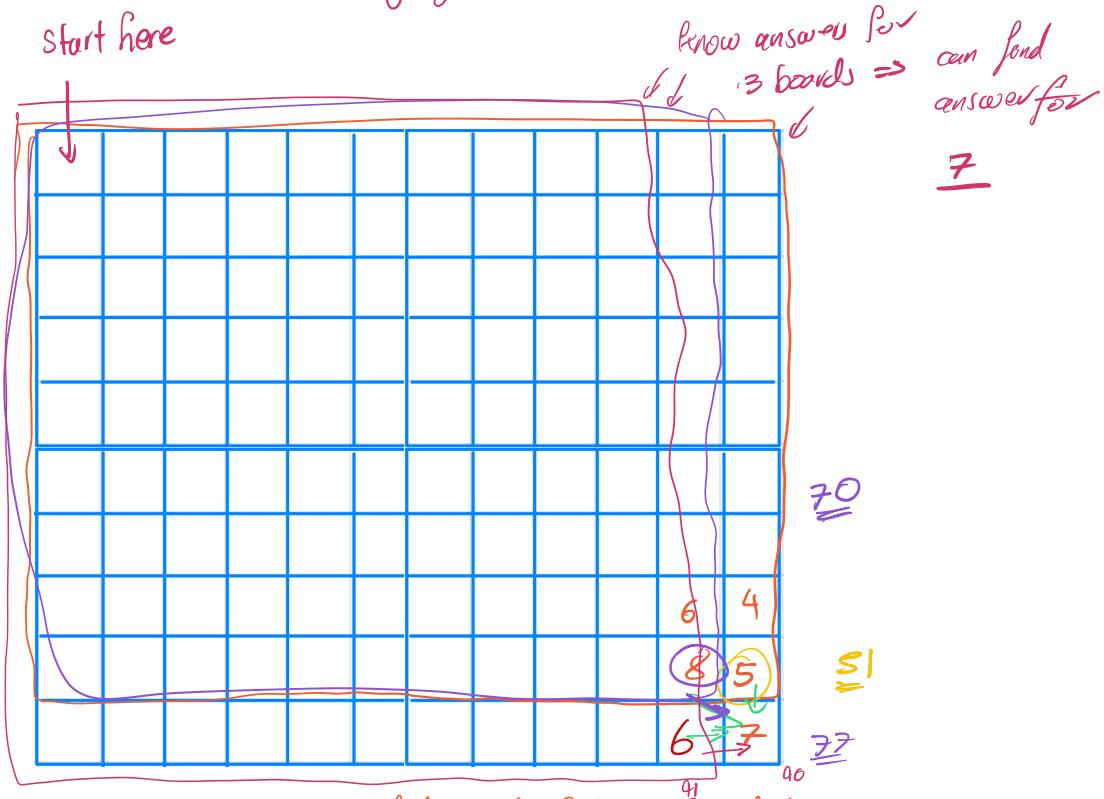
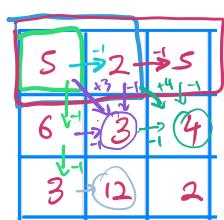


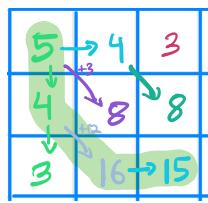
Total money you can make?



Dynamic Programming → sub solutions to find main solution



Game Board



Dynamic Prog Table
→ track highest money for any score

Now, Backtrack & find best path
→ recursive

Memory
how much additional memory to store numbers?

Time

max money for each square

Max money for cost.

Computational Complexity \rightarrow abstraction to quantify time algorithm takes
 \downarrow
 \rightarrow more complex \rightarrow \uparrow time complexity
algorithm

✓ quantifies how long it takes

↓ form of equation → e.g. linear, quadratic etc
D is "D" ↗

Complexity:

Big "O" notation

$O(k)$ linear coefficient

algorithm $\Rightarrow O(r \cdot c)$
rows \times columns
... linear to $r \cdot c$

Time complexity is linear to r.c

Memory Complexity (space)

Memory Components → depend on algorithm
of Memory requirements

Time & Memory seq

$$\text{Time of insertion} \rightarrow O(r.c) \rightarrow O(n^2)$$

→ all data stored $\Rightarrow O(n)$
 → sum kept $\Rightarrow O(1) \rightarrow O(n)$

