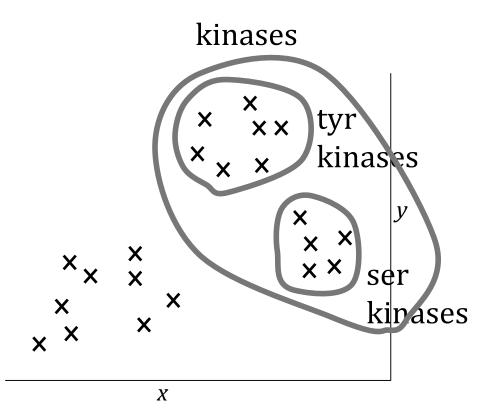
Protein spaces

Why?

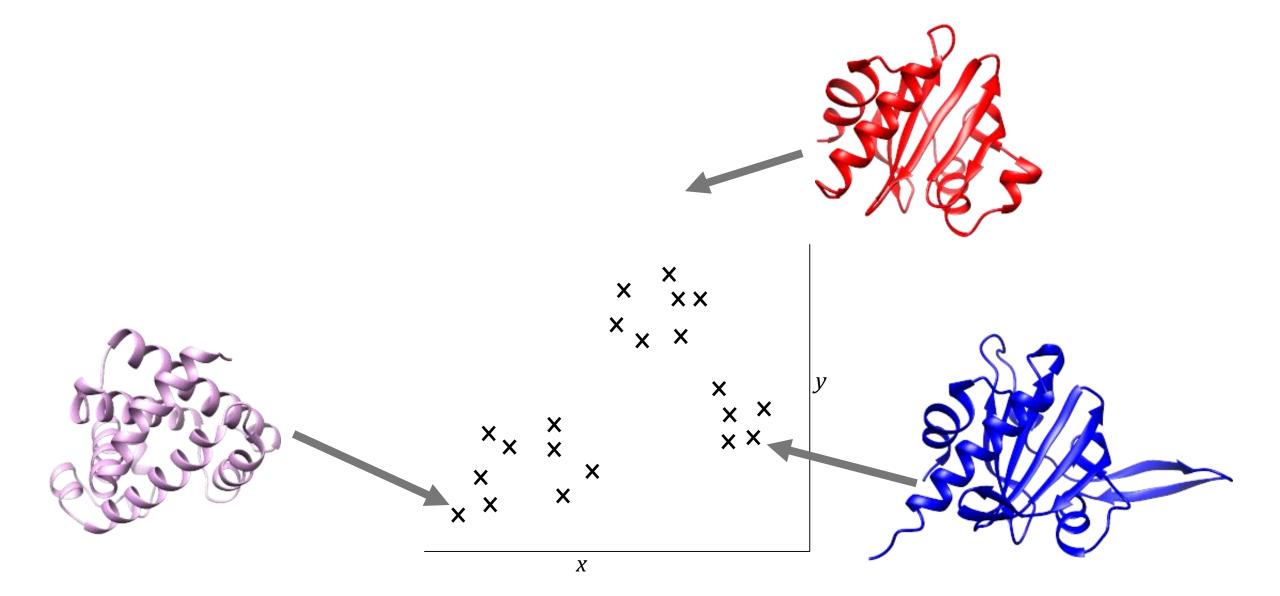
- We like to compare objects prediction of properties
- groups of proteins do they exist?

Spaces – do they exist?

- what is protein space ?
- who cares ?



A space of protein structures ?



who talks about spaces ?

Here

- sequence space (proteins)
- structure space (proteins)

Others

- small molecule space drug space
- tree space
- the set of solutions to a combinatorial problem (not really spaces)
 - how many paths does the travelling salesman problem offer ?

What does a space mean to me?

• usually a classic vector space / rarely a discrete space

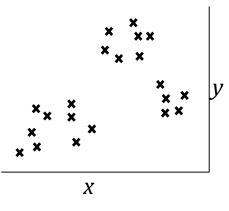
The questions

- I want spaces that are
- objective
- reproducible
- accurate (you cannot have everything)

Proteins

- sequence space (discrete)
- structure space (continuous)
 - sequence and other spaces continuous





Spaces

Conventional spaces

- 1D (x), 2D (x, y), 3D (x, y, z)
 - 4D (*x*, *y*, *z*, *w*), ...
- let us estimate how big a space or problem is examples

Sequence alignments – picking penalties

- 1. gap opening
- 2. gap widening

The optimal parameters are a point in a 2D space (one point)

Discrete spaces

Discrete space

- how many variables do I have ? (*a*, *b*, *c*, ...)
- how many values can each variable have ?
 - *a* 3 values, *b* 4 values, *c* 5
 - number of points in space = 3 × 4 × 5

Representing a Sequence

Protein sequence and structural coordinates

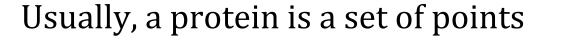
	1	2	3	4	5	6	7	N _{res}
X	1.2	2.3						10.3
у	2.4	3.5						11.1
Ζ	1.7	2.9						15.5
seq	W	Α	С	А	А			D

