

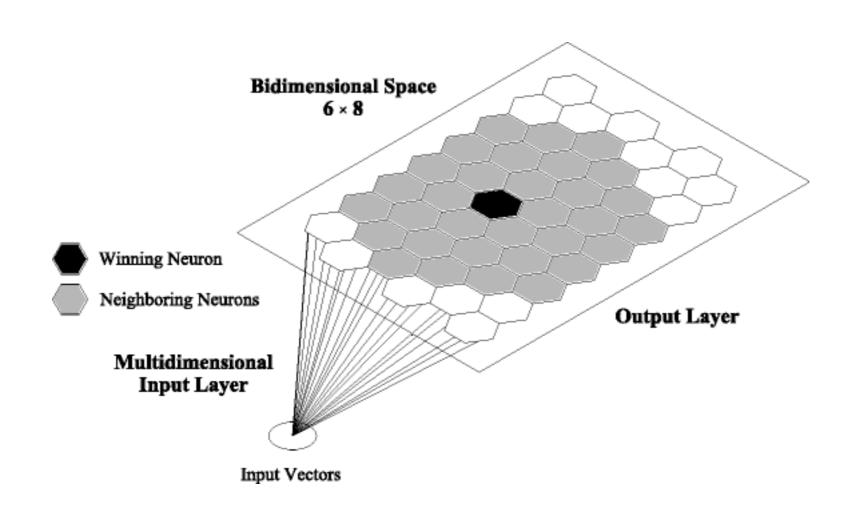


- A neural network approach to unsupervised machine learning.
- Appealing visual presentation of learned results as a 2D map.
- Learned result represents a partitioning of the input space according to a similarity metric.

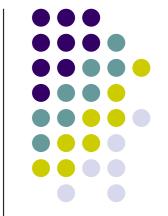
T. Kohonen, Self-organizing maps, 3rd ed. Berlin; New York: Springer, 2001.

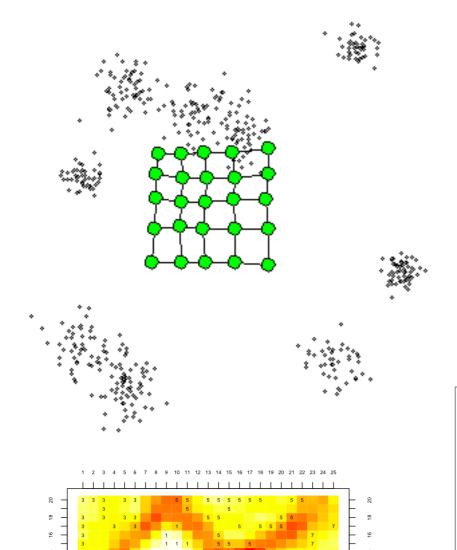
#### **SOM Architecture**





# Insight: SOMs Sample the Data Space





- Given some distribution in the data space, SOM will try to construct a sample that looks like it was drawn from the same distribution.
- It will construct the sample using interpolation (neighborhood function) and constraints (the map grid).
- We can then measure the quality of the map using a statistical two sample approach.

#### Algorithm:

```
Repeat until Done

For each training observation Do

Find the neuron that best matches the observation.

Make that neuron look more like the observation.

Smooth the immediate neighborhood of that neuron.

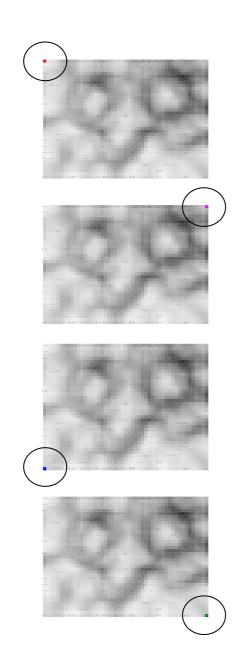
End For

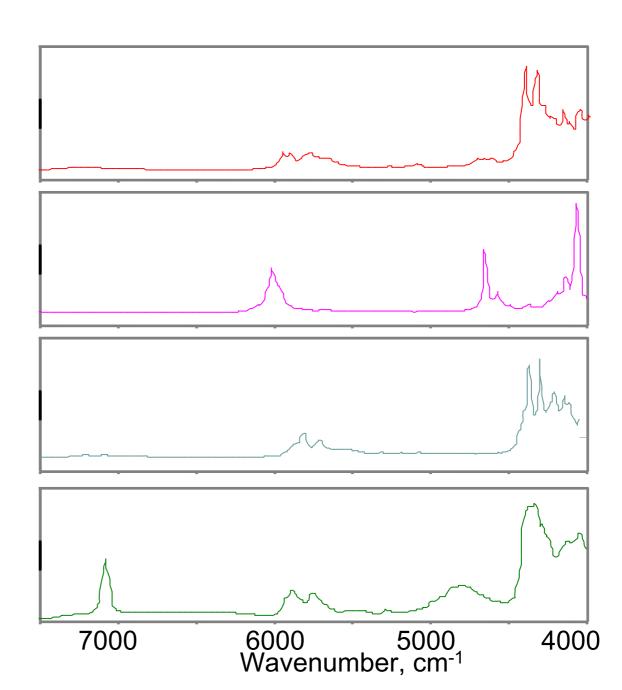
End Repeat
```



