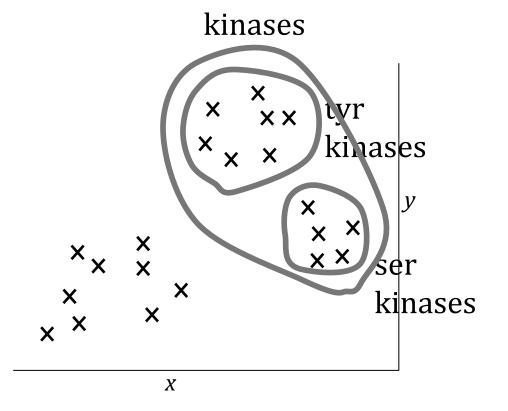
## **Protein spaces**

### Why?

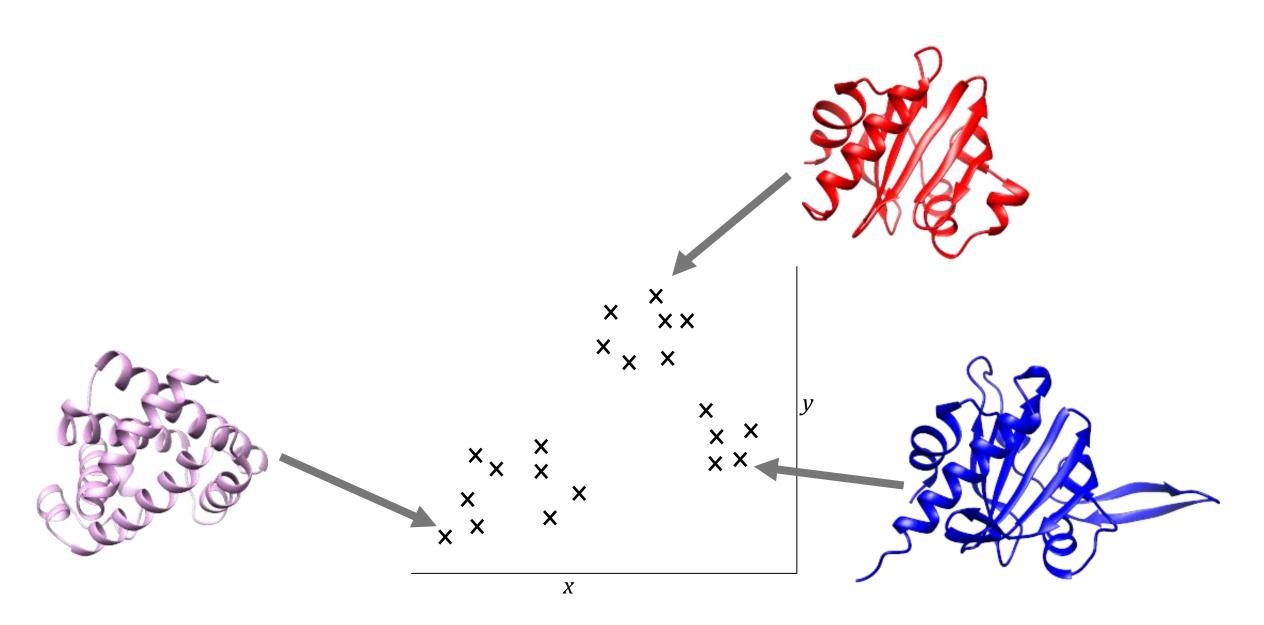
- We like to compare objects prediction of properties
- groups of proteins do they exist?

