# Chapter 6

# Dynamic Programming

CS 573: Algorithms, Fall 2014

September 11, 2014

## 6.1 Total Recall on DAGs

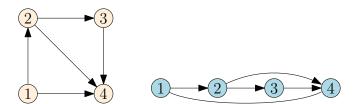
#### 6.1.0.1 DAGs

Definition 6.1.1. A DAG is a directed acyclic graph. That is a directed graph with no cycles.

## 6.1.1 Topological Ordering/Sorting

## 6.1.1.1 You should already know that...

Definition 6.1.2. A topological ordering/topological sorting of G = (V, E) is an ordering  $\prec$  on V such that if  $(u, v) \in E$  then  $u \prec v$ .



**Lemma 6.1.3.** A directed graph G can be topologically ordered iff it is a DAG.

**Lemma 6.1.4.** A DAG G = (V, E) with n vertices and m edges can be topologically sorted in O(n+m) time.

## 6.1.2 More things you should already know

#### 6.1.2.1 You should already know that...

**Lemma 6.1.5.** The strong connected components of a directed graph, can be computed in O(n+m) time.

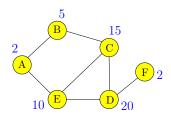
## 6.2 Maximum Weighted Independent Set in Trees

## 6.2.0.2 Maximum Weight Independent Set Problem

Problem 6.2.1 (Max Weight Independent Set).

**Input:** Graph G = (V, E) and weights  $w(v) \ge 0$  for each  $v \in V$ .

Goal: Find maximum weight independent set in G.



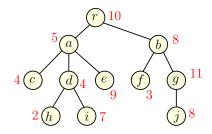
Maximum weight independent set in above graph:  $\{B, D\}$ .

## 6.2.0.3 Maximum Weight Independent Set in a Tree

Problem 6.2.2 (Max W. Independent Set in Tree).

**Input:** Tree T = (V, E) and weights  $w(v) \ge 0$  for each  $v \in V$ .

Goal: Find maximum weight independent set in T.



Maximum weight independent set in above tree: ??

## 6.2.0.4 Towards a Recursive Solution

- (A) For an arbitrary graph G:
  - (A) Number vertices as  $v_1, v_2, \ldots, v_n$
  - (B) Find recursively optimum solutions without  $v_n$  (recurse on  $\mathsf{G} v_n$ ) and with  $v_n$  (recurse on  $\mathsf{G} v_n N(v_n)$  & include  $v_n$ ).
  - (C) Saw that if graph G is arbitrary there was no good ordering that resulted in a small number of subproblems.
- (B) What about a tree?
- (C) Natural candidate for  $v_n$  is root r of T?

#### 6.2.1 Towards a Recursive Solution

#### 6.2.1.1 Maximum Weight Independent Set in a Tree

- (A) Natural candidate for  $v_n$  is root r of T?
- (B) Let  $\mathcal{O}$  be an optimum solution to the whole problem.

Case  $r \notin \mathcal{O}$ :  $\mathcal{O}$  contains optimum solution for each subtree hanging from a child of r.

Case  $r \in \mathcal{O}$ : None of children of r are in  $\mathcal{O}$ .

 $\mathcal{O}\setminus\{r\}$  contains an optimum solution for each subtree hanging at a grandchild of r.

(C) Subproblems? Subtrees of T hanging at nodes in T.

#### 6.2.1.2 A Recursive Solution

- (A) T(u): subtree of T hanging at node u.
- (B) OPT(u): max weighted independent set value in T(u).

(C) 
$$OPT(u) = \max \begin{cases} \sum_{v \text{ child of } u} OPT(v), \\ w(u) + \sum_{v \text{ grandchild of } u} OPT(v) \end{cases}$$

## 6.2.1.3 Iterative Algorithm

- (A) Compute OPT(u) bottom up.
- (B) To evaluate OPT(u) need to have computed values of all children and grandchildren of u
- (C) What is an ordering of nodes of a tree T to achieve above?
- (D) Post-order traversal of a tree.

## 6.2.1.4 Iterative Algorithm

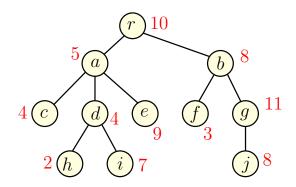
$$\begin{aligned} \textbf{MIS-Tree}(\mathsf{T}) : \\ & \text{Let } v_1, v_2, \dots, v_n \text{ be a post-order traversal of nodes of T} \\ & \textbf{for } i = 1 \text{ to } n \text{ do} \\ & M[v_i] = \max \left( \begin{array}{c} \sum_{v_j \text{ child of } v_i} M[v_j], \\ w(v_i) + \sum_{v_j \text{ grandchild of } v_i} M[v_j] \end{array} \right) \\ & \textbf{return } M[v_n] \text{ // Note: } v_n \text{ is the root of T} \end{aligned}$$

**Space:** O(n) to store the value at each node of T

#### Running time:

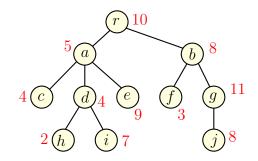
- (A) Naive bound:  $O(n^2)$  since each  $M[v_i]$  evaluation may take O(n) time and there are n evaluations.
- (B) Better bound: O(n). A value  $M[v_i]$  is accessed only by its parent and grand parent.