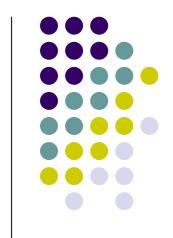
#### **SOM Neurons**

SOM neurons capture information about the input data – similar to the centroids in k-means.





## Training a SOM



	small	medium	big	Two legs	l	Hair	Hooves	Mane	Feathers	Hunt	Run	Fly	Swim
Dove	1	0	0	1	0	0	0	0	1	0	0	1	0
Hen	1	0	0	1	0	0	0	0	1	0	0	0	0
Duck	1	0	0	1	0	0	0	0	1	0	0	0	1
Goose	1	0	0	1	0	0	0	0	1	0	0	1	1
Owe	1	0	0	1	0	0	0	0	1	1	0	1	0
Hawk	1	0	0	1	0	0	0	0	1	1	0	1	0
Eagle	0	1	0	1	0	0	0	0	1	1	0	1	0
Fox	0	1	0	0	1	1	0	0	0	1	0	0	0
Dog	0	1	0	0	1	1	0	0	0	0	1	0	0
Wolf	0	1	0	0	1	1	0	1	0	1	1	0	0
Cat	1	0	0	0	1	1	0	0	0	1	0	0	0
Tiger	0	0	1	0	1	1	0	0	0	1	1	0	0
Lion	0	0	1	0	1	1	0	1	0	1	1	0	0
Horse	0	0	1	0	1	1	1	1	0	0	1	0	0
Zebra	0	0	1	0	1	1	1	1	0	0	1	0	0
Cow	0	0	1	0	1	1	1	0	0	0	0	0	0

Table of Feature Vectors

#### Algorithm:

Repeat until Done

For each row in Data Table Do

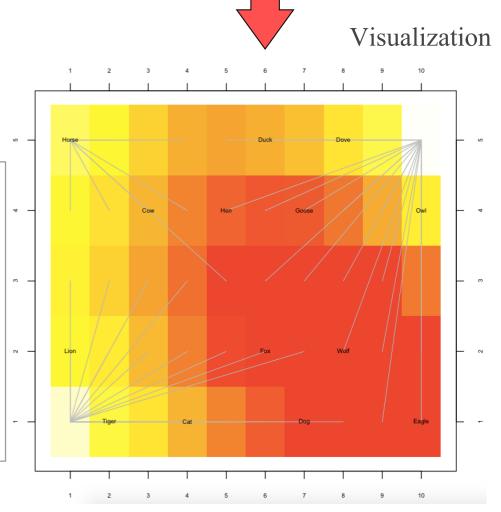
Find the neuron that best describes the row.

Make that neuron look more like the row.

Smooth the immediate neighborhood of that neuron.

