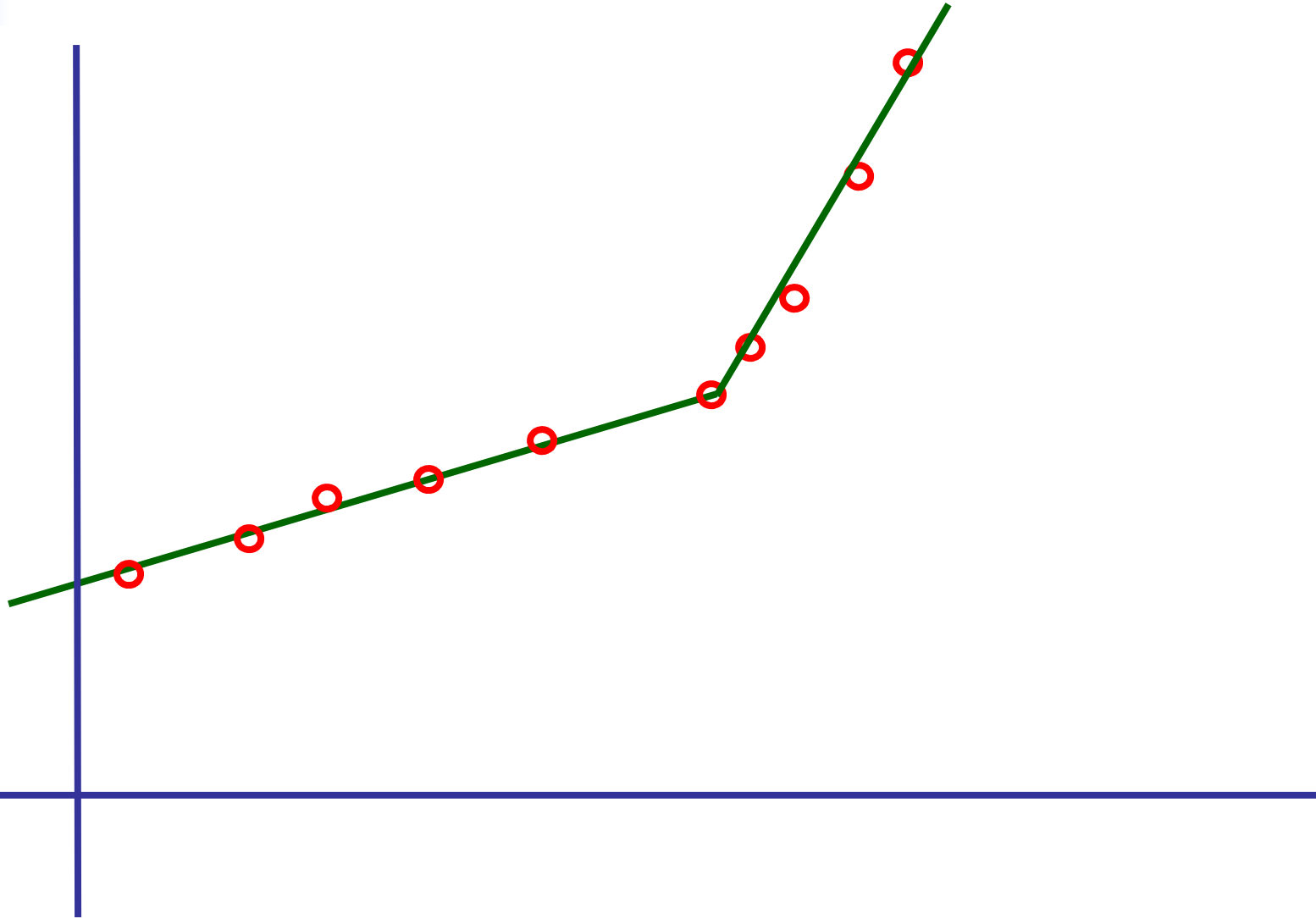




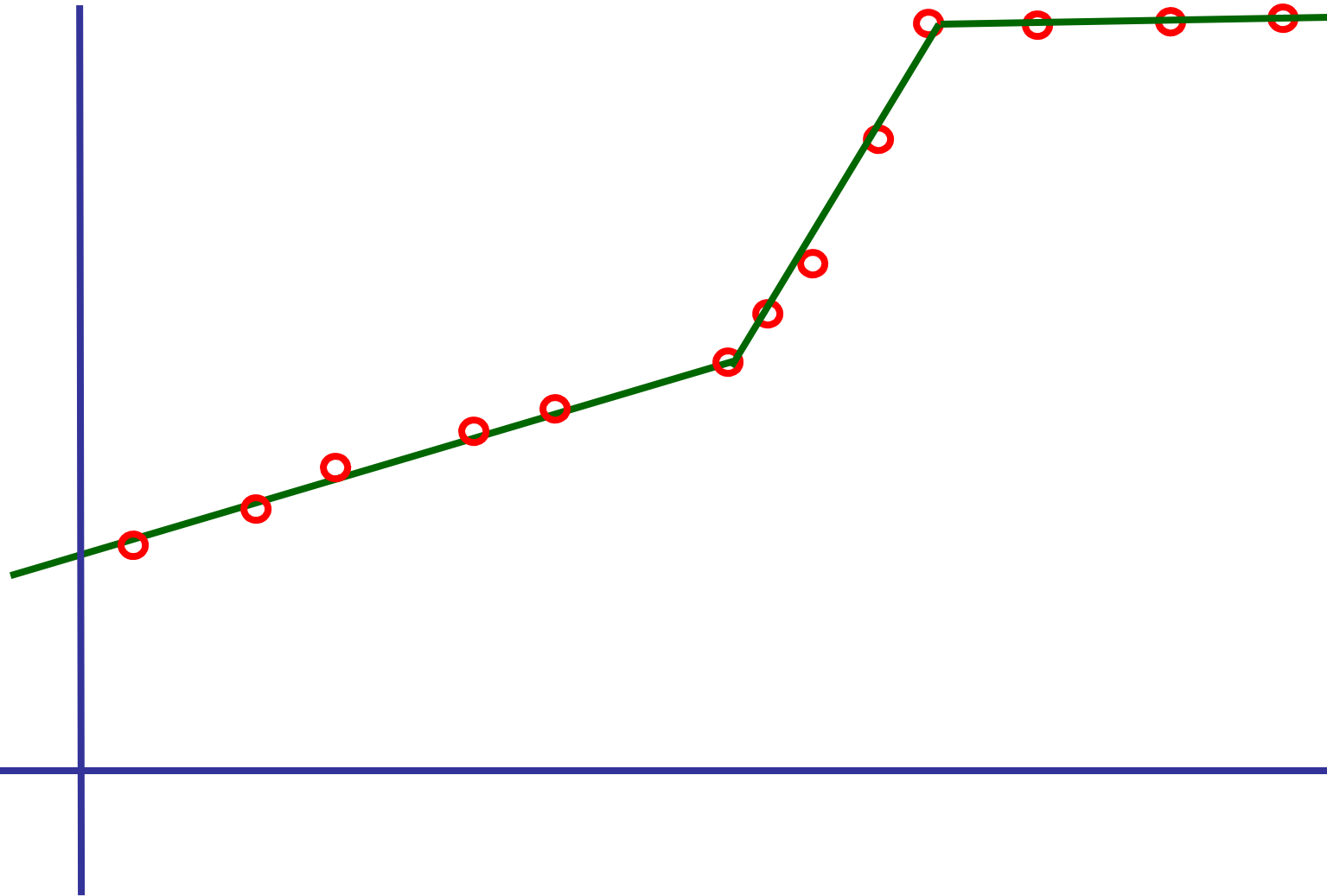
Segmented Least Squares

- What if data seems to follow a piece-wise linear model?

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- If we chose **$n-1$** pieces we could fit with **0** error
 - Not a fair measure of data fit



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- What if data seems to follow a piece-wise linear model?
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 - Not a fair measure of data fit
- Add a penalty of **C** times the number of pieces to the error to get a **total penalty**



Segmented Least Squares

- What if data seems to follow a piece-wise linear model?
- Number of pieces to choose is not obvious
- If we chose **$n-1$** pieces we could fit with **0** error
 - Not a fair measure of data fit
- Add a penalty of **C** times the number of pieces to the error to get a **total penalty**
- How do we compute a solution with the smallest possible total penalty?