A protein is a set of 3D points

A protein is a set of 4D points / descriptors if we add sequence

- 4th dimension is not continuous
- This is NOT sequence space

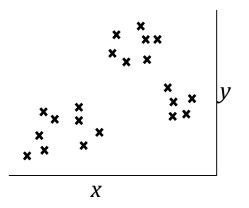
The sequence points



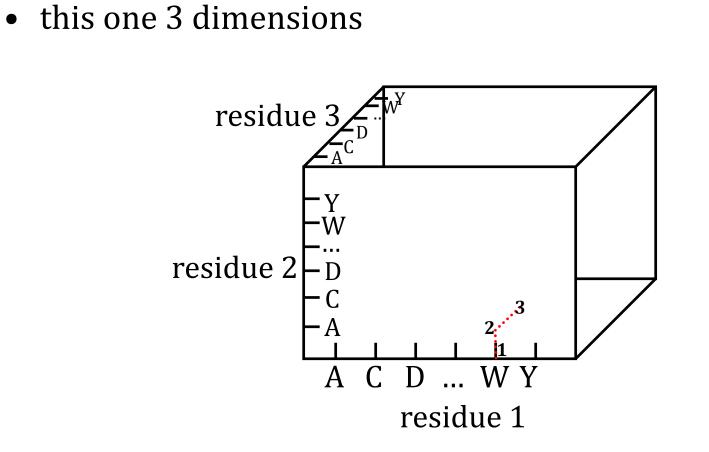
I want one point = one protein

Consider proteins of length N_{res}

• look at the first few (3) points



Finding a Sequence in This Space



Real diagram is a box of N_{res} dimensions

• looking for sequences...

1								
z seq	1.7 W	2.9 A	 С	A	A			15.5 D
	2.4							11.1
Х	1.2	2.3						10.3
	1	2	3	4	5	6	7	 Nres

Families in Sequence Space

Similar sequences are near each other

How realistic?

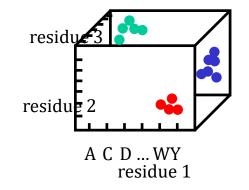
- only works for $N_{seq1} = N_{seq2}$ Conceptual or practical
- important for discussions about protein families (conceptual)
- would you use it directly ? maybe with multiple sequence alignments

What is really ugly ?

• there is no natural ordering on axes

Summary

• we have a discrete space in which every protein is a point

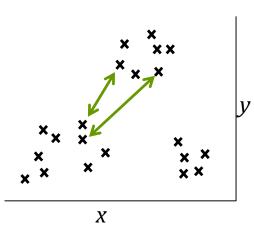


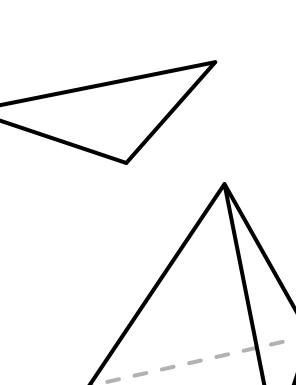
General continuous spaces

My sequence space

- conceptually useful / practically less so
- A generally useful approach
- 2 points fit in 1D (or less)
- 3 points fit in 2D (or less)
- N points can always fit into N-1 dimensions (maybe less)

• my diagrams are usually 2D





Some protein spaces

Do I have a measure of similarity ? Many

Sequence-based

- % sequence similarity
- alignment scores
- *k*-mer similarity, ..

Structure-based

- superimpose and look at geometry
- count similarities in secondary structure elements

General rule

• If I can define similarities there is an implied space

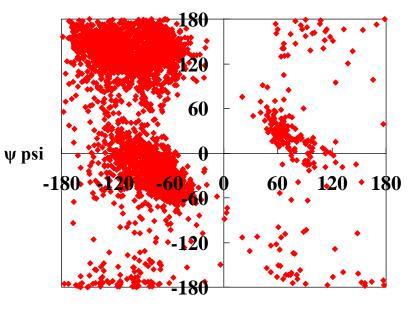
How big?

Sequence space ? (discrete)

• $20 \times 20 \times 20 \dots = 20^N$

Conformational space – how to argue

- for each residue, there are at least 2 major groups (really more)
- maybe chop plot into 3 or 5 pieces
 - say there are *c* conformational possibilities
- c^N for some c
- these spaces grow exponentially in the size of the protein

