Sequence comparison 1/25

Sequence comparison amounts to comparing the content of 2 sequences.

In simple term, a sequence is a string of letters.

- DNA sequence : A T G G C (4 letters)
- Protein sequence : W C Y L V I (20 letters)

Approach 1: look at sequence composition.

DNA: single nucleotide content
→ weak relationship to specie

Protein: content as tripeptide (3 AA)
- distinguish domains on the tree of life (bacteria, eukaryots ...)

All this is qualitative at best, and a more quantitative approach is needed.

Approach 2 quantification.

We need to start with quantifying the difference between two letters in the sequence alphabet.

(1) Identity.

Build a table:

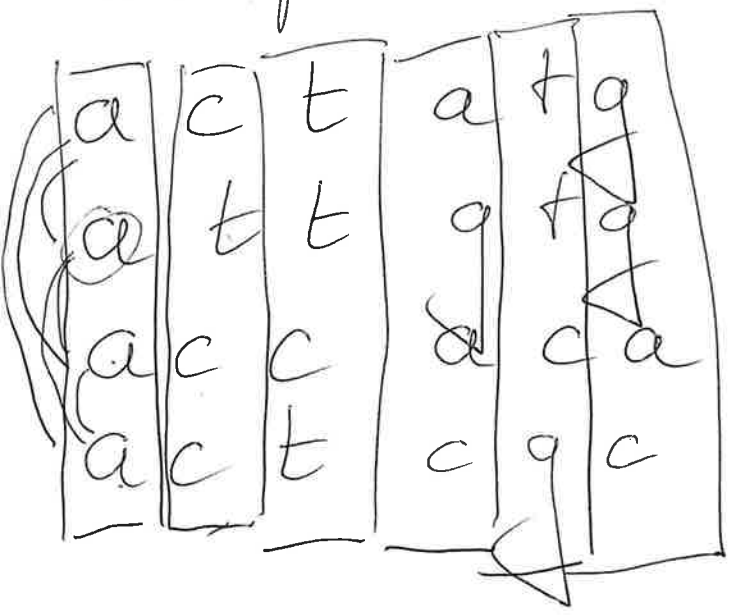
	A	T	G	C
A	1	0	0	0
T	0	1	0	0
G	0	0	1	0
C	0	0	0	1

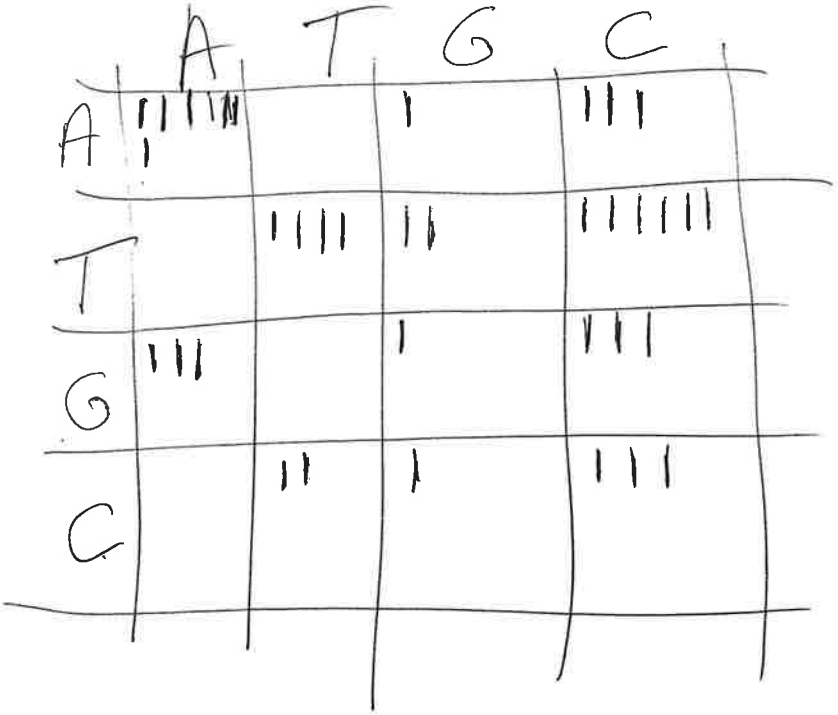
(2) Account for chemistry

	A	T	G	C
A	1	0	0.5	0
T	0	1	0	0.5
G	0.5	0	1	0
C	0	0.5	0	1

(3) Probe nature

Example:





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

	A	T	G	C
A	$\frac{7}{11}$	0	$\frac{1}{11}$	$\frac{3}{11}$
T	0	$\frac{1}{3}$	$\frac{1}{6}$	$\frac{1}{2}$
G	$\frac{3}{7}$	0	$\frac{1}{7}$	$\frac{3}{7}$
C	0	$\frac{1}{3}$	$\frac{1}{6}$	$\frac{1}{2}$

In the table above, I was able to define quantities of the form P_{ij} that represents the probability to replace a letter i with a letter j.

I really want:

$$S_{ij} = \frac{P_{ij}}{P_i P_j}$$

For convenience,

$$B_{ij} = \log S_{ij} = \log \frac{P_{ij}}{P_i P_j}$$

Which sequences should we use to build those preferences?

The most common sequences and corresponding tables are referred to as BLOSUM:

a) Use sequences that are at least 90% identical

→ BLOSUM 90

b) Use sequences that are at least 50% identical

→ BLOSUM 50

The best "universal" matrix is BLOSUM 62