## **6.2.1.5** Example



#### 6.2.1.6 Dominating set

Definition 6.2.3. G = (V, E). The set  $X \subseteq V$  is a **dominating set**, if any vertex  $v \in V$  is either in X or is adjacent to a vertex in X.



Problem 6.2.4. Given weights on vertices, compute the *minimum* weight dominating set in G.

Dominating Set is NP-Hard!

## 6.3 DAGs and Dynamic Programming

#### 6.3.0.7 Recursion and DAGs

**Observation 6.3.1.** A: recursive algorithm for problem  $\Pi$ . For each instance I of  $\Pi$  there is an associated DAG G(I).

- (A) Create directed graph G(I) as follows...
- (B) For each sub-problem in the execution of A on I create a node.
- (C) If sub-problem v depends on or recursively calls sub-problem u add directed edge (u, v) to graph.
- (D) G(I) is a DAG. Why? If G(I) has a cycle then A will not terminate on I.

## 6.3.1 Iterative Algorithm for...

## 6.3.1.1 Dynamic Programming and DAGs

**Observation 6.3.2.** An iterative algorithm B obtained from a recursive algorithm A for a problem  $\Pi$  does the following:

For each instance I of  $\Pi$ , it computes a topological sort of G(I) and evaluates subproblems according to the topological ordering.

- (A) Sometimes the  $\overline{\mathsf{DAG}}$   $\mathsf{G}(I)$  can be obtained directly without thinking about the recursive algorithm A
- (B) In some cases (**not all**) the computation of an optimal solution reduces to a shortest/longest path in DAG G(I)
- (C) Topological sort based shortest/longest path computation is dynamic programming!