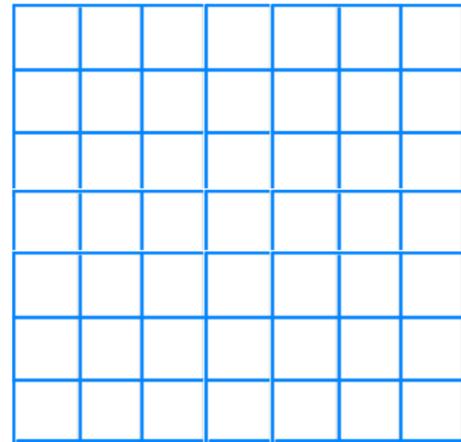
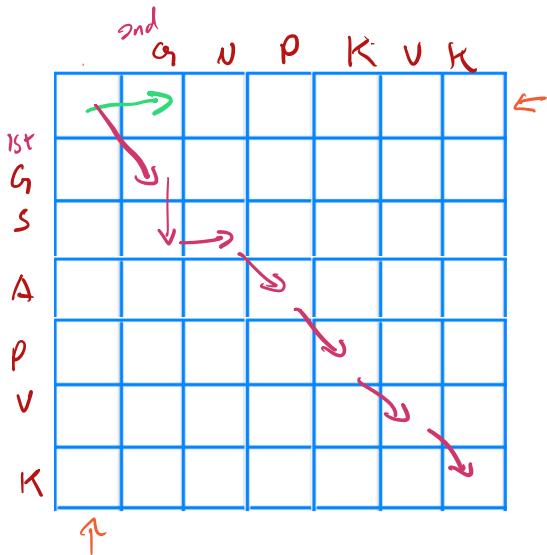
\rightarrow rows deleted, columns kept $\Rightarrow O(n) \rightarrow$ On

→ rows deleted
② columns kept, rows deleted $\Rightarrow O(r) \rightarrow$ On

→ columns kept rows deleted

let $r = n$, $c = n$

Game Board for Sequence Alignment



empty columns
allow gaps

$\rightarrow \downarrow = \text{gap}$
 $\rightarrow \swarrow = \text{match/mismatch}$

$Q S - A P V K$
 $Q - N P K V K$

$\downarrow \Rightarrow \text{gap in second}$
 $\rightarrow \Rightarrow \text{gap in first}$

	Q	N	P	K	V	K	
Q	1	0	0	0	0	0	
S	0	0	0	0	0	0	
A	0	0	0	0	0	0	
P	0	0	1	0	0	0	
V	0	0	0	0	1	0	
K	0	0	0	1	0	1	

match there $\Rightarrow +1$

all alignments modeled as paths through board

match 1 gap -1

diagonal \Rightarrow match/mismatch $\rightarrow +1$ or $+0$
& design board so you can have max score

Gap move \rightarrow ignore game board
 \rightarrow pay gap penalty
included for DP Table

	a	n	p	k	v	k
q	0	0	0	0	0	0
s	0	0	0	0	0	0
a	0	0	0	0	0	0
p	0	0	0	1	0	0
v	0	0	0	0	0	1
k	0	0	0	0	1	0

zero :- these numbers will never get picked up

	a	n	p	k	v	k
q	0	-1	-2	-3	-4	-5
s	-1	1				
a	-2					
p	-3					
v	-4					
k	-5					

do not need game board for seq align
 ↳ just go through DP table
 ↳ if match use match score

if given substitution matrix
 use that to decide scores (game board)

$$S_{i,j} = \max \left\{ \begin{array}{l} S_{i-1, j-1} + s(x_i, y_j) \\ S_{i-1, j} + g \\ S_{i, j-1} + g \end{array} \right.$$

↑ substitution matrix (game board)
 ↓ store in var, find max

gameboard needs gap penalty

- fill in 1st row & col of DP Table
- begin for loop in $DP[1,1]$ not $DP[0,0]$

semi-global alignment (free end gaps)

shift sequences or
 \Rightarrow DONT penalize gaps in last row & last column
 \rightarrow let algorithm run to end
 \rightarrow identify max val in last row & col \Rightarrow max align score
 \rightarrow back-track from there

local alignment

no penalties for both seq
 endgaps \rightarrow outer rows, columns
 \rightarrow to align shorter sequence, get to start for free, reach end
 \rightarrow local align \Rightarrow start anywhere, stop anywhere

change

$$S_{i,j} = \max \begin{cases} S_{i-1,j-1} + s(x_i, y_j) \\ S_{i-1,j} + g \\ S_{i,j-1} + g \\ 0 \end{cases}$$

↑
new starting points
↓
more can start anywhere on board

\rightarrow fill board
 \rightarrow find highest value in entire board
 \Rightarrow That is maximum local align score
 \Rightarrow back-tracking stops at zeros

multiple LB

\rightarrow run algorithm
 \rightarrow set max path = 0 \Rightarrow no longer allowed
 \Rightarrow find next max \Rightarrow select

set a threshold for maxima

Game Board (Scoring Matrix/Dictionary)

3	7	8	8	7	6	8
5	3	7	4	2	7	3
6	7	1	8	8	6	2
5	7	7	5	3	5	7
4	1	4	1	10	7	5
6	3	10	2	3	7	5
10	3	1	8	6	7	7