Segmented Least Squares

- Recursive idea
 - If we knew the point p_j where the **last** line segment began then we could solve the problem optimally for points p_1, \dots, p_j and combine that with the last segment to get a global optimal solution



Segmented Least Squares

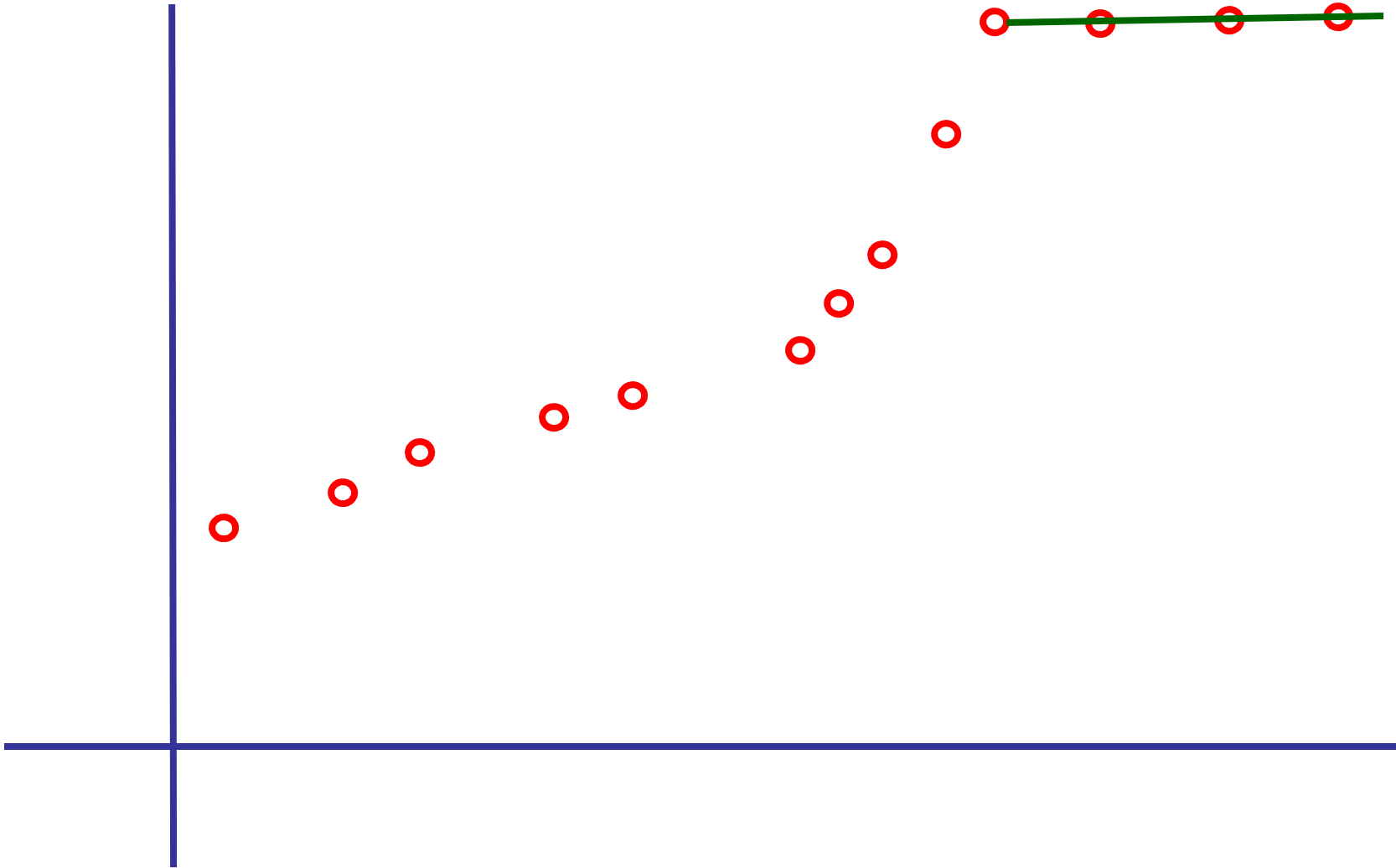
- Recursive idea
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 - Let $OPT(j)$ be the optimal penalty for points $\{p_1, \dots, p_j\}$



Segmented Least Squares

- Recursive idea
 - If we knew the point p_j where the **last** line segment began then we could solve the problem optimally for points p_1, \dots, p_j and combine that with the last segment to get a global optimal solution
 - Let $OPT(j)$ be the optimal penalty for points $\{p_1, \dots, p_j\}$
 - Total penalty for this solution would be $Error(\{p_j, \dots, p_n\}) + C + OPT(j-1)$

Segmented Least Squares





Segmented Least Squares

- Recursive idea
 - We don't know which point is p_j
 - But we do know that $1 \leq j \leq n$
 - The optimal choice will simply be the best among these possibilities



Segmented Least Squares

- Recursive idea
 - We don't know which point is \mathbf{p}_j
 - But we do know that $1 \leq j \leq n$
 - The optimal choice will simply be the best among these possibilities
 - Therefore

$$\text{OPT}(\mathbf{n}) = \min_{1 \leq j \leq n} \{ \text{Error}(\{\mathbf{p}_j, \dots, \mathbf{p}_n\}) + \mathbf{C} + \text{OPT}(j-1) \}$$



Dynamic Programming Solution

SegmentedLeastSquares(**n**)
array **OPT**[0..**n**]

OPT[0] ← 0

for **i**=1 to **n**

OPT[**i**] ← **Error**{(**p**₁, ..., **p**_{**i**})} + **C**

 for **j**=2 to **i**-1

e ← **Error**{(**p**_{**j**}, ..., **p**_{**i**})} + **C** + **OPT**[**j**-1]

 if **e** < **OPT**[**i**] then

OPT[**i**] ← **e**

 endif

 endfor

endfor

return(**OPT**[**n**])



Dynamic Programming Solution

```
SegmentedLeastSquares(n)
  array OPT[0..n]
  array Begin[1..n]
  OPT[0] ← 0
  for i=1 to n
    OPT[i] ← Error{(p1, ..., pi)} + C
    Begin[i] ← 1
    for j=2 to i-1
      e ← Error{(pj, ..., pi)} + C + OPT[j-1]
      if e < OPT[i] then
        OPT[i] ← e
        Begin[i] ← j
      endif
    endfor
  endfor
  return(OPT[n])
```



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      endif
    endfor
  endfor
  return(OPT[n])
```

```
FindSegments
i ← n
S ← ∅
while i > 1 do
  compute Line({pBegin[i], ..., pi})
  output (pBegin[i], pi), Line
  i ← Begin[i]
endwhile
```



Knapsack (Subset-Sum) Problem

- Given:
 - integer **W** (knapsack size)
 - **n** object sizes **x_1, x_2, \dots, x_n**
- Find:
 - Subset **S** of **$\{1, \dots, n\}$** such that $\sum_{i \in S} x_i \leq W$
but $\sum_{i \in S} x_i$ is as large as possible



Recursive Algorithm

- Let $K(n, W)$ denote the problem to solve for W and x_1, x_2, \dots, x_n
- For $n > 0$,
 - The optimal solution for $K(n, W)$ is the better of the optimal solution for either $K(n-1, W)$ or $x_n + K(n-1, W - x_n)$

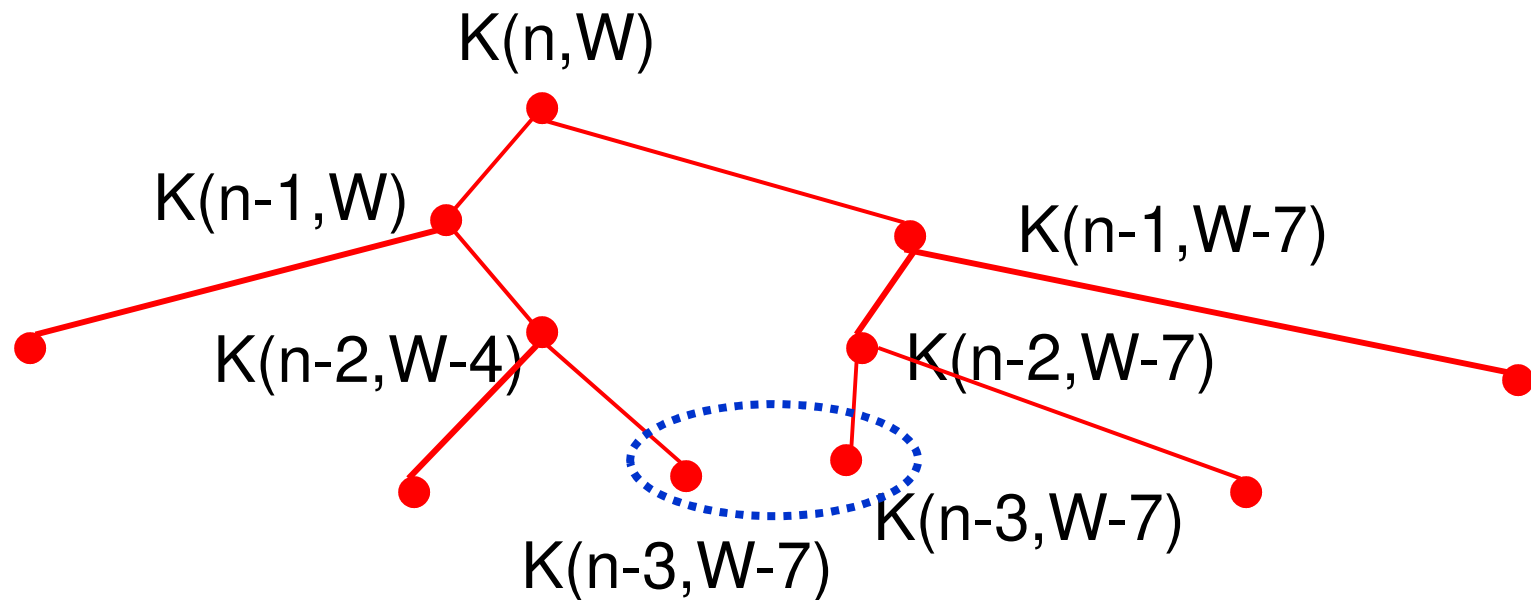


Recursive Algorithm

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- For $n > 0$,
 - The optimal solution for $K(n, W)$ is the better of the optimal solution for either $K(n-1, W)$ or $x_n + K(n-1, W - x_n)$
- For $n = 0$
 - $K(0, W)$ has a trivial solution of an empty set S with weight 0

Recursive calls

- Recursive calls on list ..., 3, 4, 7





Common Sub-problems

- Only sub-problems are $K(i,w)$ for
 - $i = 0, 1, \dots, n$
 - $w = 0, 1, \dots, W$
- Dynamic programming solution
 - Table entry for each $K(i,w)$
 - **OPT** - value of optimal soln for first i objects and weight w
 - **belong** flag - is x_i a part of this solution?



Common Sub-problems

- Only sub-problems are $K(i,w)$ for
 - $i = 0,1,\dots, n$
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- Dynamic programming solution
 - Table entry for each $K(i,w)$
 - **OPT** - value of optimal soln for first i objects and weight w
 - **belong** flag - is x_i a part of this solution?
 - Initialize **OPT**[0, w] for $w=0,\dots,W$
 - Compute all **OPT**[$i,*$] from **OPT**[$i-1,*$] for $i>0$



Dynamic Knapsack Algorithm

