

MIT Dynamic Programming (DP)

* DP \approx "careful brute force."

DP \approx subproblems + "reuse"

Fibonacci numbers

$$\begin{cases} F_1 = F_2 = 1 \\ F_n = F_{n-1} + F_{n-2} \end{cases}$$

Goal: compute F_n

Naive Recursive algorithm: $\text{fib}()$:

Exponential Time

$$T(n) = T(n-1) + T(n-2) + \Theta(1)$$

if $n \leq 2$: $f = 1$

else $f = \text{fib}(n-1) + \text{fib}(n-2)$

return f

Memoized Dynamic Programming algorithm

memo = { }

$\text{fib}(n)$:

if n in memo : return memo[n]

if $n \leq 2$: $f = 1$

else : $f = \text{fib}(n-1) + \text{fib}(n-2)$

memo[n] = f

return f

$\text{fib}(k)$ only recurses the first time it's called, $\forall k$

→ memoized calls cost $\Theta(1)$

→ #non-memoized calls is n :

$\text{fib}(1), \text{fib}(2), \dots, \text{fib}(n)$

— non-recursive work per call = $\Theta(1)$

⇒ time = $\Theta(n)$

In General DP is recursion + memoization.

→ memoize (remember)
& re-use solutions to subproblems that help solve the problem

⇒ time # subproblem. n $\frac{\text{time / subproblem}}{O(1)}$

Bottom-up DP algorithm.

fib = { }

for k in range(~~n~~)

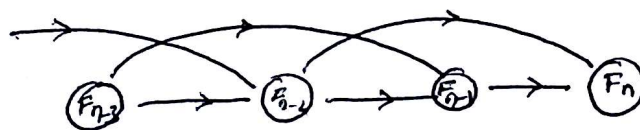
[if $k \leq 2$ $f = 1$

[else $f = \text{fib}[k-1] + \text{fib}[k-2]$

fib[k] = f

return fib[n]

- exact same computation as memoization
- topological sort of subproblem dependency DAG.



- Can often save space.

Shortest path algorithm

Bellman-Ford algorithm.

Bellman-Ford (G, w, s)

Initialize - Single-source (G, s)

for $i \leftarrow 1$ to $|V[G]| - 1$

do for each edge $(u, v) \in E[G]$

do Relax(u, v, w)

for each edge $(u, v) \in E[G]$

do if $d[v] > d[u] + w(u, v)$

then return False

return True

Relax(u, v, w)

if $d[v] > d[u] + w(u, v)$

then $d[v] \leftarrow d[u] + w(u, v)$

$\pi[v] \leftarrow u$

6.23. If $i > 0$ then

$$OPT(i, v) = \min(OPT(i-1, v), \min_{w \in V} (OPT(i-1, w) + C_w w))$$

Shortest-Path (G, s, t)

n = number of nodes in G

Array $M[0 \dots n-1, V]$

define $M[0, t] = 0$ and $M[0, v] = \infty$ for all $v \in V$

For $i = 1 \dots n-1$.

For $v \in V$ in any order

compute $M[i, v]$ using 6.23

Endfor

Endfor

Return $M[n-1, s]$

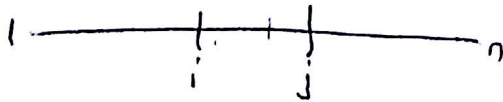
RNA Secondary structure: DP over intervals

value of $OPT(i, j) \Rightarrow$ Secondary structure that maximizes number of pairs when we restrict attention to the subsequence between i and j

want $OPT(1, n)$

$$i \geq 1 \quad j \leq n$$

$$i \leq j$$



Recurrence

Base case $OPT(i, j) = 0$ if $i \geq j - 4$.

$$OPT(i, j) = \max \left[OPT(i, j-1), \right.$$

$$\left. \max(1 + OPT(i, t-1) + OPT(t+1, j-1)) \right]$$

if char in pos t and j can pair

Weighted Interval Scheduling

$$OPT(j) = \max(v_j + OPT(P(j)), OPT(j-1))$$

Segmented Least Squares

$$OPT(j) = \min_{1 \leq i \leq j} (e_{ij} + C + OPT(i-1))$$

Knapsack

If $w \leq w_i$ then $OPT(i, w) = OPT(i-1, w)$ otherwise

$$OPT(i, w) = \max(OPT(i-1, w), w_i + OPT(i-1, w-w_i))$$