## 6.3.2 A quick reminder...

## 3.3.2.1 A Recursive Algorithm for weighted interval scheduling

Let  $O_i$  be value of an optimal schedule for the first i jobs.

```
 \begin{aligned} & \textbf{Schedule}(n): \\ & \textbf{if } n = 0 \textbf{ then return } 0 \\ & \textbf{if } n = 1 \textbf{ then return } w(v_1) \\ & O_{p(n)} \leftarrow \textbf{Schedule}(p(n)) \\ & O_{n-1} \leftarrow \textbf{Schedule}(n-1) \\ & \textbf{if } (O_{p(n)} + w(v_n) < O_{n-1}) \textbf{ then } \\ & O_n = O_{n-1} \\ & \textbf{else} \\ & O_n = O_{p(n)} + w(v_n) \\ & \textbf{return } O_n \end{aligned}
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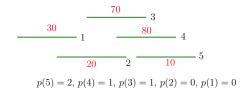
## 6.3.3 Weighted Interval Scheduling via...

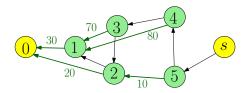
## 6.3.3.1 Longest Path in a DAG

Given intervals, create a DAG as follows:

- (A) Create one node for each interval, plus a dummy sink node 0 for interval 0, plus a dummy source node s.
- (B) For each interval i add edge (i, p(i)) of the length/weight of  $v_i$ .
- (C) Add an edge from s to n of length 0.
- (D) For each interval i add edge (i, i 1) of length 0.

### **6.3.3.2** Example





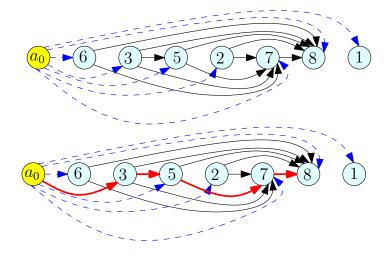
## 6.3.3.3 Relating Optimum Solution

- (A) Given interval problem instance I let G(I) denote the DAG constructed as described.
- (B) Claim 6.3.3. Optimum solution to weighted interval scheduling instance I is given by longest path from s to 0 in G(I).
- (C) Assuming claim is true,
  - (A) If I has n intervals,  $\overline{DAG}$  G(I) has n+2 nodes and O(n) edges. Creating G(I) takes  $O(n \log n)$  time: to find p(i) for each i. How?
  - (B) Longest path can be computed in O(n) time recall O(m+n) algorithm for shortest/longest paths in DAGs.

## 6.3.3.4 DAG for Longest Increasing Sequence

Given sequence  $a_1, a_2, \ldots, a_n$  create DAG as follows:

- (A) add sentinel  $a_0$  to sequence where  $a_0$  is less than smallest element in sequence
- (B) for each i there is a node  $v_i$
- (C) if i < j and  $a_i < a_j$  add an edge  $(v_i, v_j)$
- (D) find longest path from  $v_0$



## 6.4 Edit Distance and Sequence Alignment

## 6.4.0.5 Spell Checking Problem

- (A) Given a string "exponen" that is not in the dictionary, how should a spell checker suggest a *nearby* string?
- (B) What does nearness mean?
- (C) Question: Given two strings  $x_1x_2...x_n$  and  $y_1y_2...y_m$  what is a distance between them?
- (D) Edit Distance: minimum number of "edits" to transform x into y.

#### 6.4.0.6 Edit Distance

Definition 6.4.1. Edit distance between two words X and Y is the number of letter insertions, letter deletions and letter substitutions required to obtain Y from X.

Example 6.4.2. The edit distance between FOOD and MONEY is at most 4:

$$FOOD \rightarrow MOOD \rightarrow MONOD \rightarrow MONED \rightarrow MONEY$$

#### 6.4.0.7 Edit Distance: Alternate View

Alignment Place words one on top of the other, with gaps in the first word indicating insertions, and gaps in the second word indicating deletions.

Formally, an **alignment** is a set M of pairs (i, j) such that each index appears at most once, and there is no "crossing": i < i' and i is matched to j implies i' is matched to j' > j. In the above example, this is  $M = \{(1, 1), (2, 2), (3, 3), (4, 5)\}$ . Cost of an alignment is the number of mismatched columns plus number of unmatched indices in both strings.

#### 6.4.0.8 Edit Distance Problem

Problem Given two words, find the edit distance between them, i.e., an alignment of smallest cost.

## 6.4.0.9 Applications

- (A) Spell-checkers and Dictionaries
- (B) Unix diff
- (C) DNA sequence alignment ... but, we need a new metric

## 6.4.0.10 Similarity Metric

Definition 6.4.3. For two strings X and Y, the cost of alignment M is

- (A) [Gap penalty] For each gap in the alignment, we incur a cost  $\delta$ .
- (B) [Mismatch cost] For each pair p and q that have been matched in M, we incur cost  $\alpha_{pq}$ ; typically  $\alpha_{pp} = 0$ .

Edit distance is special case when  $\delta = \alpha_{pq} = 1$ .

## 6.4.0.11 An Example

Example 6.4.4.

Alternative:

Or a really stupid solution (delete string, insert other string):

