

MEKKITGYTT  
NMFFVSANPW  
MLNELQQ  
MNAHTAFL

```
'0': {'1': -9, '2': -9, '3': -4}, sum = -22
'1': {'0': -9, '2': -9, '3': -4}, sum = -22
'2': {'0': -9, '1': -9, '3': -4}, sum = -22
'3': {'0': -4, '1': -4, '2': -4}, sum = -12
```

```
Centre = seqs[3]: {'0': -4, '1': -4, '2': -4}
Orders: ['0', '1', '2']
```

```
global_align(seqs[0], centre, match, mismatch,
gap_open)
```

```
('MEKKITGYTT', 'M-NAHT-AFL', -4)
Centre = M-NAHT-AFL
```

```
global_align(seqs[1], centre, match, mismatch,
gap_open)
```

```
('NM-FFVS-ANPW', '-M-NAHT-A-FL', -8)
Centre = -M-NAHT-A-FL
```

```
global_align(seqs[2], centre, match, mismatch,
gap_open)
('-MLN--E-L-QQ', ' -M-NAHT-A-FL', -12)
Centre = -M-NAHT-A-FL
```

```
initial score: -45
initial alignment
```

```
-MEKKITGY-TT
NM-FFVS-ANPW
-MLN--E-L-QQ
-M-NAHT-A-FL
```

Blocks = 2:11

```
-MEKKITGYTT
NM-FFVSANPW
-MLN--EL-QQ
-M-NAHTA-FL
```

```
Blocks= 2:10
No improvment
```

```
S3: M-NAHT-AFL
S0: MEKKITGYTT
```

```
S3: -M-NAHT-A-FL
S0: -MEKKITGY-TT
S1: NM-FFVS-ANPW
```

```
S3: -M-NAHT-A-FL
S0: -MEKKITGY-TT
S1: NM-FFVS-ANPW
S2: -MLN--E-L-QQ
```

Example of SP Score with Gap



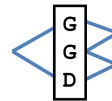
## Example of SP Score with Gap

	F	Y	G	D
F	5	-2	-2	-1
Y		7	1	-5
G			4	-3
D				5

BLOSUM 60

$$m = \begin{matrix} \boxed{F} & \boxed{-} & \boxed{G} \\ \boxed{F} & \boxed{Y} & \boxed{D} \\ \boxed{F} & \boxed{-} & \boxed{D} \end{matrix}$$

Gap penalty = -8  
 $s(-,-) = 0$



$$\begin{aligned} S(m) &= S(m_1) + S(m_2) + S(m_3) \\ &= 3s(F,F) + 2s(-,Y) + s(-,-) + s(G,G) + 2s(G,D) \\ &= 15 - 16 + 0 + 4 - 6 = -3 \end{aligned}$$

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## Star Alignment

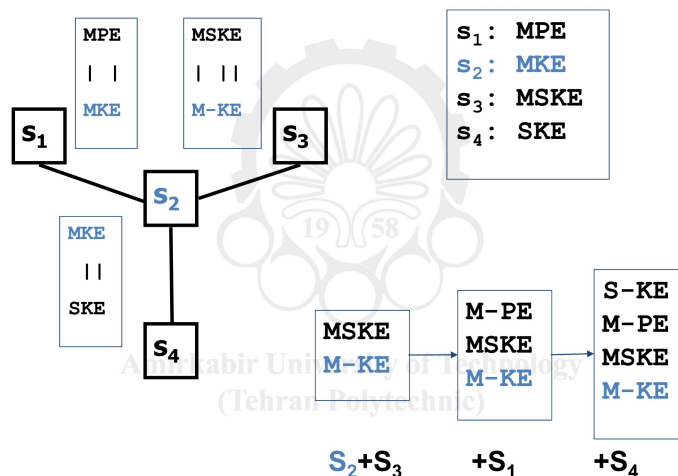
- Fast heuristic to compute MSA
- Good approximation of *optimal* MSA, if scoring scheme satisfies triangle inequality

### Algorithm:

1. Compute pairwise similarities
2. Select center  $s_c$  that maximizes  $\sum_{i \neq c} S(s_c, s_i)$
3. Add sequences in decreasing order of similarity to center  $s_c$ 
  - Rule: “once a gap, always a gap”

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## Step 3 - Add sequences in decreasing order



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## Star-Alignment Example (Cont.)

$S_1$		A	T	T	G	C	C	A	T	T		
$S_2$		A	T	G	G	C	C	A	T	T		
$S_1$		A	T	T	G	C	C	A	T	T	-	-
$S_3$		A	T	C	-	C	A	A	T	T	T	T
$S_1$		A	T	T	G	C	C	A	T	T		
$S_4$		A	T	C	T	T	C	-	T	T		
$S_1$		A	T	T	G	C	C	A	T	T		
$S_5$		A	C	T	G	A	C	C	-	-		

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## Star-Alignment Example (Cont.)

Let's use the alignment of  $S_1$  and  $S_2$ .

$S_1$		A	T	T	G	C	C	A	T	T		
$S_2$		A	T	G	G	C	C	A	T	T		

$S_1$  and  $S_2$  are aligned

Now, let's add  $S_3$ , using its alignment to  $S_1$ .

$S_1$		A	T	T	G	C	C	A	T	T	-	-
$S_2$		A	T	G	G	C	C	A	T	T	-	-
$S_3$		A	T	C	-	C	A	A	T	T	T	T

$S_1$ ,  $S_2$ , and  $S_3$  are aligned

Then, let's add  $S_4$ , using its alignment to  $S_1$ .

$S_1$		A	T	T	G	C	C	A	T	T	-	-
$S_2$		A	T	G	G	C	C	A	T	T	-	-
$S_3$		A	T	C	-	C	A	A	T	T	T	T
$S_4$		A	T	C	T	T	C	-	T	T	-	-

$S_1$ ,  $S_2$ ,  $S_3$ , and  $S_4$  are aligned

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## Star-Alignment Example (Cont.)

Finally, let's add  $S_5$ , using its alignment to  $S_1$ .

$S_1$		A	T	T	G	C	C	A	T	T	-	-
$S_2$		A	T	G	G	C	C	A	T	T	-	-
$S_3$		A	T	C	-	C	A	A	T	T	T	T
$S_4$		A	T	C	T	T	C	-	T	T	-	-
$S_5$		A	C	T	G	A	C	C	-	-	-	-

$S_1$ ,  $S_2$ ,  $S_3$ ,  $S_4$  and  $S_5$  are aligned

For consistency, once a gap is added, it is never removed.