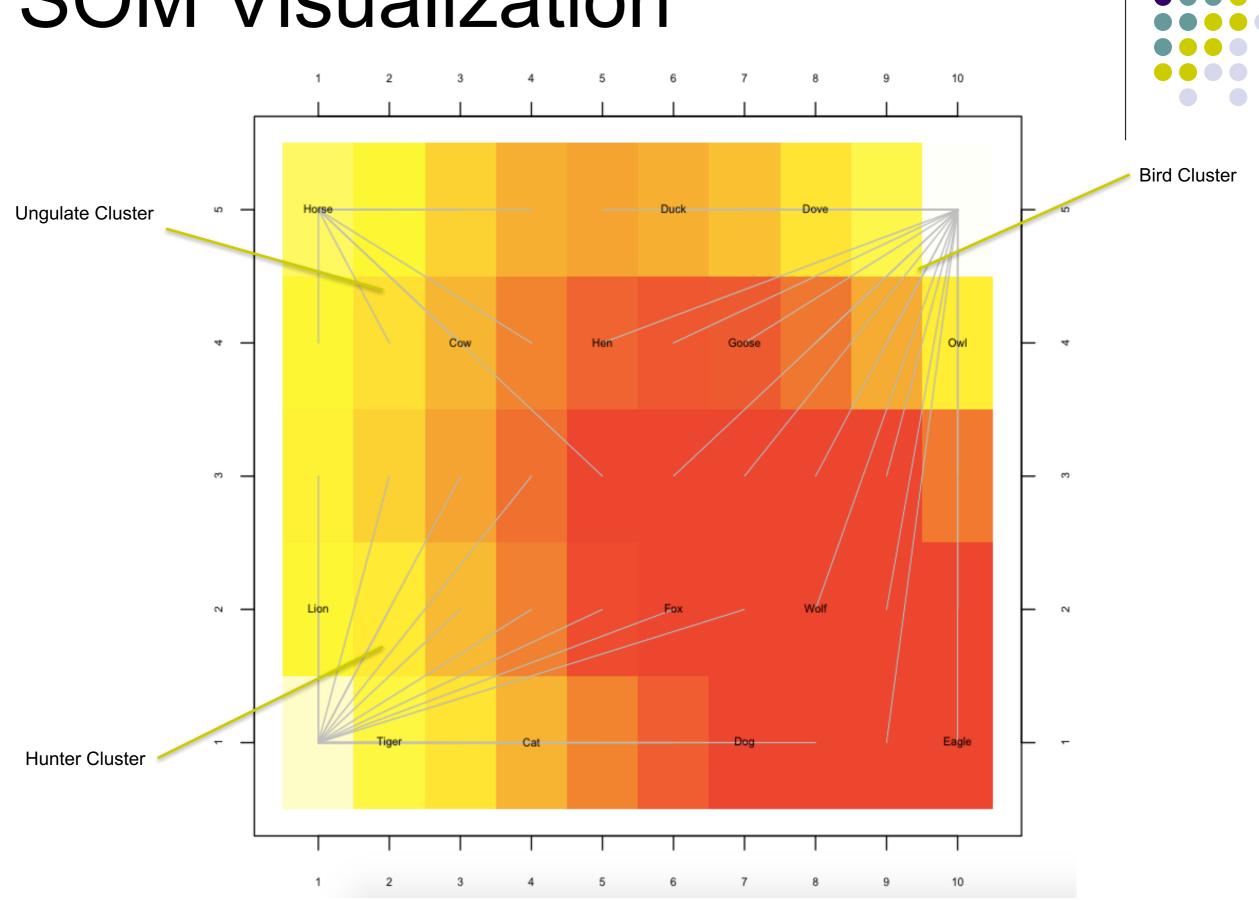
End For

End Repeat



"Grid of Neurons"

### **SOM Visualization**



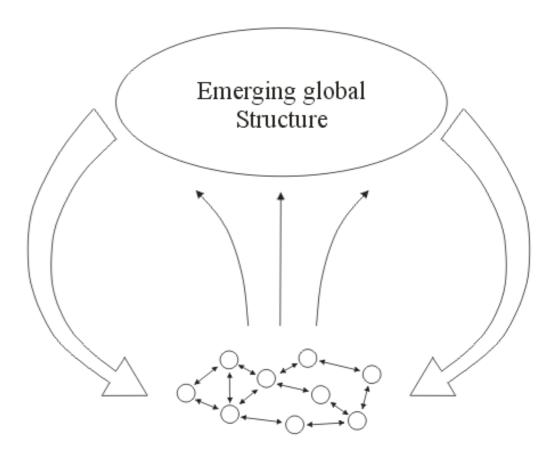
# Self-Organization and Learning



- Self-organization refers to a process in which the internal organization of a system increases automatically without being guided or managed by an outside source.
- This process is due to <u>local</u> interaction with <u>simple rules</u>.
- Local interaction gives rise to global structure.

We can interpret emerging global structures as <u>learned</u> structures.

Learned structures appear as <u>clusters</u> of similar objects.