## 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

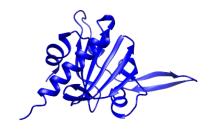
### I want spaces that are

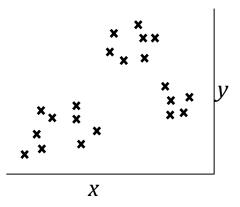
- objective
- reproducible
- accurate (you cannot have everything)

#### **Proteins**

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







# **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 =  $3 \times 4 \times 5$

## 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
$\boldsymbol{Z}$	1.7	2.9						15.5
seq	W	A	С	A	A			D

A protein is a set of 3D points

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

- 4<sup>th</sup> dimension is not continuous
- This is NOT sequence space

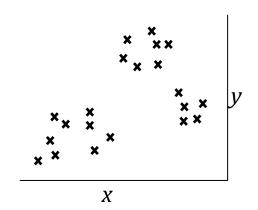
## The sequence points

Usually, a protein is a set of points

I want one point = one protein

Consider proteins of length  $N_{res}$ 

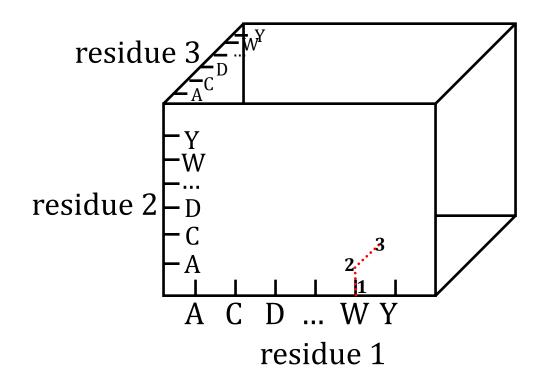
• look at the first few (3) points

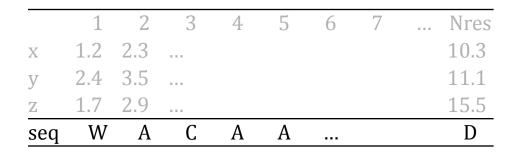


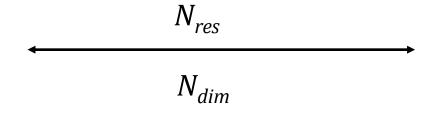
## Finding a Sequence in This Space

Real diagram is a box of  $N_{res}$  dimensions

• this one 3 dimensions







• looking for sequences...

# **Families in Sequence Space**

Similar sequences are near each other

#### How realistic?

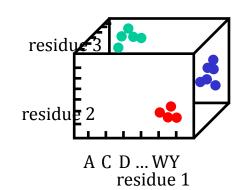
- only works for  $N_{seq1} = N_{seq2}$
- Conceptual or practical
- important for discussions about protein families (conceptual)
- would you use it directly? maybe with multiple sequence alignments

### 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

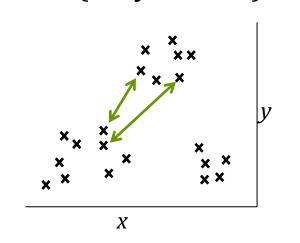
### My sequence space

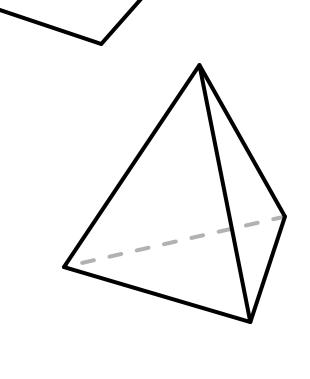
conceptually useful / practically less so

### 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

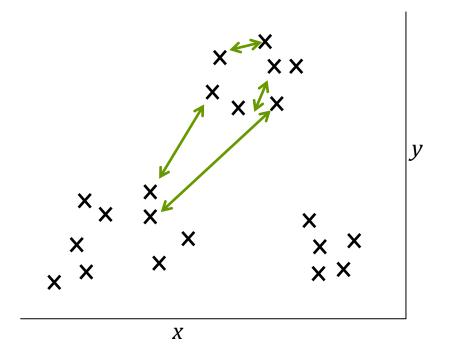
- % sequence similarity
- alignment scores
- *k*-mer similarity, ..

#### 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



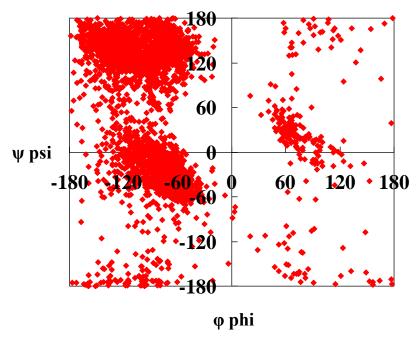
# How big?