 $Cost = 19\delta$ .

## 6.4.0.12 Sequence Alignment

Problem 6.4.5 (Sequence Alignment).

**Input:** Given two words X and Y, and gap penalty  $\delta$  and mismatch costs  $\alpha_{pq}$ .

Goal: Find alignment of minimum cost.

## 6.4.1 Edit distance

#### 6.4.1.1 Basic observation

Let  $X = \alpha x$  and  $Y = \beta y$ 

 $\alpha, \beta$ : strings.

x and y single characters.

Think about optimal edit distance between X and Y as alignment, and consider last column of alignment of the two strings:

$\alpha$	x
β	u

or

$\alpha$	x
$\beta y$	

or

$\alpha x$	
β	y

Observation 6.4.6. Prefixes must have optimal alignment!

#### 6.4.1.2 Problem Structure

**Observation 6.4.7.** Let  $X = x_1x_2 \cdots x_m$  and  $Y = y_1y_2 \cdots y_n$ . If (m, n) are not matched then either the mth position of X remains unmatched or the nth position of Y remains unmatched.

- (A) Case  $x_m$  and  $y_n$  are matched.
  - (A) Pay mismatch cost  $\alpha_{x_m y_n}$  plus cost of aligning strings  $x_1 \cdots x_{m-1}$  and  $y_1 \cdots y_{n-1}$
- (B) Case  $x_m$  is unmatched.
  - (A) Pay gap penalty plus cost of aligning  $x_1 \cdots x_{m-1}$  and  $y_1 \cdots y_n$
- (C) Case  $y_n$  is unmatched.
  - (A) Pay gap penalty plus cost of aligning  $x_1 \cdots x_m$  and  $y_1 \cdots y_{n-1}$

## 6.4.1.3 Subproblems and Recurrence

Optimal Costs Let Opt(i,j) be optimal cost of aligning  $x_1 \cdots x_i$  and  $y_1 \cdots y_i$ . Then

$$Opt(i, j) = \min \begin{cases} \alpha_{x_i y_j} + Opt(i - 1, j - 1), \\ \delta + Opt(i - 1, j), \\ \delta + Opt(i, j - 1) \end{cases}$$

Base Cases:  $\operatorname{Opt}(i,0) = \delta \cdot i$  and  $\operatorname{Opt}(0,j) = \delta \cdot j$ 

## 6.4.1.4 Dynamic Programming Solution

$$\begin{aligned} &\textbf{for all } i \ \textbf{do} \ M[i,0] = i\delta \\ &\textbf{for all } j \ \textbf{do} \ M[0,j] = j\delta \end{aligned} \\ &\textbf{for } i = 1 \ \textbf{to} \ m \ \textbf{do} \\ &\textbf{for } j = 1 \ \textbf{to} \ n \ \textbf{do} \\ &M[i,j] = \min \begin{cases} \alpha_{x_iy_j} + M[i-1,j-1], \\ \delta + M[i-1,j], \\ \delta + M[i,j-1] \end{cases} \end{aligned}$$

Analysis

- (A) Running time is O(mn).
- (B) Space used is O(mn).

## 6.4.1.5 Matrix and DAG of Computation

## 6.4.1.6 Sequence Alignment in Practice

- (A) Typically the DNA sequences that are aligned are about 10<sup>5</sup> letters long!
- (B) So about  $10^{10}$  operations and  $10^{10}$  bytes needed
- (C) The killer is the 10GB storage
- (D) Can we reduce space requirements?

## 6.4.1.7 Optimizing Space

(A) Recall

$$M(i,j) = \min \begin{cases} \alpha_{x_i y_j} + M(i-1, j-1), \\ \delta + M(i-1, j), \\ \delta + M(i, j-1) \end{cases}$$

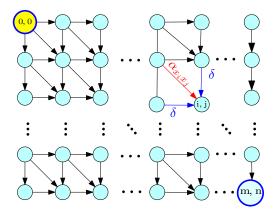


Figure 6.1: Iterative algorithm in previous slide computes values in row order. Optimal value is a shortest path from (0,0) to (m,n) in DAG.

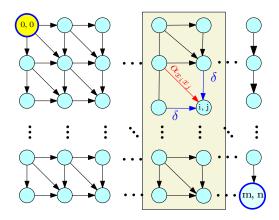


Figure 6.2: M(i, j) only depends on previous column values. Keep only two columns and compute in column order.

- (B) Entries in jth column only depend on (j-1)st column and earlier entries in jth column
- (C) Only store the current column and the previous column reusing space; N(i,0) stores M(i,j-1) and N(i,1) stores M(i,j)

## 6.4.1.8 Computing in column order to save space

#### 6.4.1.9 Space Efficient Algorithm

$$\begin{aligned} & \textbf{for all } i \ \textbf{ do } N[i,0] = i\delta \\ & \textbf{for } j = 1 \ \textbf{to } n \ \textbf{ do} \\ & N[0,1] = j\delta \ (* \ \textbf{corresponds to } M(0,j) \ *) \\ & \textbf{for } i = 1 \ \textbf{to } m \ \textbf{ do} \\ & N[i,1] = \min \begin{cases} \alpha_{x_iy_j} + N[i-1,0] \\ \delta + N[i-1,1] \\ \delta + N[i,0] \end{cases} \\ & \textbf{for } i = 1 \ \textbf{to } m \ \textbf{ do} \\ & \texttt{Copy } N[i,0] = N[i,1] \end{aligned}$$

Analysis Running time is O(mn) and space used is O(2m) = O(m)

## 6.4.1.10 Analyzing Space Efficiency

- (A) From the  $m \times n$  matrix M we can construct the actual alignment (exercise)
- (B) Matrix N computes cost of optimal alignment but no way to construct the actual alignment
- (C) Space efficient computation of alignment? More complicated algorithm see text book.