Local Interaction

Complexity: Life at the Edge of Chaos, Roger Lewin, University Of Chicago Press; 2nd edition, 2000





#### Algorithm 1 Iterative SOM training algorithm.

```
1: given:
           \mathbf{m}_i \leftarrow \{\text{neurons for } i = 1, \dots, n\}
           \mathbf{x}_k \leftarrow \{\text{training instances for } k = 1, \dots, l\}
           \eta \leftarrow \{\text{learning rate, } 0 < \eta < 1\}
          h(c,i) \leftarrow \{ \text{loss function for neurons } c \text{ and } i \}
 5:
 6:
 7: repeat
          /*** Select a training instance ***/
          \mathbf{x}_k for some k=1,\ldots,l:
10:
          /*** Find the winning neuron ***/
11:
          c \leftarrow 1
12:
          v \leftarrow ||\mathbf{m}_1 - \mathbf{x}_1||^2
13:
          for i=2, n do
14:
                                                               Compare input to
               d \leftarrow ||\mathbf{m}_i - \mathbf{x}_k||^2
15:
                                                               each neuron
               if d < v then
16:
                    c \leftarrow i
17:
                    v \leftarrow d
18:
               end if
19:
20:
          end for
21:
          /*** Update neighborhood ***/
22:
          for i=1, n do
23:
               \mathbf{m}_i \leftarrow \mathbf{m}_i - \eta(\mathbf{m}_i - \mathbf{x}_k)h(c, i)
                                                                                update all neurons
24:
          end for
25:
26: until done
27: return \mathbf{m}_i for all i = 1, \ldots, n
```

The Loss Function:

$$h(c,i) = \begin{cases} 1 & \text{if } i \in \Gamma(c), \\ 0 & \text{otherwise,} \end{cases}$$

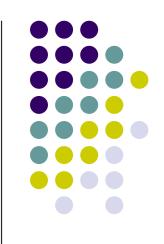
The Neighborhood Function (time dependent):

$$\Gamma(c)|_{t=0} = \{1, 2, \dots, n\}.$$

$$\Gamma(c)|_{t\gg 0} = \{c\}.$$

This is the "standard" training algorithm for SOMs implemented in most packages.

### **Applications of SOM**

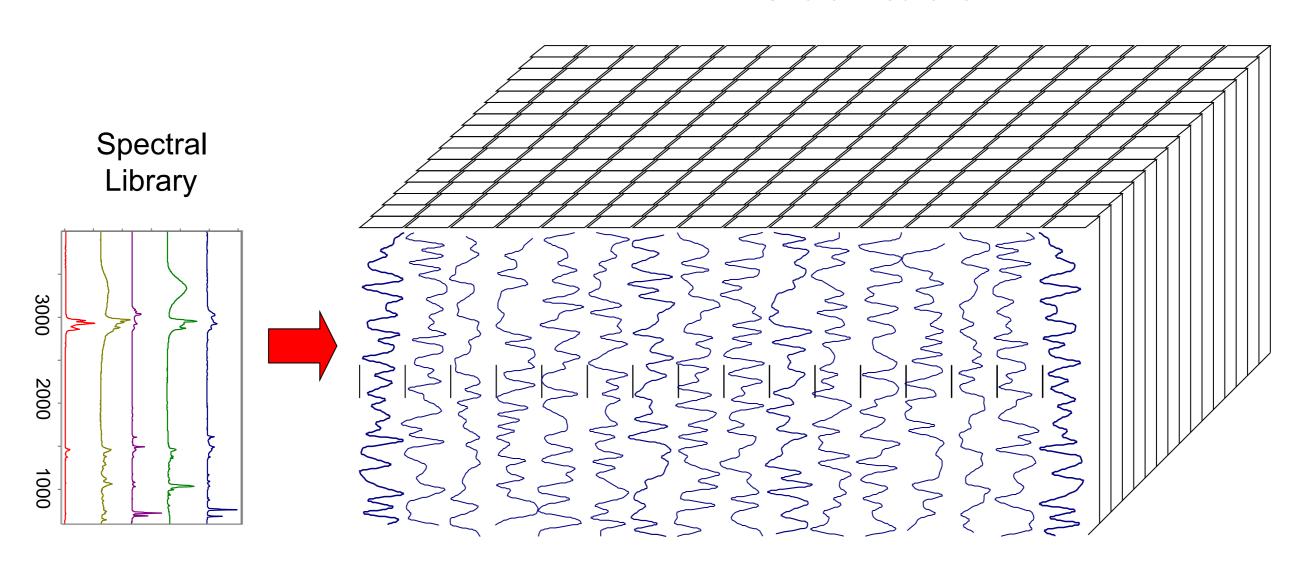


- Infrared Spectroscopy
  - Goal: to find out if compounds are chemically related without performing an expensive chemical analysis.
  - Each compound is tested for light absorbency in the infrared spectrum.
  - Specific chemical structures absorb specific ranges in the infrared spectrum.
  - This means, each compound has a specific "spectral signature".
  - We can use SOMs to investigate similarity.