```
for  $w=0$  to  $W$ ;  $OPT[0,w] \leftarrow 0$ ; end for
for  $i=1$  to  $n$  do
  for  $w=0$  to  $W$  do
     $OPT[i,w] \leftarrow OPT[i-1,w]$ 
     $belong[i,w] \leftarrow 0$ 
    if  $w \geq x_i$  then
       $val \leftarrow x_i + OPT[i-1,w-x_i]$ 
      if  $val > OPT[i,w]$  then
         $OPT[i,w] \leftarrow val$ 
         $belong[i,w] \leftarrow 1$ 
    end for
  end for
end for
return( $OPT[n,W]$ )
```

Time $O(nW)$



Sample execution on 2, 3, 4, 7 with $W=15$



Saving Space

- To compute the value **OPT** of the solution only need to keep the last two rows of **OPT** at each step
- What about determining the set **S**?
 - Follow the **belong** flags **O(n)** time
 - What about space?



Three Steps to Dynamic Programming

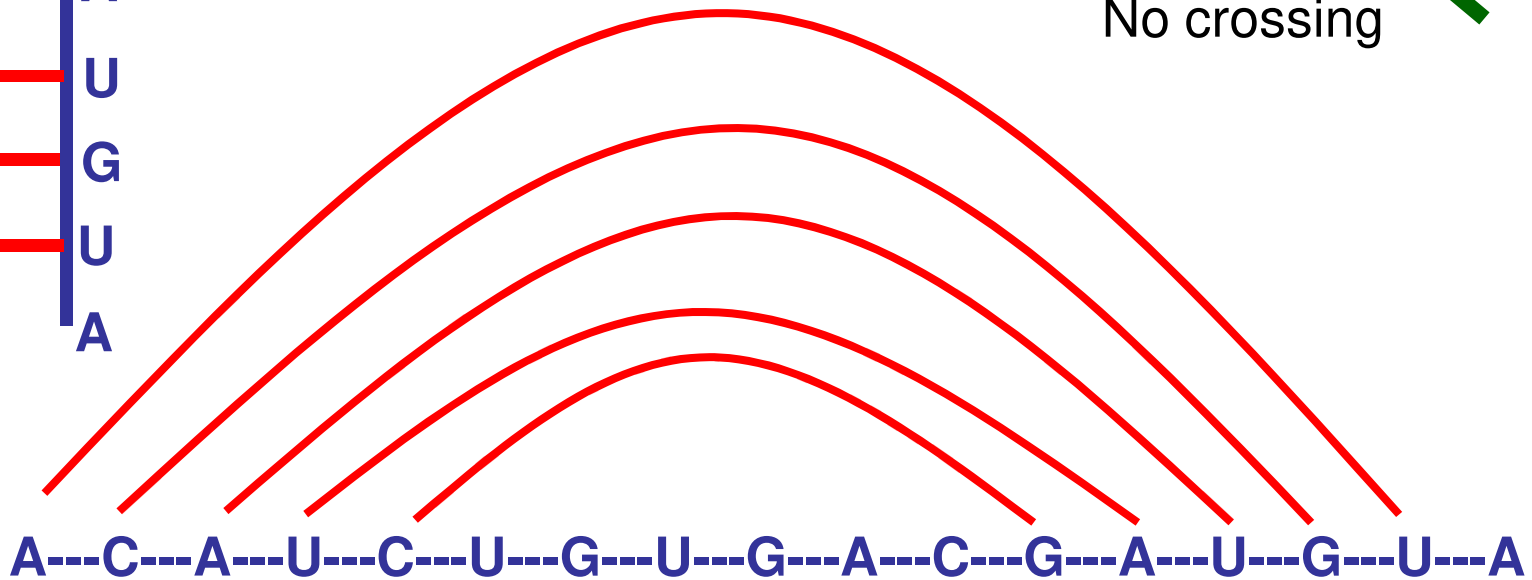
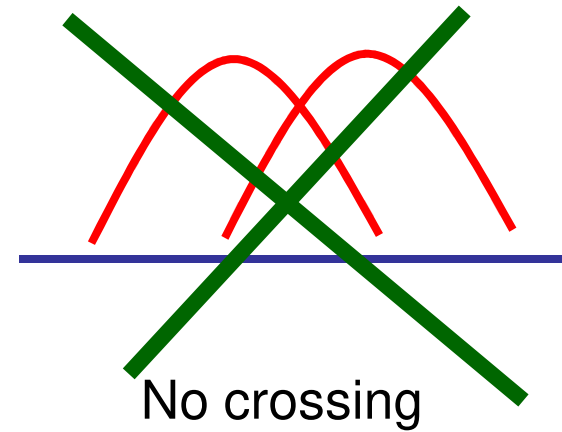
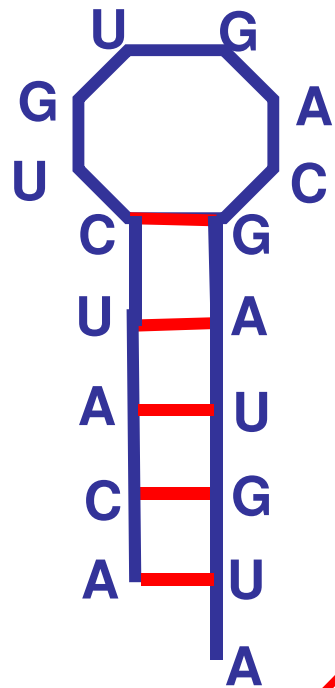
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- Show that the number of different values of parameters in the recursive algorithm is “small”
 - e.g., bounded by a low-degree polynomial
- Specify an order of evaluation for the recurrence so that you already have the partial results ready when you need them.



RNA Secondary Structure: Dynamic Programming on Intervals

- RNA: sequence of bases
 - String over alphabet {**A**, **C**, **G**, **U**}
 - U-G-U-A-C-C-G-G-U-A-G-U-A-C-A**
- RNA folds and sticks to itself like a zipper
 - **A** bonds to **U**
 - **C** bonds to **G**
 - Bends can't be sharp
 - No twisting or criss-crossing
- How the bonds line up is called the **RNA secondary structure**

Another view of RNA Secondary Structure



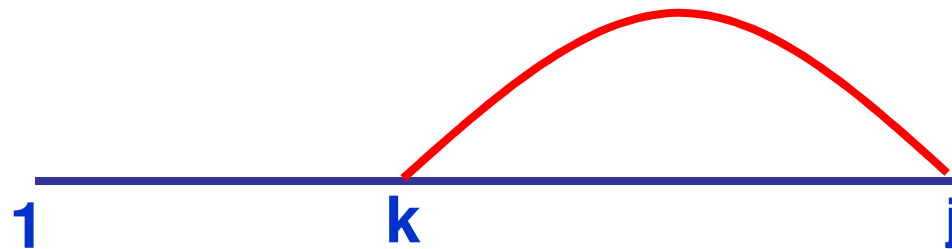


RNA Secondary Structure

- **Input:** String $x_1 \dots x_n \in \{A, C, G, U\}^*$
- **Output:** Maximum size set **S** of pairs (i, j) such that
 - $\{x_i, x_j\} = \{A, U\}$ or $\{x_i, x_j\} = \{C, G\}$
 - The pairs in **S** form a matching
 - $i < j - 4$ (no sharp bends)
 - No crossing pairs
 - If (i, j) and (k, l) are in **S** then it is not the case that they cross as in $i < k < j < l$