### 6.4.1.11 Takeaway Points

- (A) Dynamic programming is based on finding a recursive way to solve the problem. Need a recursion that generates a small number of subproblems.
- (B) Given a recursive algorithm there is a natural DAG associated with the subproblems that are generated for given instance; this is the dependency graph. An iterative algorithm simply evaluates the subproblems in some topological sort of this DAG.
- (C) The space required to evaluate the answer can be reduced in some cases by a careful examination of that dependency DAG of the subproblems and keeping only a subset of the DAG at any time.

## 6.5 All Pairs Shortest Paths

#### 6.5.0.12 Shortest Path Problems

Shortest Path Problems

**Input** A (undirected or directed) graph G = (V, E) with edge lengths (or costs). For edge e = (u, v),  $\ell(e) = \ell(u, v)$  is its length.

- (A) Given nodes s, t find shortest path from s to t.
- (B) Given node s find shortest path from s to all other nodes.
- (C) Find shortest paths for all pairs of nodes.

#### 6.5.0.13 Single-Source Shortest Paths

Single-Source Shortest Path Problems

**Input** A (undirected or directed) graph G = (V, E) with edge lengths. For edge  $e = (u, v), \ell(e) = \ell(u, v)$  is its length.

- (A) Given nodes s, t find shortest path from s to t.
- (B) Given node s find shortest path from s to all other nodes.

**Dijkstra's algorithm** for non-negative edge lengths. Running time:  $O((m+n)\log n)$  with heaps and  $O(m+n\log n)$  with advanced priority queues.

**Bellman-Ford algorithm** for arbitrary edge lengths. Running time: O(nm).

## 6.5.0.14 All-Pairs Shortest Paths

All-Pairs Shortest Path Problem

**Input** A (undirected or directed) graph G = (V, E) with edge lengths. For edge e = (u, v),  $\ell(e) = \ell(u, v)$  is its length.

(A) Find shortest paths for all pairs of nodes. Apply single-source algorithms n times, once for each vertex.

(A) Non-negative lengths.  $O(nm \log n)$  with heaps and  $O(nm+n^2 \log n)$  using advanced priority queues.

(B) Arbitrary edge lengths:  $O(n^2m)$ .  $\Theta(n^4)$  if  $m = \Omega(n^2)$ .

Can we do better?

#### 6.5.0.15 Shortest Paths and Recursion

- (A) Compute the shortest path distance from s to t recursively?
- (B) What are the smaller sub-problems?

**Lemma 6.5.1.** Let G be a directed graph with arbitrary edge lengths. If  $s = v_0 \rightarrow v_1 \rightarrow v_2 \rightarrow \ldots \rightarrow v_k$  is a shortest path from s to  $v_k$  then for  $1 \leq i < k$ :

(A)  $s = v_0 \rightarrow v_1 \rightarrow v_2 \rightarrow \ldots \rightarrow v_i$  is a shortest path from s to  $v_i$ 

Sub-problem idea: paths of fewer hops/edges

## 6.5.0.16 Hop-based Recur': Single-Source Shortest Paths

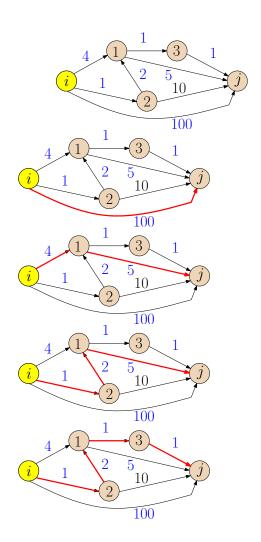
- (A) Single-source problem: fix source s.
- (B) OPT(v, k): shortest path dist. from s to v using at most k edges.
- (C) Note: dist(s, v) = OPT(v, n 1).
- (D) Recursion for OPT(v, k):

$$OPT(v,k) = \min \begin{cases} \min_{u \in V} (OPT(u,k-1) + c(u,v)). \\ OPT(v,k-1) \end{cases}$$

- (E) Base case: OPT(v, 1) = c(s, v) if  $(s, v) \in E$  otherwise  $\infty$
- (F) Leads to Bellman-Ford algorithm see text book.
- (G) OPT(v, k) values are also of independent interest: shortest paths with at most k hops.

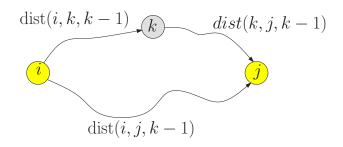
## 6.5.0.17 All-Pairs: Recursion on index of intermediate nodes

- (A) Number vertices arbitrarily as  $v_1, v_2, \ldots, v_n$
- (B) dist(i, j, k): shortest path distance between  $v_i$  and  $v_j$  among all paths in which the largest index of an *intermediate node* is at most k



$$dist(i, j, 0) = 100$$
  
 $dist(i, j, 1) = 9$   
 $dist(i, j, 2) = 8$   
 $dist(i, j, 3) = 5$ 

#### 6.5.0.18 All-Pairs: Recursion on index of intermediate nodes



$$dist(i, j, k) = \min \begin{cases} dist(i, j, k - 1) \\ dist(i, k, k - 1) + dist(k, j, k - 1) \end{cases}$$

Base case: dist(i, j, 0) = c(i, j) if  $(i, j) \in E$ , otherwise  $\infty$ 

Correctness: If  $i \to j$  shortest path goes through k then k occurs only once on the path — otherwise there is a negative length cycle.

## 6.5.1 Floyd-Warshall Algorithm

#### 6.5.1.1 for All-Pairs Shortest Paths