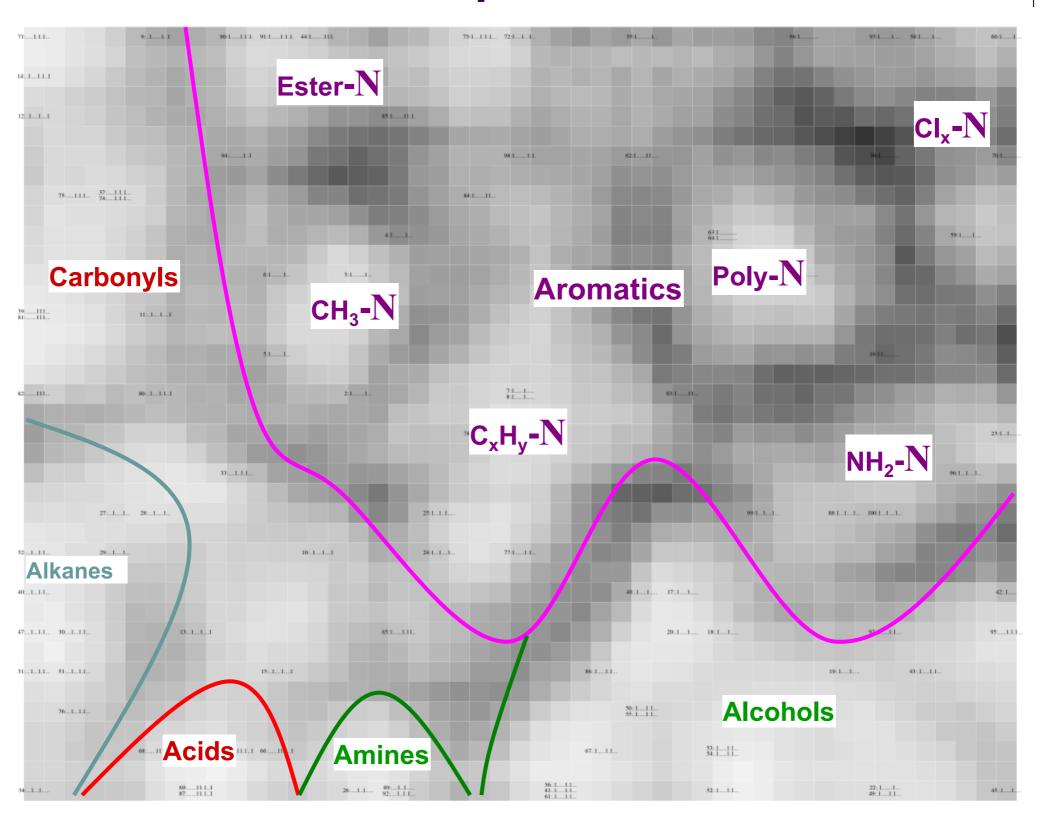
#### **Grid of Neurons**



Random Number Spectra

### **NIR SOM**

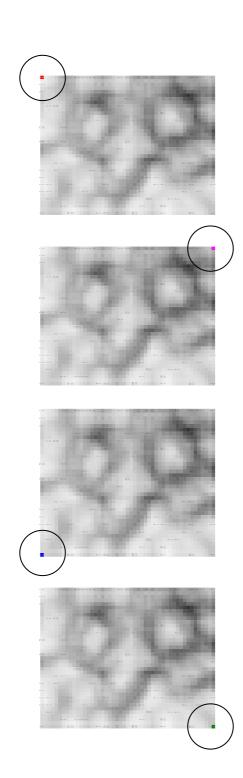
#### **Functional Groups**

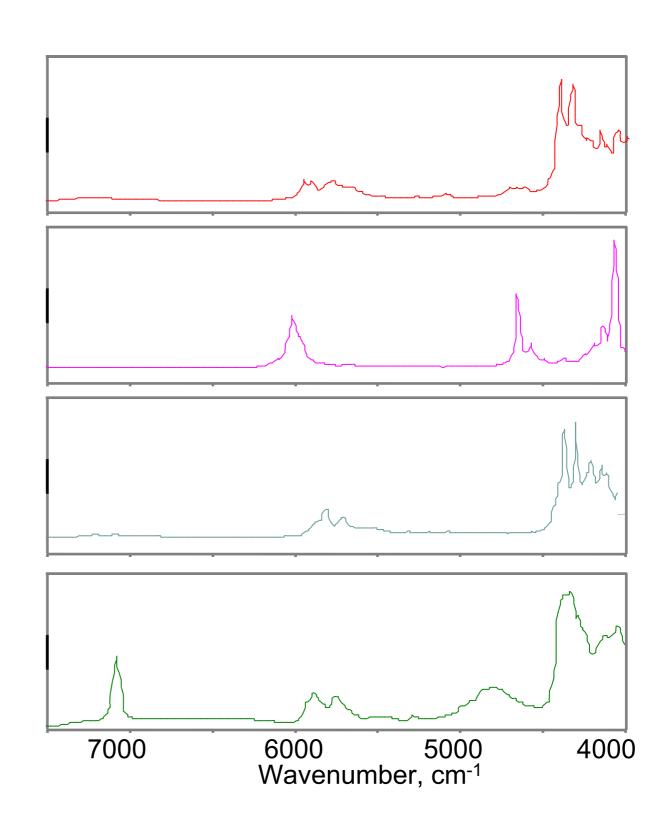




# NIR Centroid Spectra

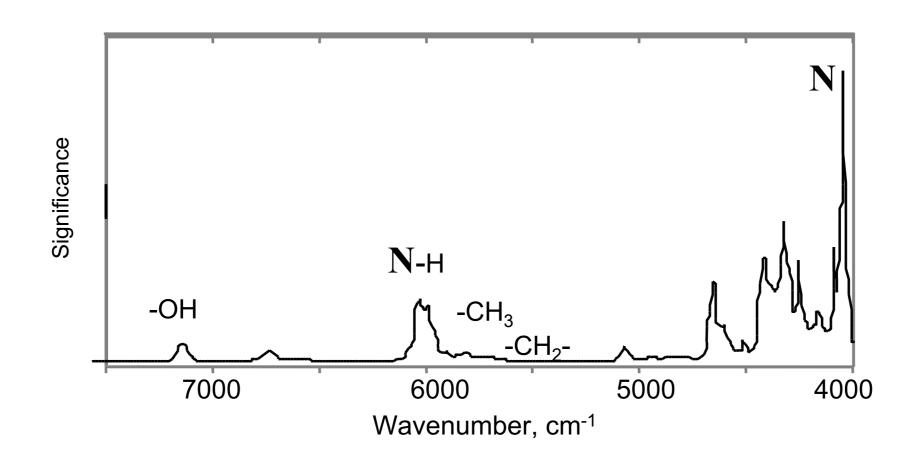