Recursion Solution

- Try all possible matches for the last bas

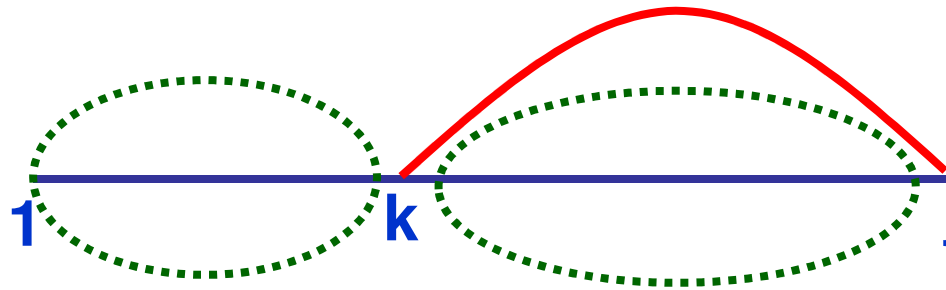


$$\text{OPT}(1..j) = \text{MAX}(\text{OPT}(1..j-1), 1 + \text{MAX}_{k=1..j-5} (\text{OPT}(1..k-1) + \text{OPT}(k+1..j-1)))$$

x_k matches x_j

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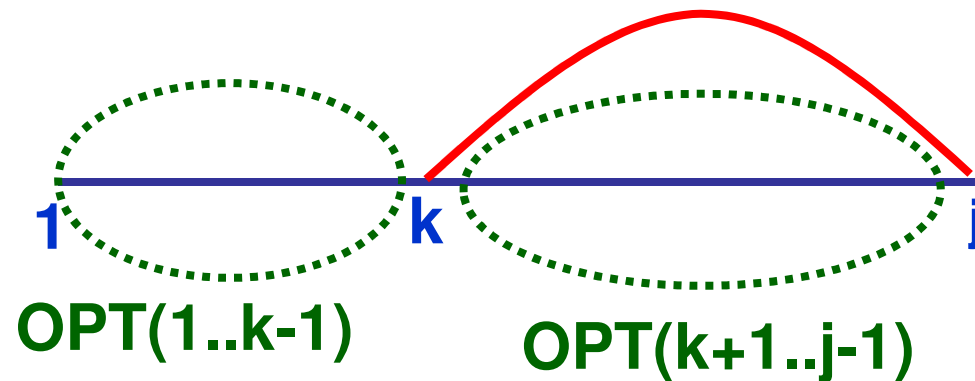


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- Try all possible matches for the last base

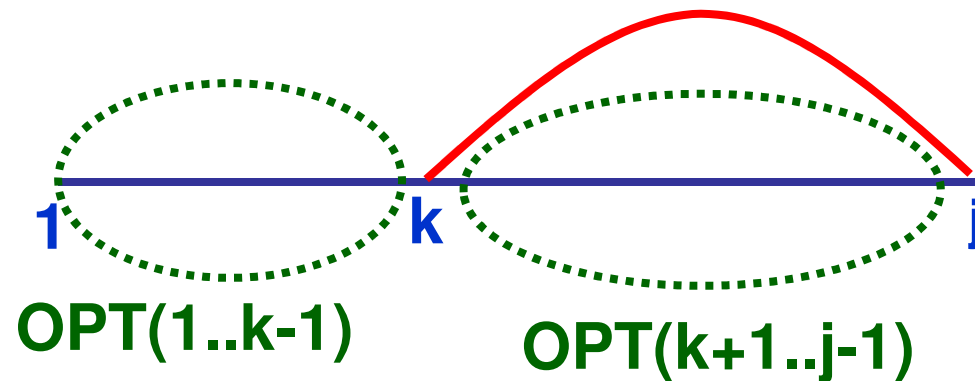


$$OPT(1..j) = \text{MAX}(OPT(1..j-1), 1 + \text{MAX}_{k=1..j-5} (OPT(1..k-1) + OPT(k+1..j-1)))$$

x_k matches x_j

Recursion Solution

- Try all possible matches for the last base



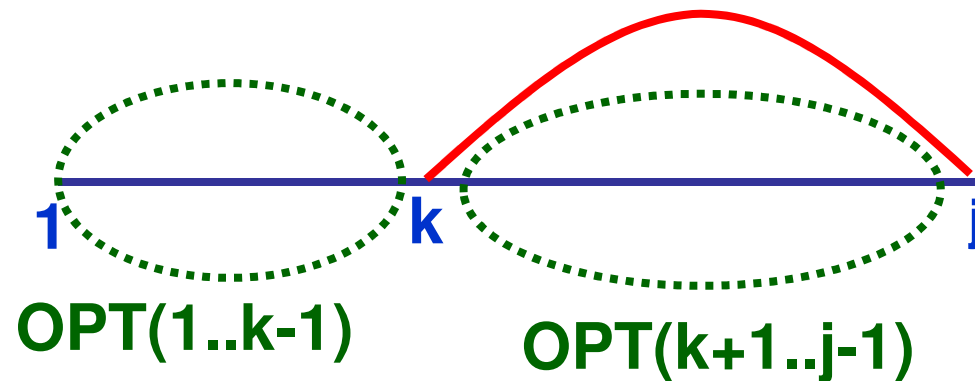
$$OPT(1..j) = \text{MAX}(OPT(1..j-1), 1 + \text{MAX}_{k=1..j-5} (OPT(1..k-1) + OPT(k+1..j-1)))$$

x_k matches x_j

Doesn't start at 1

Recursion Solution

- Try all possible matches for the last base



$$\text{OPT}(1..j) = \text{MAX}(\text{OPT}(1..j-1), 1 + \text{MAX}_{k=1..j-5} (\text{OPT}(1..k-1) + \text{OPT}(k+1..j-1)))$$

x_k matches x_j

Doesn't start at 1

General form:

$$\text{OPT}(i..j) = \text{MAX}(\text{OPT}(i..j-1), 1 + \text{MAX}_{k=i..j-5} (\text{OPT}(i..k-1) + \text{OPT}(k+1..j-1)))$$

x_k matches x_j



RNA Secondary Structure

- 2D Array **OPT(i,j)** for $i \leq j$ represents optimal # of matches entirely for segment $i..j$
- For $j-i \leq 4$ set **OPT(i,j)=0** (no sharp bends)
- Then compute **OPT(i,j)** values when $j-i=5,6,\dots,n-1$ in turn using recurrence.
- Return **OPT(1,n)**
- Total of **$O(n^3)$** time
- Can also record matches along the way to produce **S**
 - Similar polynomial-time algorithm for other problems
 - Context-Free Language recognition
 - Optimal matrix products, etc.
 - All use dynamic programming over intervals



Sequence Alignment: Edit Distance

- **Given:**

- Two strings of characters $A = a_1 a_2 \dots a_n$ and $B = b_1 b_2 \dots b_m$

- **Find:**

- The minimum number of edit steps needed to transform **A** into **B** where an edit can be:
 - **insert** a single character
 - **delete** a single character
 - **substitute** one character by another



Applications

- "diff" utility – where do two files differ
- Version control & patch distribution – save/send only changes
- Molecular biology
 - Similar sequences often have similar origin and function
 - Similarity often recognizable despite millions or billions of years of evolutionary divergence

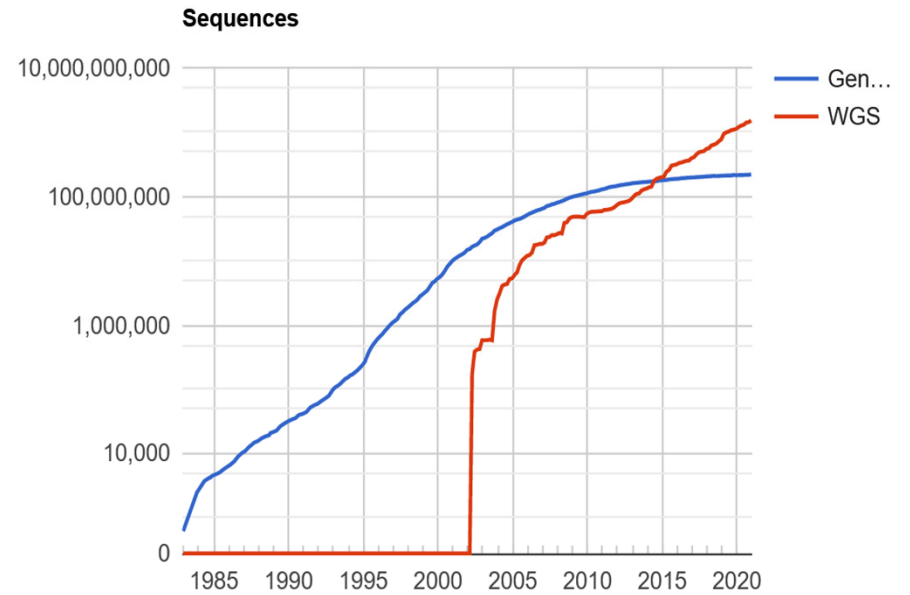
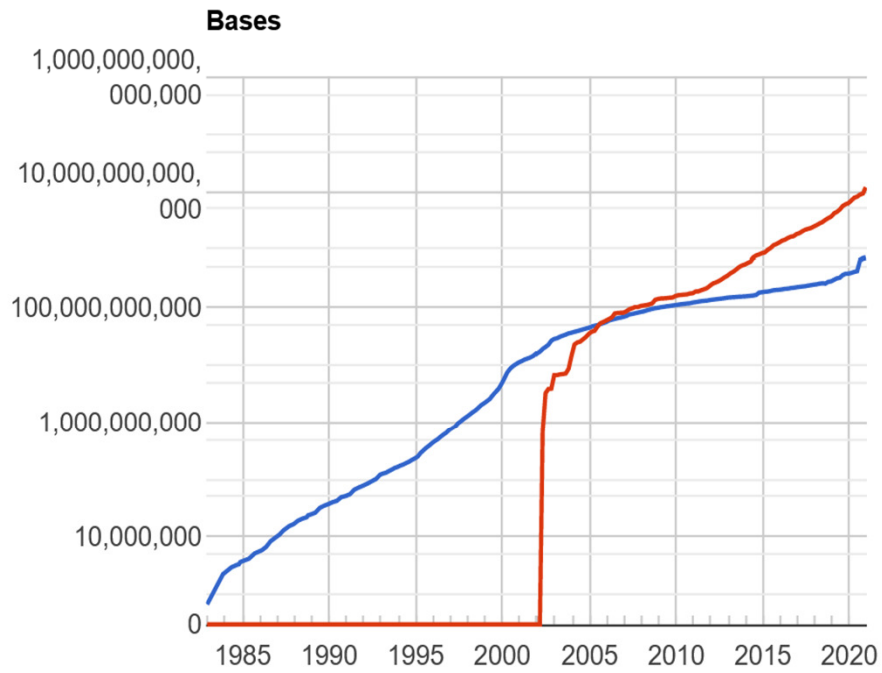
```
C A - C G T G A T
| |   | |   | X |
C A T C G - G T T
```



Sequence Alignment vs Edit Distance

- Sequence Alignment
 - Insert corresponds to aligning with a “-” in the first string
 - Cost δ (in our case **1**)
 - Delete corresponds to aligning with a “-” in the second string
 - Cost δ (in our case **1**)
 - Replacement of an **a** by a **b** corresponds to a mismatch
 - Cost α_{ab} (in our case **1** if **a**≠**b** and **0** if **a**=**b**)
- In Computational Biology this alignment algorithm is attributed to Smith & Waterman

GenBank and WGS Statistics





Recursive Solution

- **Sub-problems:** Edit distance problems for **all prefixes** of **A** and **B** that don't include all of both **A** and **B**
- Let **$D(i,j)$** be the number of edits required to transform **$a_1 a_2 \dots a_i$** into **$b_1 b_2 \dots b_j$**
- Clearly **$D(0,0)=0$**



Computing $D(n,m)$

- Imagine how best sequence handles the last characters a_n and b_m
- If best sequence of operations
 - deletes a_n then $D(n,m)=D(n-1,m)+1$
 - inserts b_m then $D(n,m)=D(n,m-1)+1$
 - replaces a_n by b_m then
$$D(n,m)=D(n-1,m-1)+1$$
 - matches a_n and b_m then
$$D(n,m)=D(n-1,m-1)$$



Recursive algorithm $D(n,m)$

if $n=0$ then

 return (m)

elseif $m=0$ then

 return(n)

else

 if $a_n=b_m$ then

replace-cost $\leftarrow 0$

 else

replace-cost $\leftarrow 1$

} cost of substitution of a_n by b_m (if used)

endif

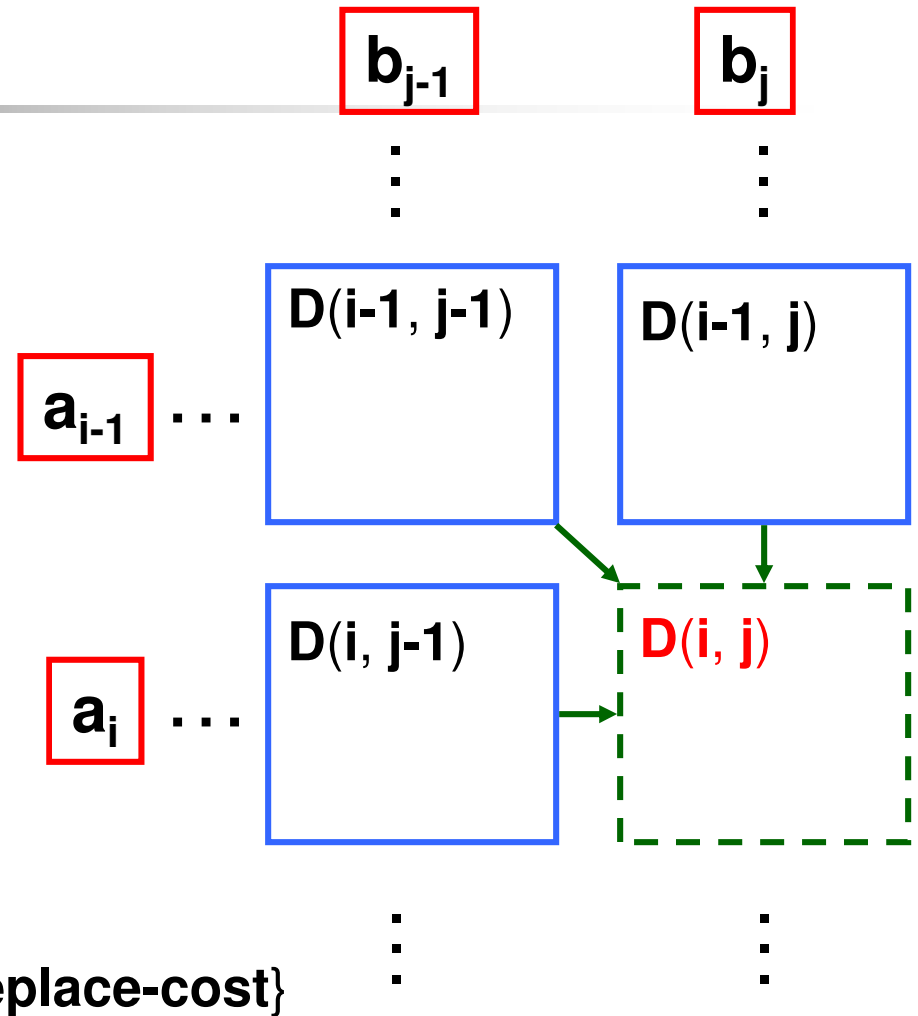
return(**min**{ $D(n-1, m) + 1$,
 $D(n, m-1) + 1$,
 $D(n-1, m-1) + \text{replace-cost}$ })

Dynamic Programming