```
Check if G has a negative cycle // Bellman-Ford: O(mn) time if there is a negative cycle then return "Negative cycle" for i=1 to n do for j=1 to n do dist(i,j,0)=c(i,j) (* c(i,j)=\infty if (i,j)\notin E, 0 if i=j*) for k=1 to n do for i=1 to n do for j=1 to n do dist(i,j,k)=\min \begin{cases} dist(i,j,k-1), \\ dist(i,k,k-1)+dist(k,j,k-1) \end{cases}
```

Correctness: Recursion works under the assumption that all shortest paths are defined (no negative length cycle).

Running Time:  $\Theta(n^3)$ , Space:  $\Theta(n^3)$ .

## 6.5.2 Floyd-Warshall Algorithm

#### 6.5.2.1 for All-Pairs Shortest Paths

Do we need a separate algorithm to check if there is negative cycle?

```
\begin{array}{l} \mbox{for } i=1 \mbox{ to } n \mbox{ do} \\ \mbox{ for } j=1 \mbox{ to } n \mbox{ do} \\ \mbox{ } dist(i,j,0)=c(i,j) \mbox{ (* } c(i,j)=\infty \mbox{ if } (i,j)\notin E \mbox{, 0 if } i=j \mbox{ **}) \\ \mbox{for } i=1 \mbox{ to } n \mbox{ do} \\ \mbox{ for } i=1 \mbox{ to } n \mbox{ do} \\ \mbox{ } for \mbox{ } j=1 \mbox{ to } n \mbox{ do} \\ \mbox{ } dist(i,j,k)=min(dist(i,j,k-1),dist(i,k,k-1)+dist(k,j,k-1)) \\ \mbox{for } i=1 \mbox{ to } n \mbox{ do} \\ \mbox{ if } (dist(i,i,n)<0) \mbox{ then} \\ \mbox{ Output that there is a negative length cycle in G} \end{array}
```

Correctness: exercise

#### 6.5.2.2 Floyd-Warshall Algorithm: Finding the Paths

- (A) **Question:** Can we find the paths in addition to the distances?
- (B) Create a  $n \times n$  array Next that stores the next vertex on shortest path for each pair of vertices
- (C) With array Next, for any pair of given vertices i, j can compute a shortest path in O(n) time.

## 6.5.3 Floyd-Warshall Algorithm

### 6.5.3.1 Finding the Paths