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



## How general

- You can usually invent a space
- High dimensional spaces are not much fun (directly)
  - what do you do with 7-dimensional coordinates?

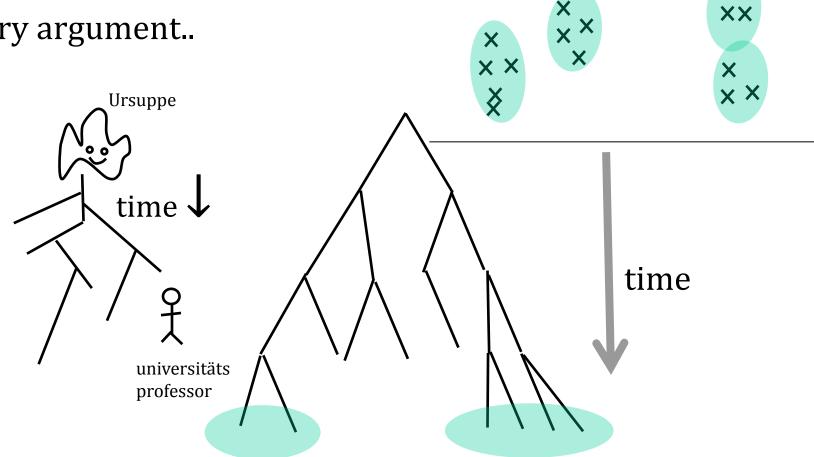
What does one normally do?

- 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?

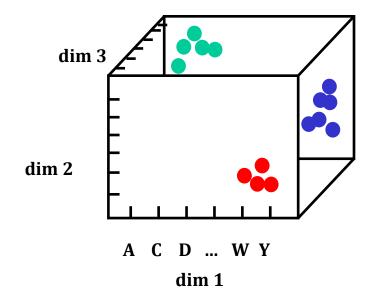
- 7 lowest level clusters
- 3 higher level clusters

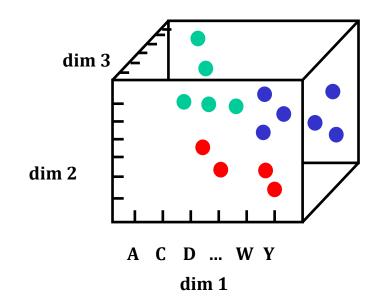




## Do we expect protein families?

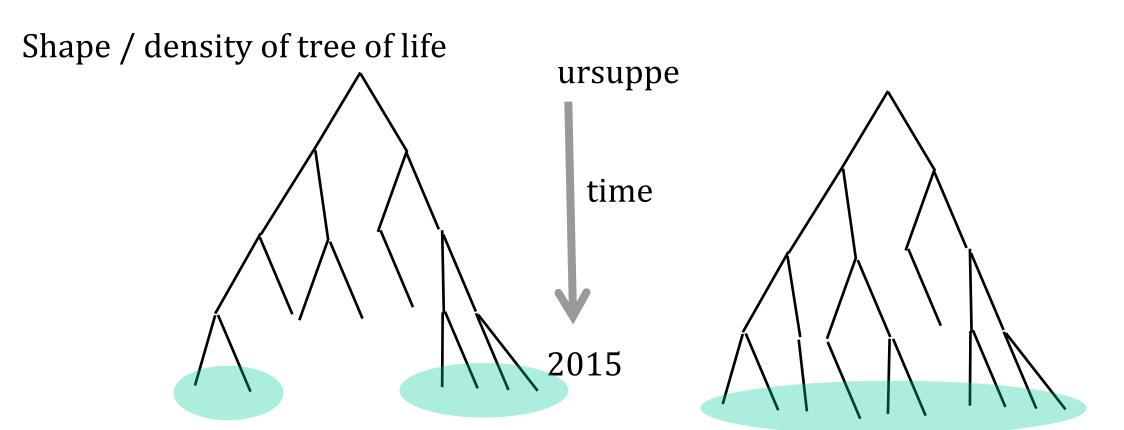
- No real answer
- we have an idea of spaces sequence or structure based
- how are proteins distributed?





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?