# NIR Significance Spectrum

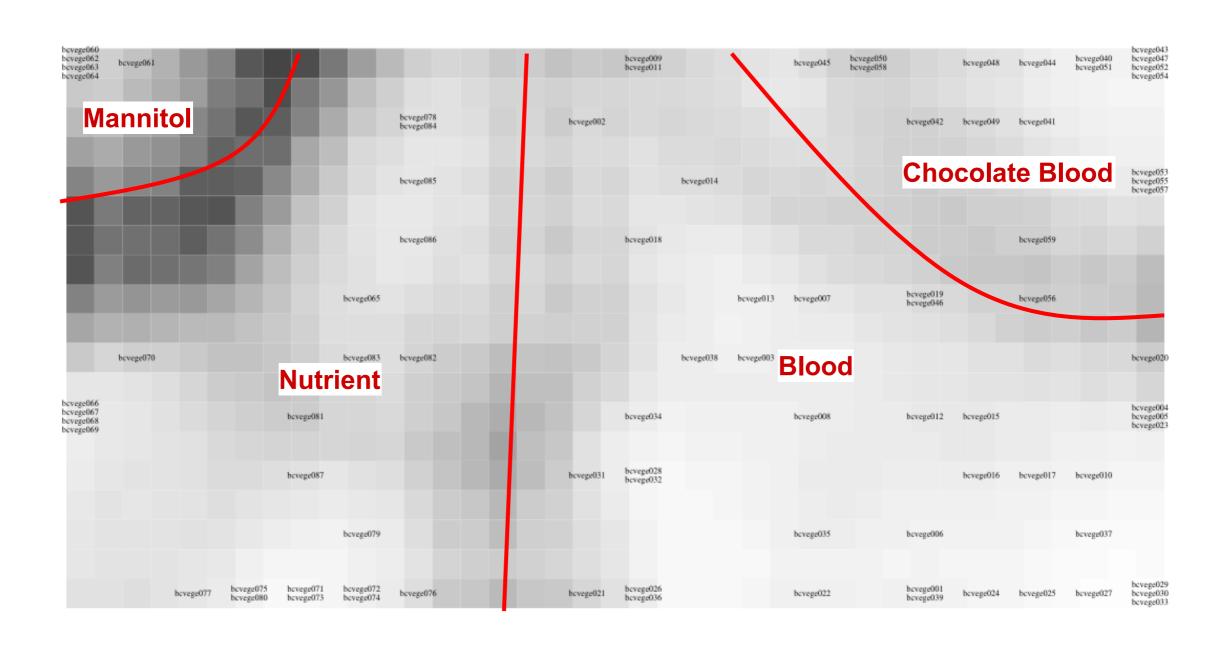




#### SOM

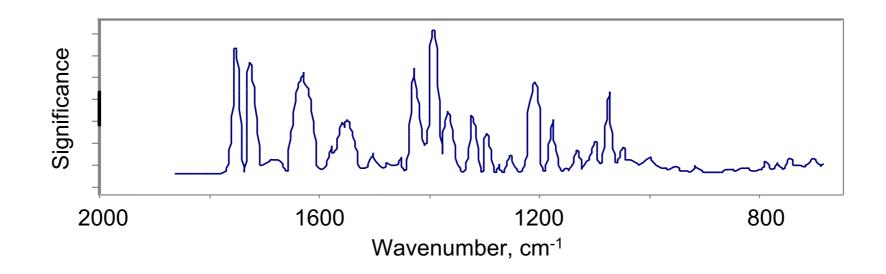
### Bacterium *b-cereus* on different agars "You are what you eat!"





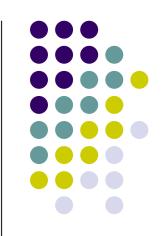
# Significance Spectrum b-cereus on different agars