```
\begin{array}{c} \mbox{for } i=1 \mbox{ to } n \mbox{ do} \\ \mbox{for } j=1 \mbox{ to } n \mbox{ do} \\ \mbox{} dist(i,j,0)=c(i,j) \mbox{ (* } c(i,j)=\infty \mbox{ if } (i,j) \mbox{ not edge, } 0 \mbox{ if } i=j \mbox{ *)} \\ \mbox{} Next(i,j)=-1 \\ \mbox{for } k=1 \mbox{ to } n \mbox{ do} \\ \mbox{for } i=1 \mbox{ to } n \mbox{ do} \\ \mbox{if } (dist(i,j,k-1)>dist(i,k,k-1)+dist(k,j,k-1)) \mbox{ then} \\ \mbox{} dist(i,j,k)=dist(i,k,k-1)+dist(k,j,k-1) \\ \mbox{} Next(i,j)=k \\ \mbox{for } i=1 \mbox{ to } n \mbox{ do} \\ \mbox{if } (dist(i,i,n)<0) \mbox{ then} \\ \mbox{Output that there is a negative length cycle in G} \end{array}
```

**Exercise:** Given Next array and any two vertices i, j describe an O(n) algorithm to find a i-j shortest path.

### 6.5.3.2 Summary of results on shortest paths

Single vertex		
No negative edges	Dijkstra	$O(n\log n + m)$
Edges cost might be negative But no negative cycles	Bellman Ford	O(nm)

All Pairs Shortest Paths

No negative edges	n * Dijkstra	$O(n^2 \log n + nm)$
No negative cycles	n * Bellman Ford	$O(n^2m) = O(n^4)$
No negative cycles	Floyd-Warshall	$O(n^3)$

## 6.6 Knapsack

#### 6.6.0.3 Knapsack Problem

Problem 6.6.1 (Knapsack).

**Input:** Given a Knapsack of capacity W lbs. and n objects with ith object having weight  $w_i$  and value  $v_i$ ; assume  $W, w_i, v_i$  are all positive integers.

Goal: Fill the Knapsack without exceeding weight limit while maximizing value.

(A) Basic problem that arises in many applications as a sub-problem.

#### 6.6.0.4 Knapsack Example

Item  $I_2$  $I_3$  $I_4$  $I_5$ Example 6.6.2. Value 6 18 22 28 2 5 7 Weight 1 6

If W = 11, the best is  $\{I_3, I_4\}$  giving value 40.

Special Case When  $v_i = w_i$ , the Knapsack problem is called the **Subset Sum Problem**.

## 6.6.0.5 Greedy Approach

- (A) Pick objects with greatest value
  - (A) Let W = 2,  $w_1 = w_2 = 1$ ,  $w_3 = 2$ ,  $v_1 = v_2 = 2$  and  $v_3 = 3$ ; greedy strategy will pick  $\{3\}$ , but the optimal is  $\{1, 2\}$
- (B) Pick objects with smallest weight
  - (A) Let  $W=2, w_1=1, w_2=2, v_1=1$  and  $v_2=3$ ; greedy strategy will pick  $\{1\}$ , but the optimal is  $\{2\}$
- (C) Pick objects with largest  $v_i/w_i$  ratio
  - (A) Let W=4,  $w_1=w_2=2$ ,  $w_3=3$ ,  $v_1=v_2=3$  and  $v_3=5$ ; greedy strategy will pick  $\{3\}$ , but the optimal is  $\{1,2\}$
  - (B) Can show that a slight modification always gives half the optimum profit: pick the better of the output of this algorithm and the largest value item. Also, the algorithms gives better approximations when all item weights are small when compared to W.

#### 6.6.0.6 Towards a Recursive Solution

First guess: Opt(i) is the optimum solution value for items  $1, \ldots, i$ .

**Observation 6.6.3.** Consider an optimal solution  $\mathcal{O}$  for  $1, \ldots, i$ 

Case item  $i \notin \mathcal{O}$   $\mathcal{O}$  is an optimal solution to items 1 to i-1

Case item  $i \in \mathcal{O}$  Then  $\mathcal{O} - \{i\}$  is an optimum solution for items 1 to n-1 in knapsack of capacity  $W - w_i$ .

Subproblems depend also on remaining capacity. Cannot write subproblem only in terms of  $Opt(1), \ldots, Opt(1)$ .

 $\operatorname{Opt}(i,w)$ : optimum profit for items 1 to i in knapsack of size w

Goal: compute Opt(n, W)

## 6.6.0.7 Dynamic Programming Solution

Definition 6.6.4. Let Opt(i, w) be the optimal way of picking items from 1 to i, with total weight not exceeding w.

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

## 6.6.0.8 An Iterative Algorithm

$$\begin{aligned} & \mathbf{for} \ \ w = 0 \ \ \mathbf{to} \ \ W \ \ \mathbf{do} \\ & M[0,w] = 0 \\ & \mathbf{for} \ \ i = 1 \ \ \mathbf{to} \ \ \mathbf{do} \\ & \mathbf{for} \ \ w = 1 \ \ \mathbf{to} \ \ W \ \ \mathbf{do} \\ & \mathbf{if} \ \ (w_i > w) \ \ \mathbf{then} \\ & M[i,w] = M[i-1,w] \\ & \mathbf{else} \\ & M[i,w] = \max(M[i-1,w],M[i-1,w-w_i] + v_i) \end{aligned}$$

Running Time

- (A) Time taken is O(nW)
- (B) Input has size  $O(n + \log W + \sum_{i=1}^{n} (\log v_i + \log w_i))$ ; so running time not polynomial but "pseudo-polynomial"!

## 6.6.0.9 Knapsack Algorithm and Polynomial time

- (A) Input size for Knapsack:  $O(n) + \log W + \sum_{i=1}^{n} (\log w_i + \log v_i)$ .
- (B) Running time of dynamic programming algorithm: O(nW).
- (C) Not a polynomial time algorithm.
- (D) Example:  $W = 2^n$  and  $w_i, v_i \in [1..2^n]$ . Input size is  $O(n^2)$ , running time is  $O(n2^n)$  arithmetic/comparisons.
- (E) Algorithm is called a *pseudo-polynomial* time algorithm because running time is polynomial if *numbers* in input are of size polynomial in the *combinatorial size* of problem.