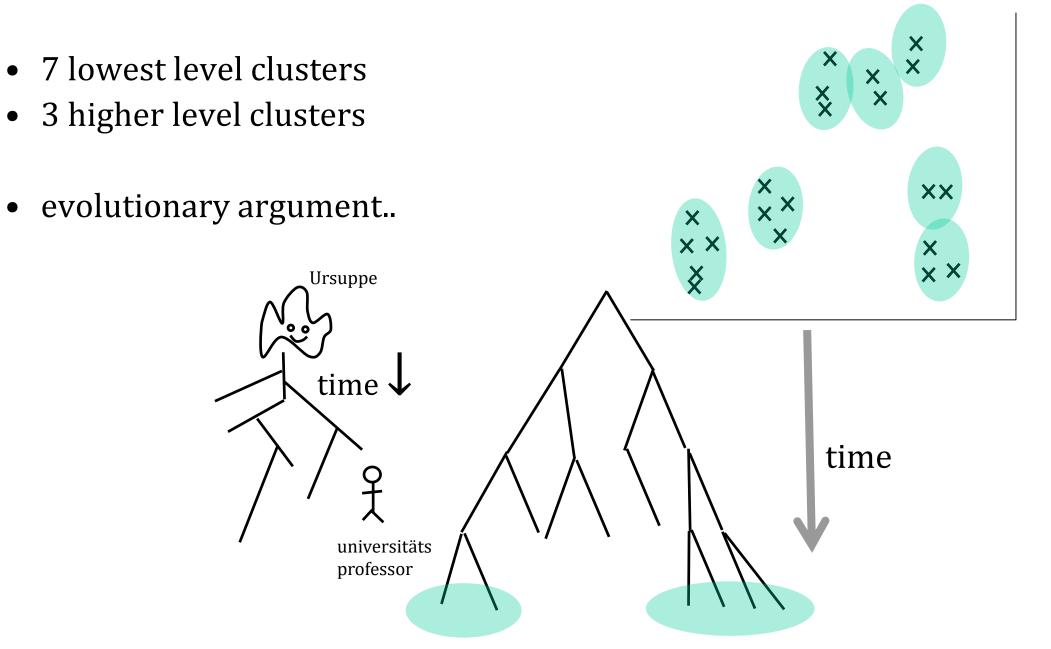


φ phi

How general

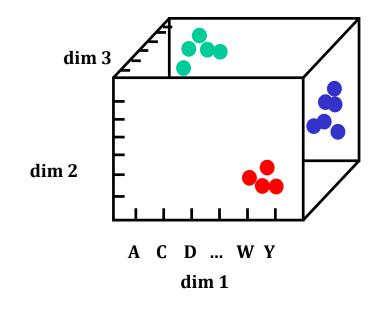
- You can usually invent a space
- High dimensional spaces are not much fun (directly)
 - what do you do with 7-dimensional coordinates?
- What does one normally do?
- reduce to fewer dimensions find the best 2 or 3-dimensional representation of the data
 - distance geometry (and others) OR
- work with distances coordinates are just something to think about

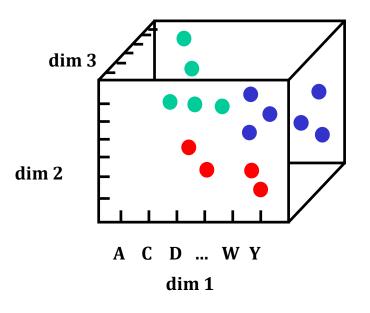
Should we expect a hierarchy ?



Do we expect protein families ?

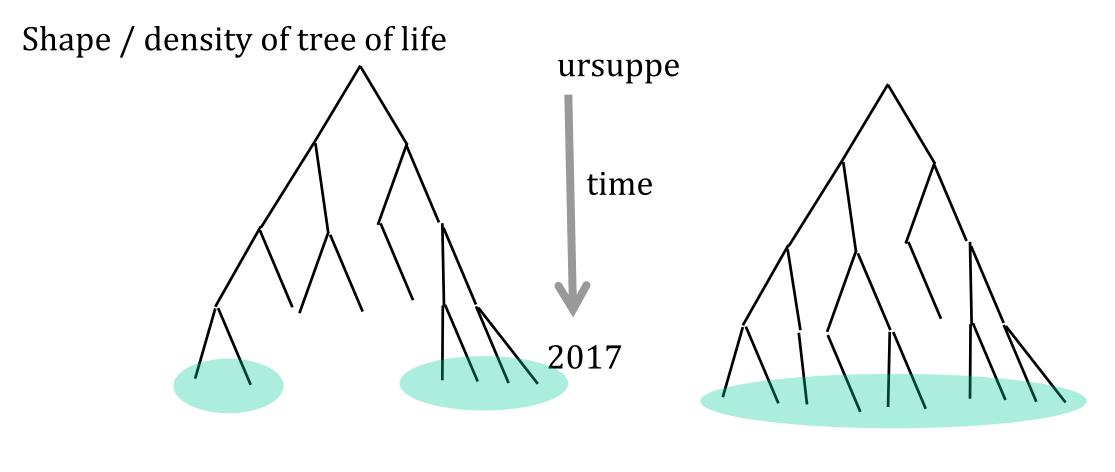
- No real answer
- we have an idea of spaces sequence or structure based
- how are proteins distributed ?





• should you expect clusters ?

Evolution and phylogeny



clear families

no families

Next

- People like to classify proteins
- Should one expect a hierarchy ? Maybe
- Should one expect clusters ? Yes
- what are the distance measures between proteins ?