Start

Row 1

Col 1

Dynamic Programming Table

3	2	1	0	-1	-2	-3
↓	↓	↑	6			
2		↓				
1		↓				
0		↓				
-1		↓				
-2		↓				
-3						

Row 1 & Col 1 are empty for sequences

Code:

```
for i in range(0, r):
    for j in range(0, c):
```

	G	C	A	G	T	
T	0	0	0	0	0	0
G	0	-2	-2	-2	-2	2
T	0	-2	-2	-2	-2	2
C	0	-2	2	-2	-2	-2
G	0	2	-2	-2	2	-2

	A	C	C	G	
A	0	-2	-4	-6	-8
C	-2	2	0	-2	-4
A	-4	0	2	-2	-4
G	-6	-2	2	-2	-4
	-8	-4	0	2	-2

match = 2

mismatch = 0

gap = -2

	G	C	A	G	T	
T	0	0	0	0	0	0
G	0					
T	0					
C	0					
G	0					

	A	C	C	G	
A	0	-2	-4	-6	-8
C	-2	2	0	-2	-4
A	-4	0	2	-2	-4
G	-6	-2	2	-2	-4
	-8	-4	0	2	-2

match = 2
mismatch = 0
gap = -2

score = 2
A C C G
A C A G

	G	C	A	G	T	
T	0	-1	-2	-3	-4	-5
G	-1	-2	-3	-4	-5	-6
T	-2	1	0	-1	-2	-3
C	-3					
C	-4	-1	2	1	0	-1
G	-5	-2	1	0	3	2

match = 2
mismatch = -2
gap = -1

- G1 - C A G1 T

	T	G	T	C	G	
T	0	-1	-2	-3	-4	-5
G	-1	-2	1	0	-1	-2
C	-2	-3	0	-1	2	1
A	-3	-4	-1	-2	1	0
G	-4	-5	-2	-3	0	3
T	-5	-2	-3	0	-1	2

match = 2
mismatch = -2

T A T C G1

G C

- G1 - C A G1 T
T A T C G1

match = 1
mismatch = -1

	G	C	A	G	T	
T	0	0	0	0	0	
G	0	-1	-1	-2	0	-1
T	0	-1	0	-2	-2	1
C	0	-1	0	-1	-3	-1
G	0	-1	1	-1	0	-2

match = 1
mismatch = -1

	T	G	T	C	G	
T	0	-2	-4	-6	-8	-10
G	-2	-1	1	-3	-5	-7
C	-4	-3	-2	-2	-5	-4
A	-6	-5	-4	-3	-3	-3
G	-8	-7	-4	-5	-4	-2
T	-10	-7	-6	-3	-5	-4

match = 1
mismatch = -1

- G C A G T
T A T C G

G1 C A G T - - -
- - - T G T C G
G1 C A G T - - -
- - - T G T C G
- - - G C A G T
T G T C G - - -

	P	I	W	A	L	I	
A	0	-6	-12	-18	-24	-30	-36
L	-6	-1	7				
I	-12						
W	-18						
T	-24						
T	-30						

	P	I	W	A	L	I	
A	0	-6	-12	-18	-24	-30	-36
L	-6	-1	7				
I	-12						
W	-18						
T	-24						
T	-30						

	G	C	A	G	T	
T	0	0	0	0	0	
G	0	-1	-1	-2	0	-1
T	0	-1	0	-2	-2	1
C	0	-1	0	-1	-3	-1
G	0	-1	1	-1	0	-2

match = 1
mismatch = -1

G1 C A G T - - -
- - - T G T C G
G1 C A G T - - -
- - - T G T C G
- - - G C A G T
T G T C G - - -

	P	I	W	A	L	I	
A	0	-6	-12	-18	-24	-30	-36
L	-6	-1	7				
I	-12						
W	-18						
T	-24						
T	-30						

Ask Dr. Saccan

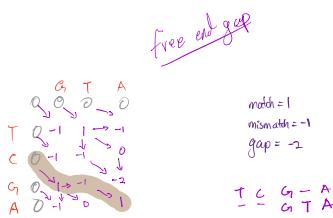
← are the three alignments valid?

gap = -2 match = 1
misMatch = -1

	G	C	A	G	T	
T	0	0	0	0	0	1
G	0	1	0	0	1	0
T	0	0	0	0	0	2
C	0	0	1	0	0	0
G	0	1	0	0	1	0

$S_{ij} = \max \begin{cases} S_{i,j-1} + s(x_i, y_j) \\ S_{i-1,j} + g \\ S_{i-1,j-1} + g \\ 0 \end{cases}$

$\sim \text{---TGT CG}$
 $\sim \text{---GCA GT ---}$



Sum of all Pairs alignment

a MWGTFNW

b MR---W

c M-GTF-N

$$S_{\text{all pair}} = S_{ab} + S_{ac} + S_{bc}$$

(= -23)

a	M	W	G	F	N	W	
b	M	R	—	—	—	W	(-2)
	5	-3	-5	-5	-5	11	
a	M	W	G	F	N	W	
c	M	—	G	F	—	N	
	5	-5	6	6	-5	-4	(-3)
b	M	R	—	—	—	W	
c	M	—	G	F	—	N	
	5	-5	-5	-5	-5	-4	(-24)

Entropy	ACA	$f_{1,A} = \frac{2}{4} = 0.5$
	AGT	$f_{1,T} = \frac{1}{4} = 0.25$
	TAT	$f_{1,C} = \frac{1}{4} = 0.25$
	GTT	$f_{1,G} = 0$

number of unrooted trees