- (F) Knapsack is NP-Hard if numbers are not polynomial in n.

## 6.7 Traveling Salesman Problem

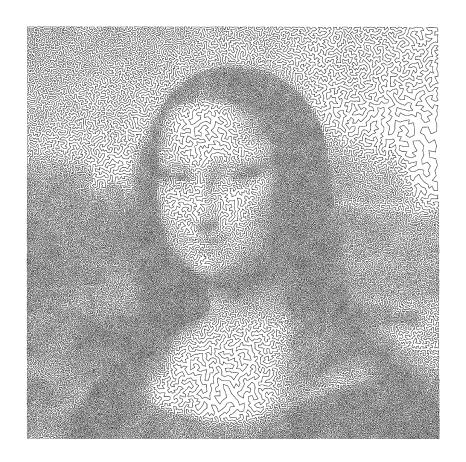
## 6.7.0.10 Traveling Salesman Problem

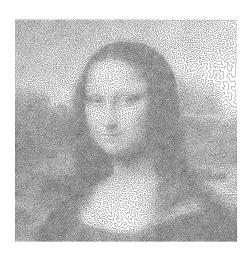
Problem 6.7.1 (**TSP**).

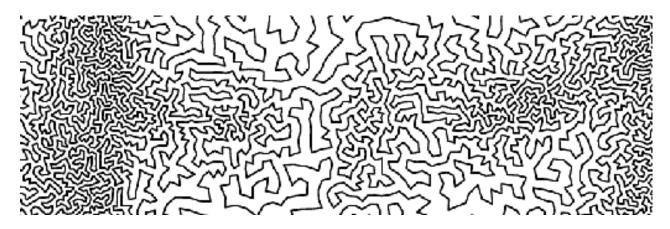
**Input:** A graph G = (V, E) with non-negative edge costs/lengths. Cost c(e) for each edge  $e \in E$ . **Goal:** Find a tour of minimum cost that visits each node.

(A) No polynomial time algorithm known. Problem is **NP-Hard**.

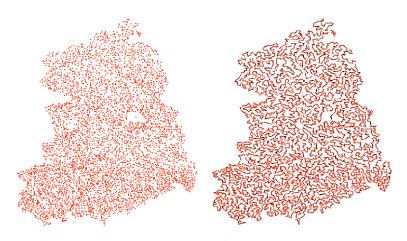
## 6.7.0.11 Drawings using TSP







6.7.0.12 Example: optimal tour for cities of a country (which one?)



### 6.7.0.13 An Exponential Time Algorithm

- (A) How many different tours are there? n!
- (B) Stirling's formula:  $n! \simeq \sqrt{n}(n/e)^n$  which is  $\Theta(2^{cn\log n})$  for some constant c>1
- (C) Can we do better? Can we get a  $2^{O(n)}$  time algorithm?

#### 6.7.0.14 Towards a Recursive Solution

- (A) Order vertices as  $v_1, v_2, \ldots, v_n$
- (B) OPT(S): optimum **TSP** tour for the vertices  $S \subseteq V$  in the graph restricted to S. Want OPT(V). Can we compute OPT(S) recursively?
- (A) Say  $v \in S$ . What are the two neighbors of v in optimum tour in S?
- (B) If u, w are neighbors of v in an optimum tour of S then removing v gives an optimum path from u to w visiting all nodes in  $S \{v\}$ .

Path from u to w is not a recursive subproblem! Need to find a more general problem to allow recursion.

#### 6.7.0.15 A More General Problem: TSP Path

(A) Problem 6.7.2 (**TSP Path**).

**Input:** A graph G = (V, E) with non-negative edge costs/lengths(c(e) for edge e) and two nodes s, t.

**Goal:** Find a path from s to t of minimum cost that visits each node exactly once.

- (B) Can solve TSP using above. Do you see how?
- (C) Recursion for optimum TSP Path problem:
  - (A) OPT(u, v, S): optimum TSP Path from u to v in the graph restricted to S (here  $u, v \in S$ ).

## 6.7.1 A More General Problem: TSP Path

#### 6.7.1.1 Continued...

- (A) What is the next node in the optimum path from u to v?
- (B) Suppose it is w. Then what is OPT(u, v, S)?
- (C)  $OPT(u, v, S) = c(u, w) + OPT(w, v, S \{u\})$
- (D) We do not know w! So try all possibilities for w.

#### 6.7.1.2 A Recursive Solution

- (A)  $OPT(u, v, S) = \min_{w \in S, w \neq u, v} (c(u, w) + OPT(w, v, S \{u\}))$
- (B) What are the subproblems for the original problem OPT(s,t,V)? OPT(u,v,S) for  $u,v \in S, S \subseteq V$ .
- (C) How many subproblems?
  - (A) number of distinct subsets S of V is at most  $2^n$
  - (B) number of pairs of nodes in a set S is at most  $n^2$
  - (C) hence number of subproblems is  $O(n^22^n)$
- (D) **Exercise:** Show that one can compute **TSP** using above dynamic program in  $O(n^3 2^n)$  time and  $O(n^2 2^n)$  space.

Disadvantage of dynamic programming solution: memory!

## 6.7.1.3 Dynamic Programming: Postscript

Dynamic Programming = Smart Recursion + Memoization

- (A) How to come up with the recursion?
- (B) How to recognize that dynamic programming may apply?

#### 6.7.1.4 Some Tips

- (A) Problems where there is a *natural* linear ordering: sequences, paths, intervals, DAGs etc. Recursion based on ordering (left to right or right to left or topological sort) usually works.
- (B) Problems involving trees: recursion based on subtrees.
- (C) More generally:
  - (A) Problem admits a natural recursive divide and conquer
  - (B) If optimal solution for whole problem can be simply composed from optimal solution for each separate pieces then plain divide and conquer works directly
  - (C) If optimal solution depends on all pieces then can apply dynamic programming if *inter-face/interaction* between pieces is *limited*. Augment recursion to not simply find an optimum solution but also an optimum solution for each possible way to interact with the other pieces.

#### 6.7.1.5 Examples

- (A) Longest Increasing Subsequence: break sequence in the middle say. What is the interaction between the two pieces in a solution?
- (B) Sequence Alignment: break both sequences in two pieces each. What is the interaction between the two sets of pieces?

- (C) Independent Set in a Tree: break tree at root into subtrees. What is the interaction between the sutrees?
- (D) Independent Set in an graph: break graph into two graphs. What is the interaction? Very high!
- (E) Knapsack: Split items into two sets of half each. What is the interaction?