```

for j = 0 to m; D(0,j) ← j; endfor
for i = 1 to n; D(i,0) ← i; endfor
for i = 1 to n
  for j = 1 to m
    if ai=bj then
      replace-cost ← 0
    else
      replace-cost ← 1
    endif
    D(i,j) ← min { D(i-1, j) + 1,
                  D(i, j-1) + 1,
                  D(i-1, j-1) + replace-cost }
  endfor
endfor

```





Example run with AGACATTG and GAGTTA

		A	G	A	C	A	T	T	G
	0	1	2	3	4	5	6	7	8
0									
G 1									
A 2									
G 3									
T 4									
T 5									
A 6									



Example run with AGACATTG and GAGTTA

	A	G	A	C	A	T	T	G	
G	0	1	2	3	4	5	6	7	8
A	1	1	1	2	3	4	5	6	7
G	2								
T	3								
T	4								
A	5								
A	6								



Example run with AGACATTG and GAGTTA

	A	G	A	C	A	T	T	G	
G	0	1	2	3	4	5	6	7	8
A	1	1	1	2	3	4	5	6	7
G	2	1	2	1					
T	3								
T	4								
A	5								
A	6								



Example run with AGACATTG and GAGTTA

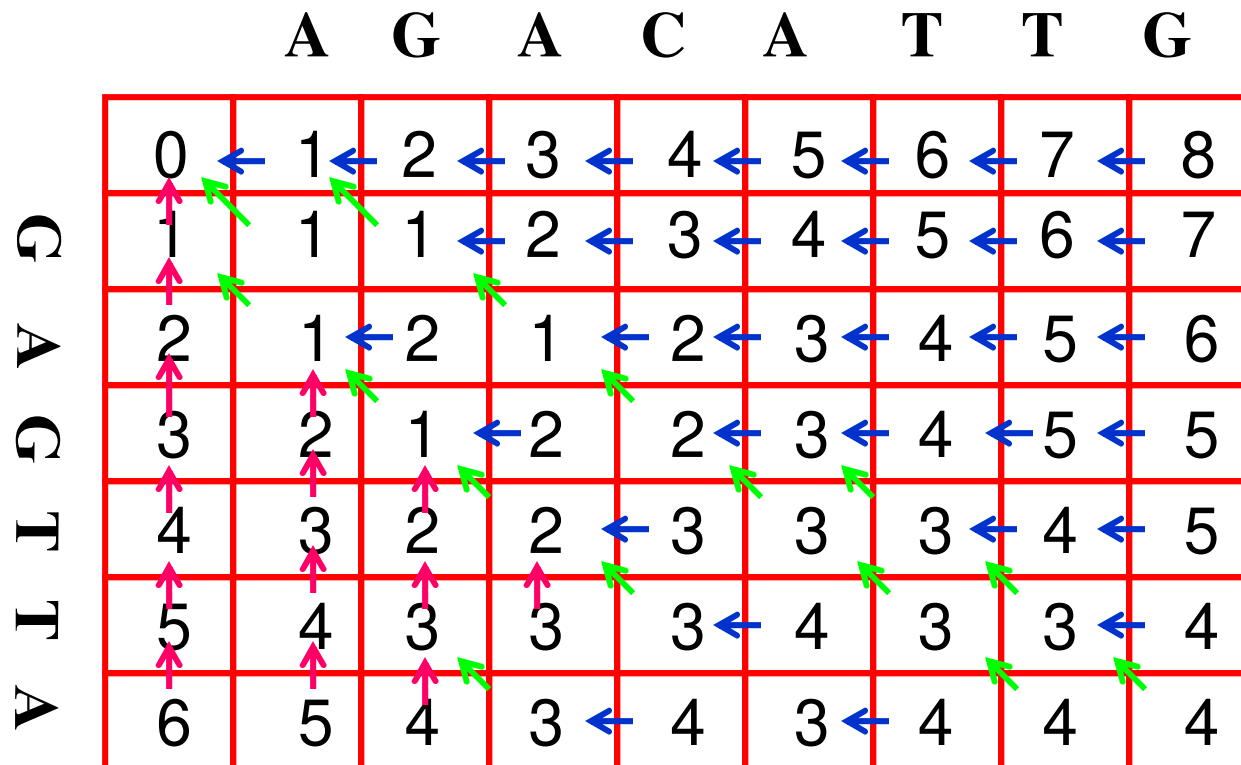
	A	G	A	C	A	T	T	G	
G	0	1	2	3	4	5	6	7	8
A	1	1	1	2	3	4	5	6	7
G	2	1	2	1	2	3	4	5	6
T	3	2	1	2	2	3	4	5	5
T	4								
A	5								
A	6								



Example run with AGACATTG and GAGTTA

	A	G	A	C	A	T	T	G	
G	0	1	2	3	4	5	6	7	8
A	1	1	1	2	3	4	5	6	7
G	2	1	2	1	2	3	4	5	6
T	3	2	1	2	2	3	4	5	5
T	4	3	2	2	3	3	3	4	5
A	5	4	3	3	3	4	3	3	4
A	6	5	4	3	4	3	4	4	4

Example run with AGACATTG and GAGTTA



Example run with AGACATTG and GAGTTA

	A	G	A	C	A	T	T	G	
G	0	1	2	3	4	5	6	7	8
A	1	1	1	2	3	4	5	6	7
G	2	1	2	1	2	3	4	5	6
A	3	2	1	2	2	3	4	5	5
G	4	3	2	2	3	3	3	4	5
A	5	4	3	3	3	4	3	3	4
G	6	5	4	3	4	3	4	4	4



Reading off the operations

- Follow the sequence and use each color of arrow to tell you what operation was performed.
- From the operations can derive an optimal alignment

A G A C A T T G
_ G A G _ T T A



Saving Space

- To compute the distance values we only need the last two rows (or columns)
 - $O(\min(m,n))$ space
- To compute the alignment/sequence of operations
 - seem to need to store all $O(mn)$ pointers/arrow colors
- Nifty divide and conquer variant that allows one to do this in $O(\min(m,n))$ space and retain $O(mn)$ time
 - In practice the algorithm is usually run on smaller chunks of a large string, e.g. m and n are lengths of genes so a few thousand characters