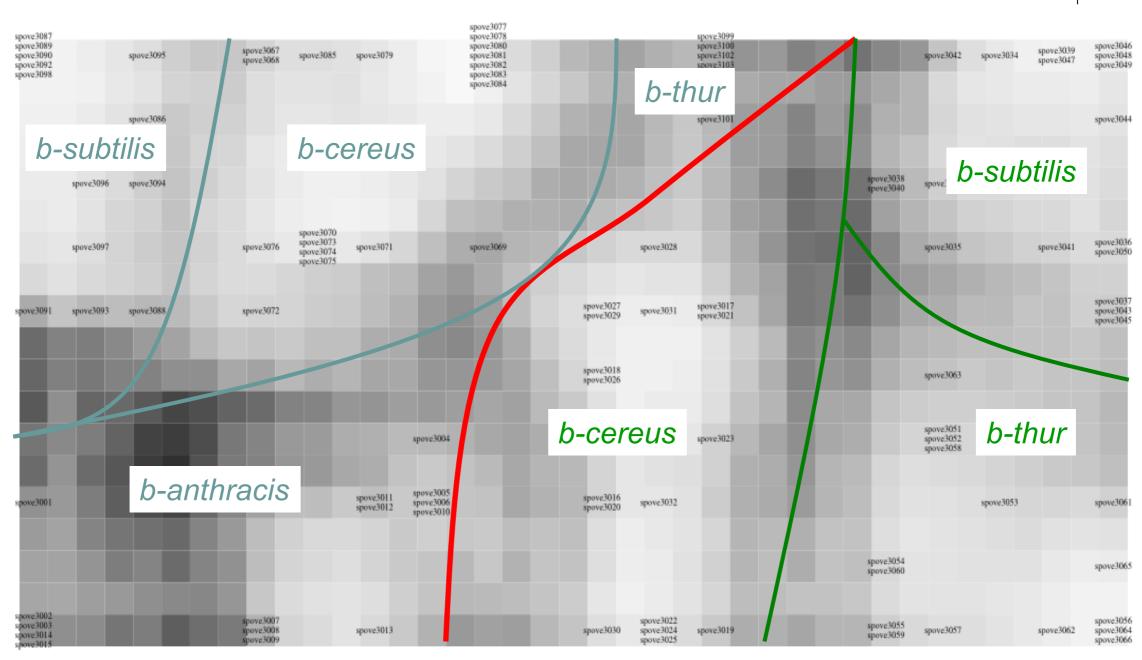




### SOM

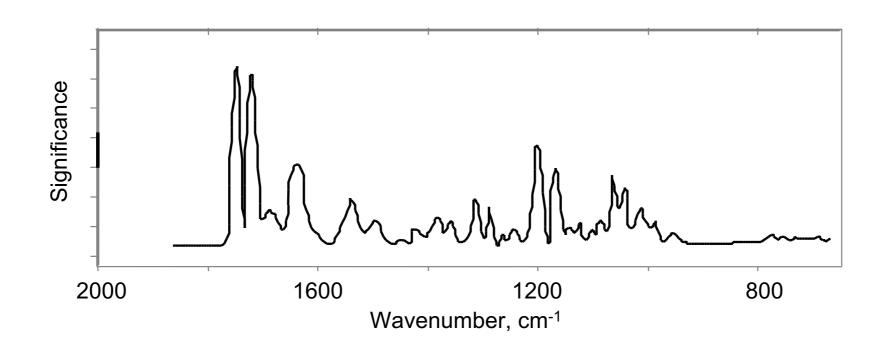
#### **Bacteria Spectra**



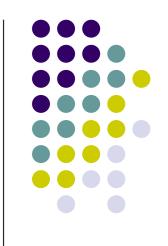


#### Significance Spectrum vs b-subtilis 1st Derivative Spectra









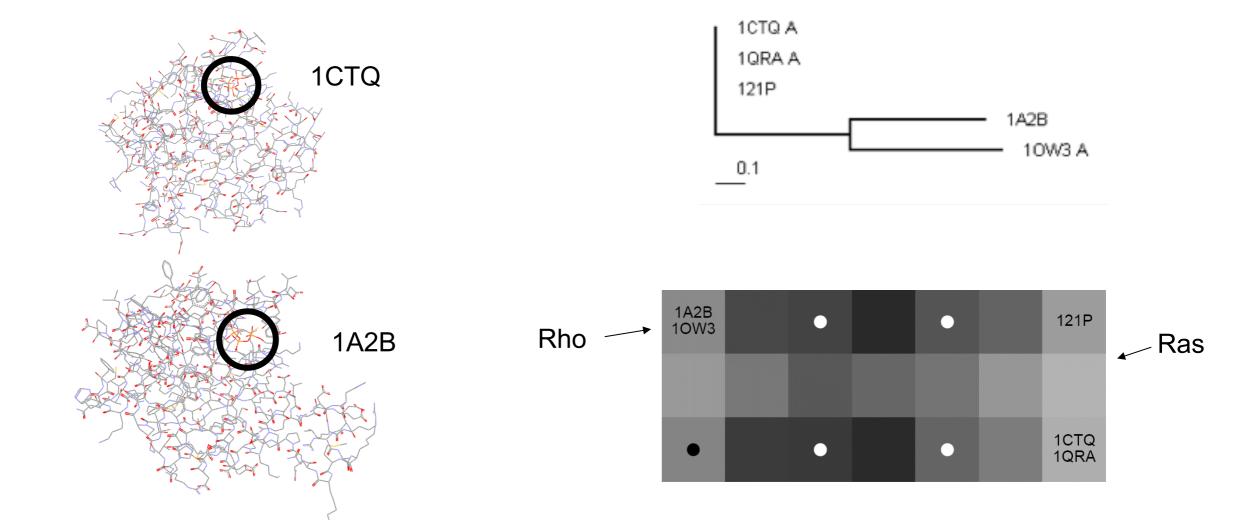
- Clustering Proteins based on the architecture of their activation loops.
  - Align the proteins under investigation.
  - Extract the functional centers.
  - Turn 3D representation into 1D feature vectors.
  - Cluster based on the feature vectors.

# **Structural Classification of GTPases**



Can we structurally distinguish between the Ras and Rho subfamilies?

- Ras: 121P, 1CTQ, and 1QRA
- Rho: 1A2B and 1OW3
- F = p-loop, r = 10Å



#### Demo



- popsom package in R
  - Very fast implementation
  - Statistically motivated convergence criterion