 - Researchers want all alignments that are close to optimal
 - Basic algorithm is run since the whole table of pointers (2 bits each) will fit in RAM
 - Ideas are neat, though



Saving space

- Alignment corresponds to a path through the table from lower right to upper left
 - Must pass through the middle column
- Recursively compute the entries for the middle column from the left
 - If we knew the cost of completing each then we could figure out where the path crossed
 - **Problem**
 - There are n possible strings to start from.
 - **Solution**
 - Recursively calculate the right half costs for each entry in this column using alignments starting at the **other** ends of the two input strings!
 - Can reuse the storage on the left when solving the right hand problem



Shortest paths with negative cost edges (Bellman-Ford)

- Dijkstra's algorithm failed with negative-cost edges
 - What can we do in this case?
 - Negative-cost cycles could result in shortest paths with length $-\infty$
- Suppose no negative-cost cycles in G
 - Shortest path from **s** to **t** has at most **n-1** edges
 - If not, there would be a repeated vertex which would create a cycle that could be removed since cycle can't have -ve cost



Shortest paths with negative cost edges (Bellman-Ford)

- We want to grow paths from **s** to **t** based on the # of edges in the path
- Let $\text{Cost}(\mathbf{s}, \mathbf{t}, \mathbf{i})$ = cost of minimum-length path from **s** to **t** using up to **i** hops.
 - $\text{Cost}(\mathbf{v}, \mathbf{t}, \mathbf{0}) = \begin{cases} \mathbf{0} & \text{if } \mathbf{v} = \mathbf{t} \\ \infty & \text{otherwise} \end{cases}$
 - $\text{Cost}(\mathbf{v}, \mathbf{t}, \mathbf{i}) = \min\{\text{Cost}(\mathbf{v}, \mathbf{t}, \mathbf{i}-1), \min_{(\mathbf{v}, \mathbf{w}) \in E} (\mathbf{c}_{\mathbf{vw}} + \text{Cost}(\mathbf{w}, \mathbf{t}, \mathbf{i}-1))\}$



Bellman-Ford

- Observe that the recursion for $\text{Cost}(\mathbf{s}, \mathbf{t}, \mathbf{i})$ doesn't change \mathbf{t}
 - Only store an entry for each \mathbf{v} and \mathbf{i}
 - Termed $\text{OPT}(\mathbf{v}, \mathbf{i})$ in the text
- Also observe that to compute $\text{OPT}(*, \mathbf{i})$ we only need $\text{OPT}(*, \mathbf{i}-1)$
 - Can store a current and previous copy in $O(\mathbf{n})$ space.



Bellman-Ford

ShortestPath(G, s, t)

for all $v \in V$

OPT[v] $\leftarrow -\infty$

OPT[t] $\leftarrow 0$

for $i=1$ to $n-1$ do

for all $v \in V$ do

O(mn) time

OPT'[v] $\leftarrow \min_{(v,w) \in E} (c_{vw} + \mathbf{OPT}[w])$

for all $v \in V$ do

OPT[v] $\leftarrow \min(\mathbf{OPT}'[v], \mathbf{OPT}[v])$

return **OPT**[s]



Negative cycles

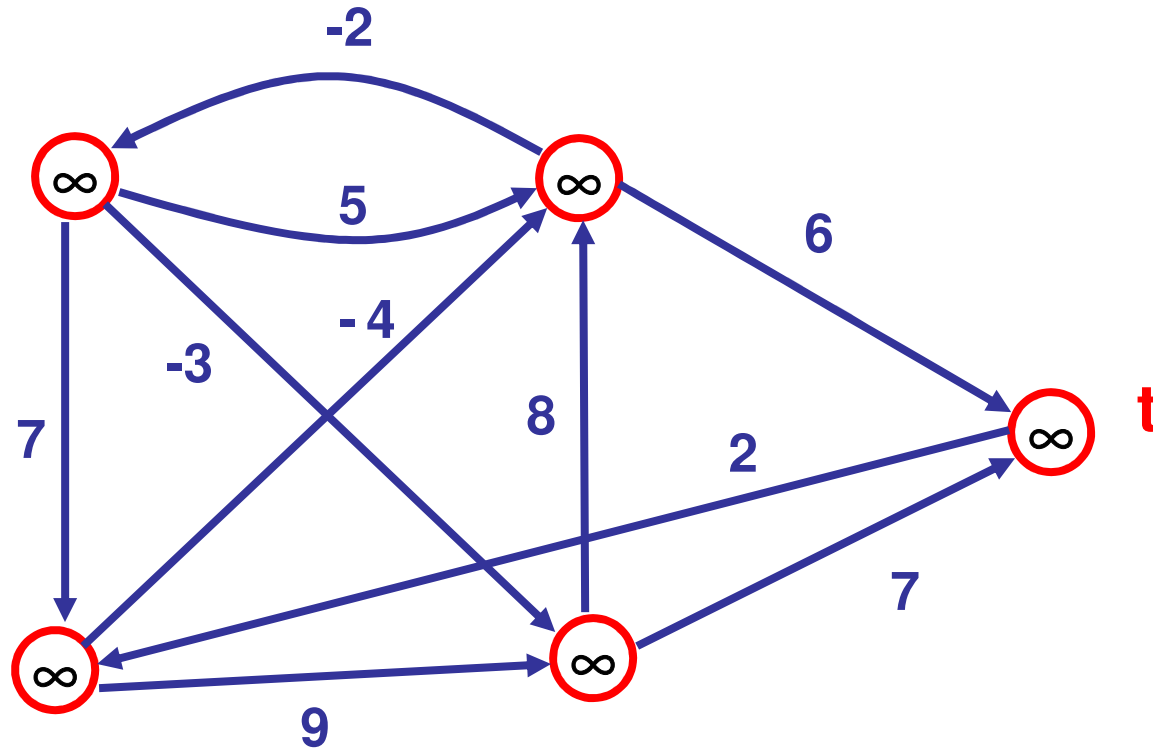
- **Claim:** There is a negative-cost cycle that can reach t iff for some vertex $v \in V$, $\text{Cost}(v,t,n) < \text{Cost}(v,t,n-1)$
- **Proof:**
 - We already know that if there aren't any then we only need paths of length up to $n-1$
 - For the other direction
 - The recurrence computes $\text{Cost}(v,t,i)$ correctly for **any** number of hops i
 - The recurrence reaches a fixed point if for every $v \in V$, $\text{Cost}(v,t,i) = \text{Cost}(v,t,i-1)$
 - A negative-cost cycle means that eventually some $\text{Cost}(v,t,i)$ gets smaller than any given bound
 - Can't have a -ve cost cycle if for every $v \in V$, $\text{Cost}(v,t,n) = \text{Cost}(v,t,n-1)$



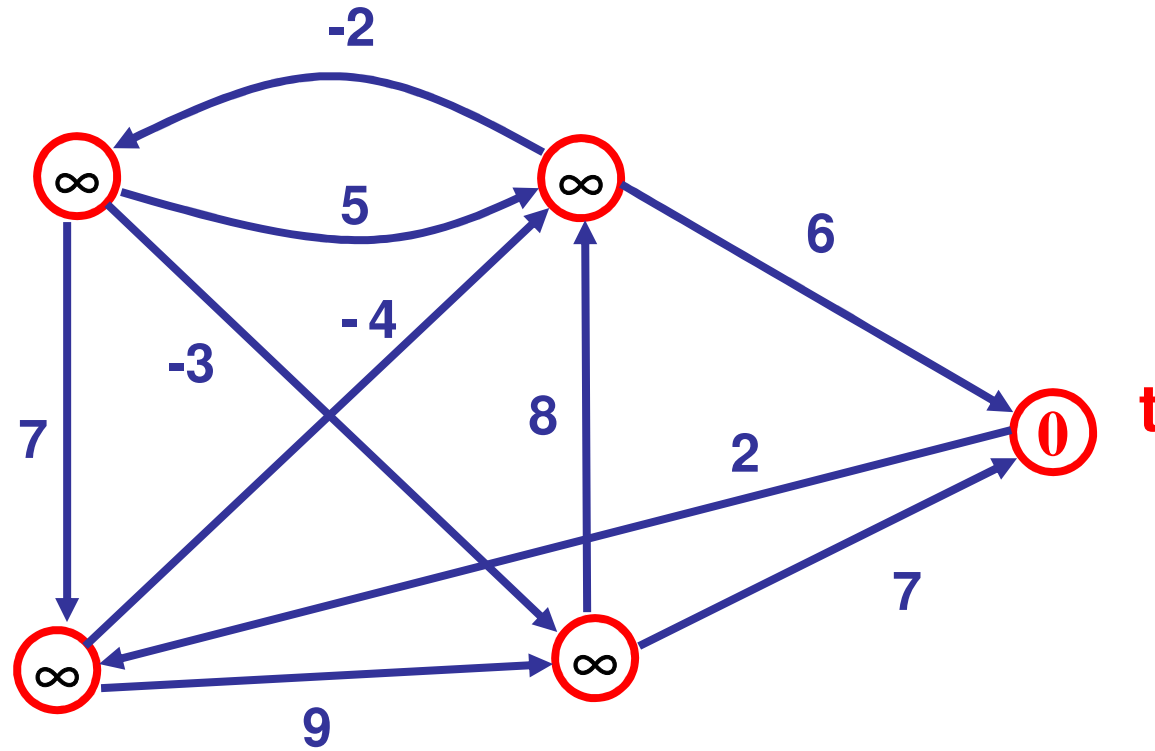
Last details

- Can run algorithm and stop early if the **OPT** and **OPT'** arrays are ever equal
 - Even better, one can update only neighbors **v** of vertices **w** with **OPT'[w] ≠ OPT[w]**
- Can store a **successor** pointer when we compute **OPT**
 - Homework assignment
- By running for step **n** we can find some vertex **v** on a negative cycle and use the successor pointers to find the cycle

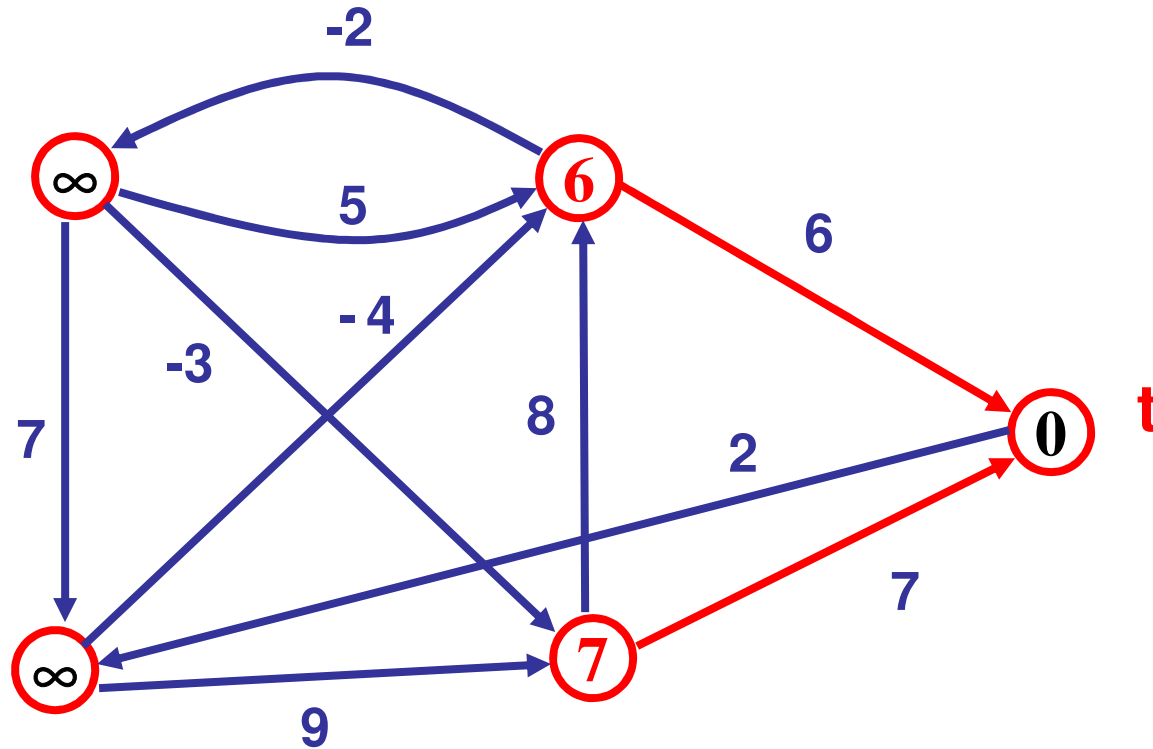
Bellman-Ford



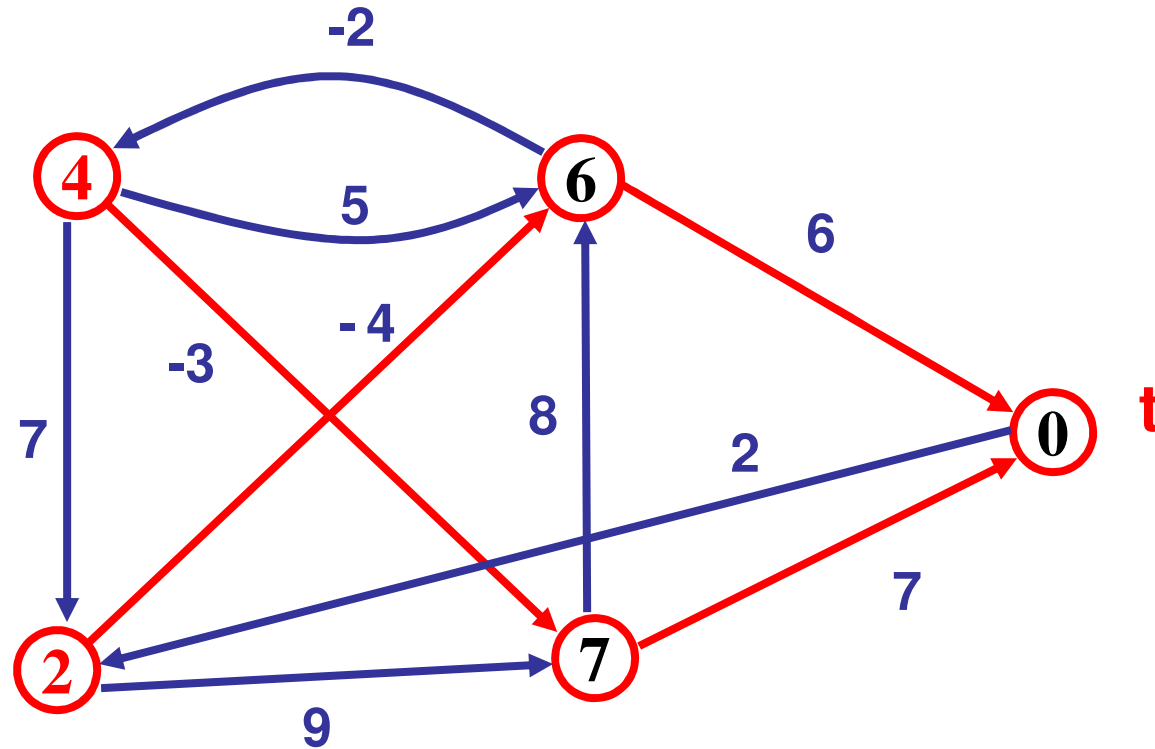
Bellman-Ford



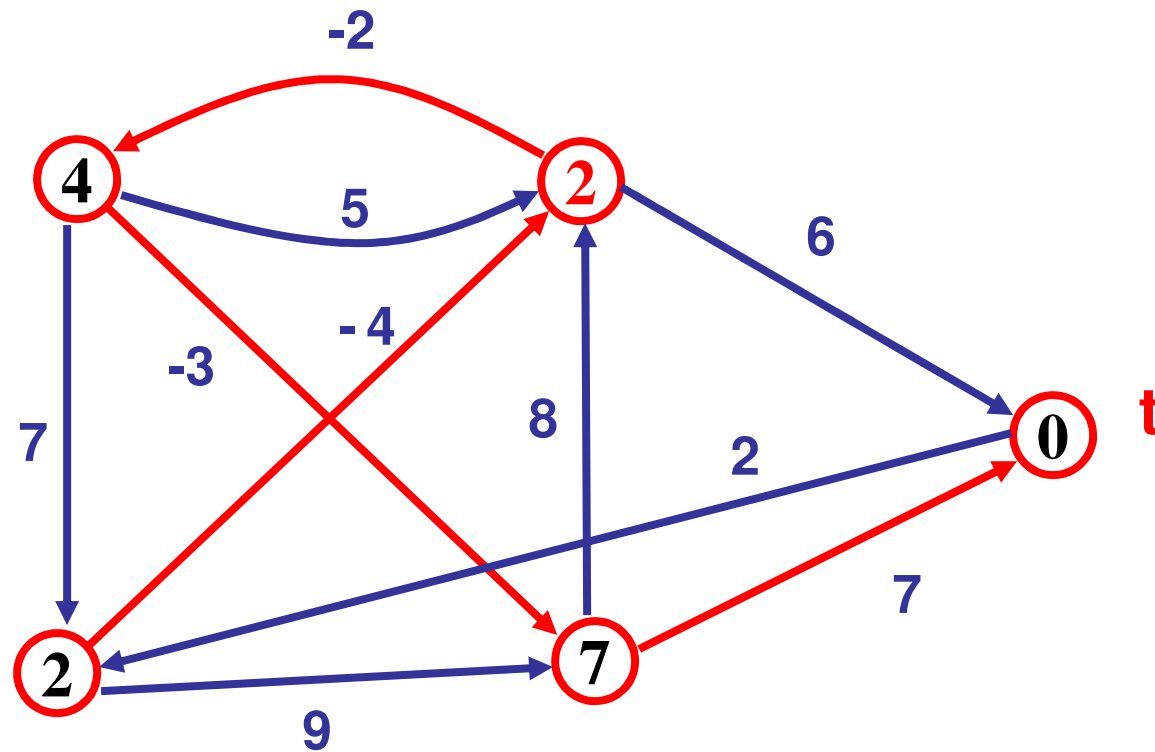
Bellman-Ford



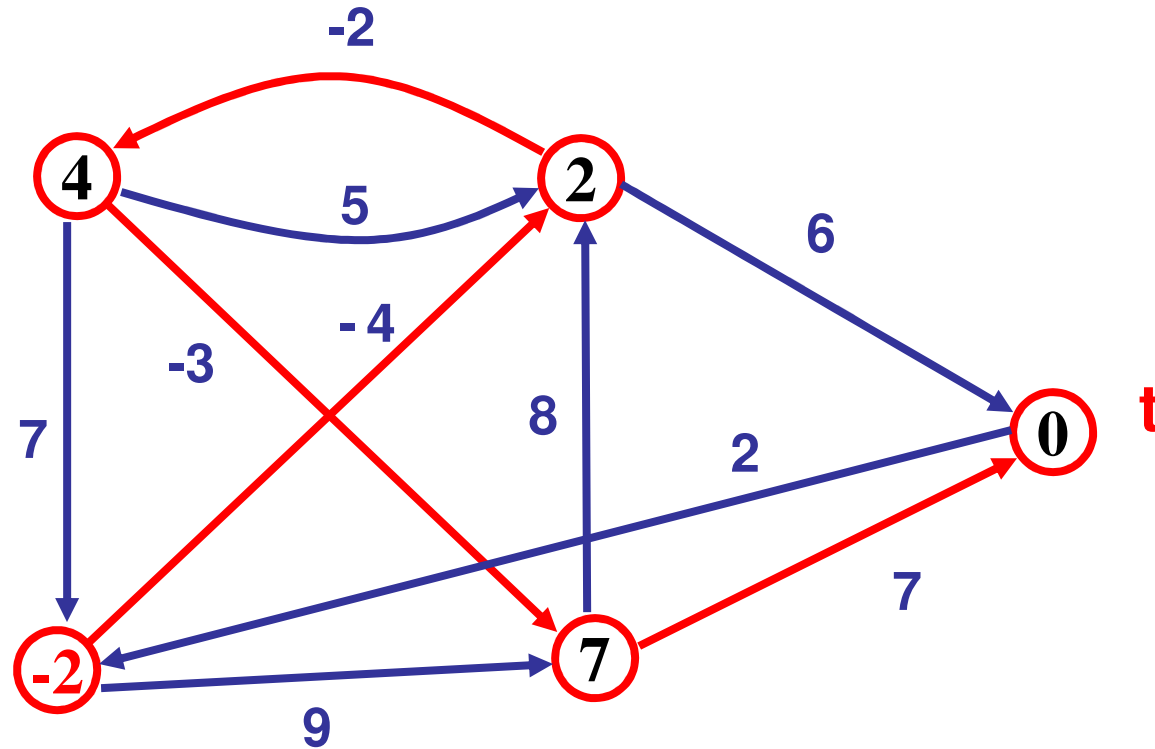
Bellman-Ford



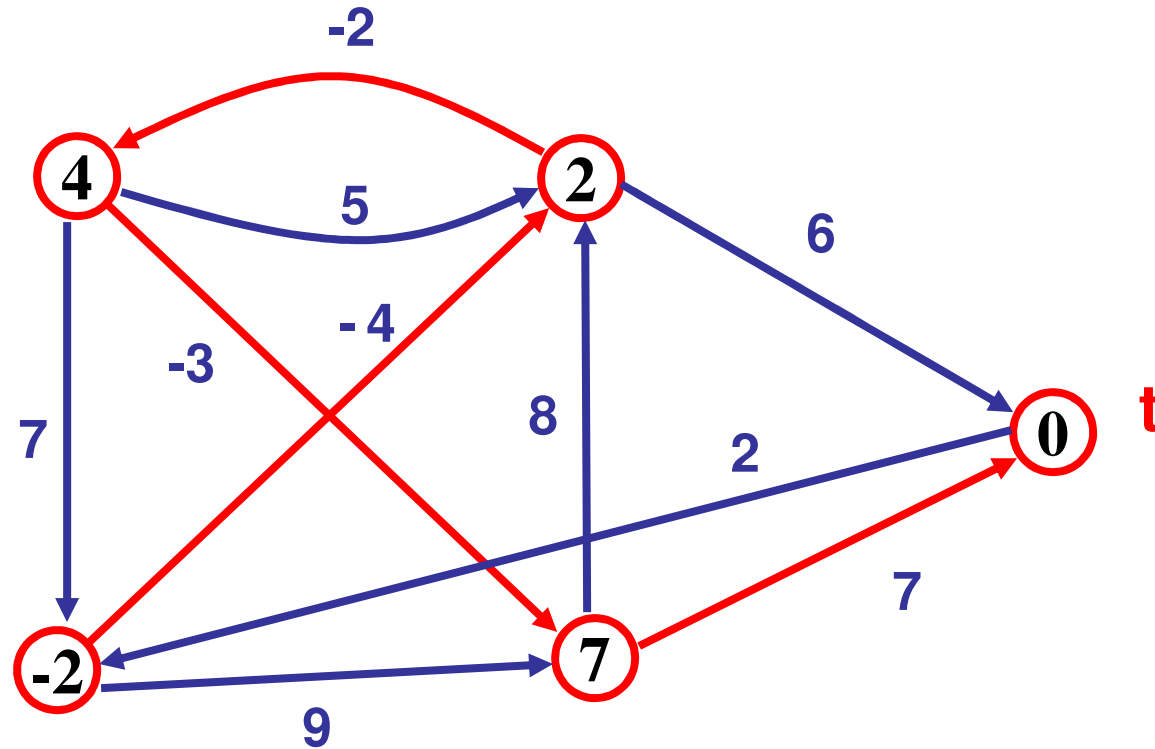
Bellman-Ford



Bellman-Ford



Bellman-Ford



Bellman-Ford with a DAG

Edges only go from lower to higher-numbered vertices

- Update distances in reverse order of topological sort
- Only one pass through vertices required
- $O(n+m)$ time

