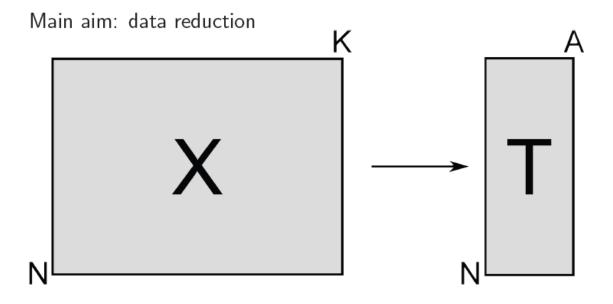
# 2. Principal Component Analysis

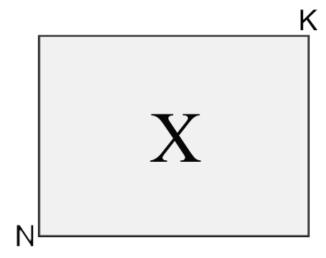
- Visualizing multivariate data
- Geometric interpretation of PCA
- Mathematical interpretation
- Example(s)

# **Principal Component Analysis**



# Visualizing Multivariate Data

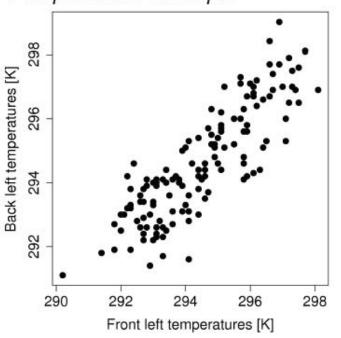
▶ PCA considers a single matrix: **X** 

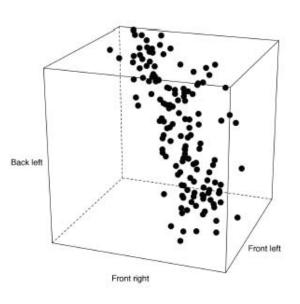


- ▶ *N* observations
- ▶ K variables
- ▶ Which variables go in **X**?

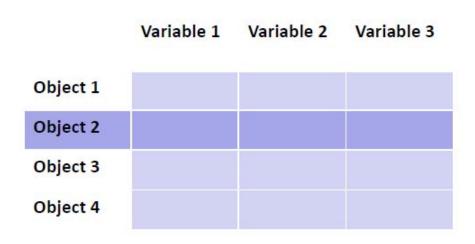
# Visualizing Multivariate Data

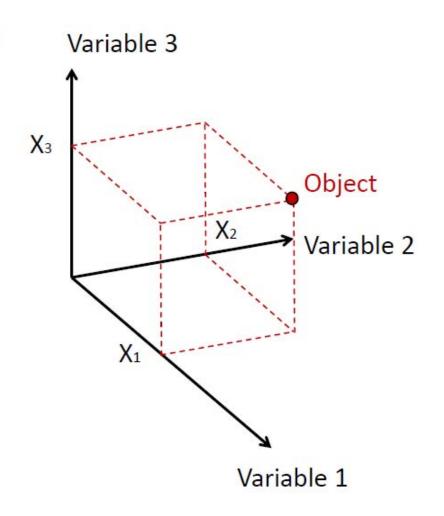
#### Temperature example





- Each variable defines an axis. A coordinate system can be made using all variables, called the variable space.
- Each object is a row in the data table (matrix) and is visualised as a point in the variable space.

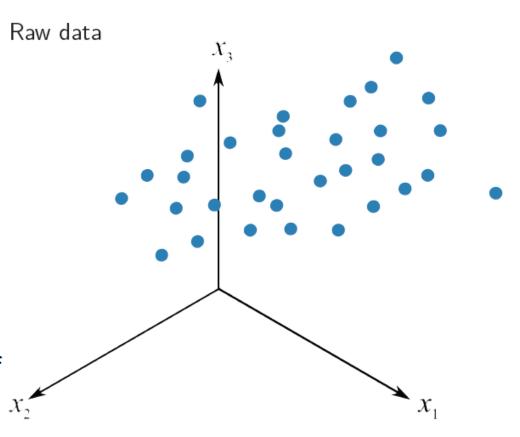




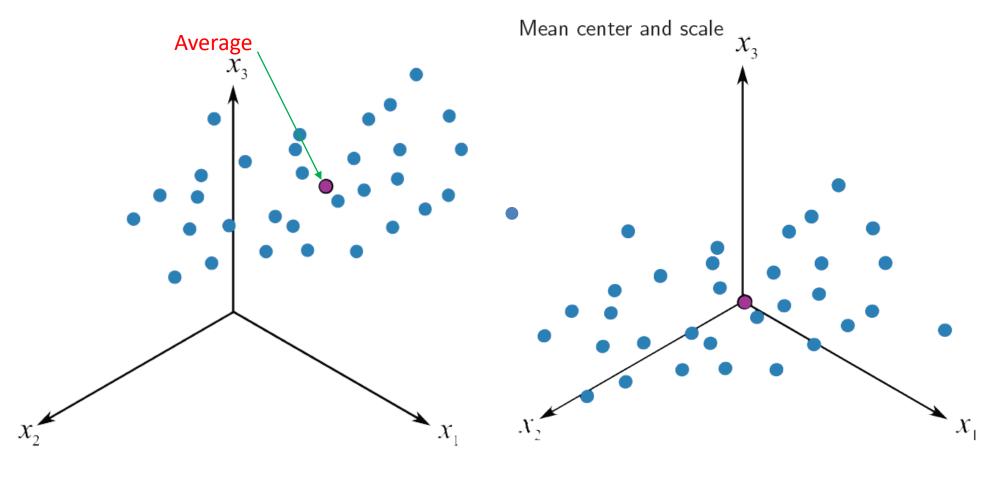
 Each sample / object is represented as a point in the variable space.

 The whole data table constitutes a swarm of points in the variable space.

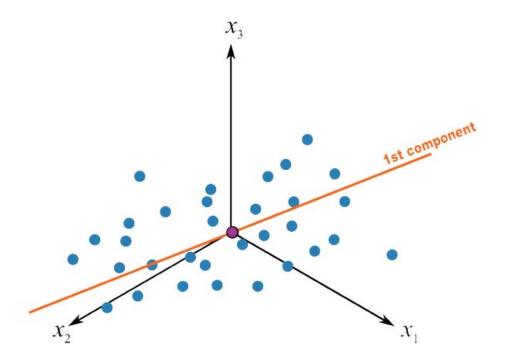
 We would like to find out more about the structure of this swarm.



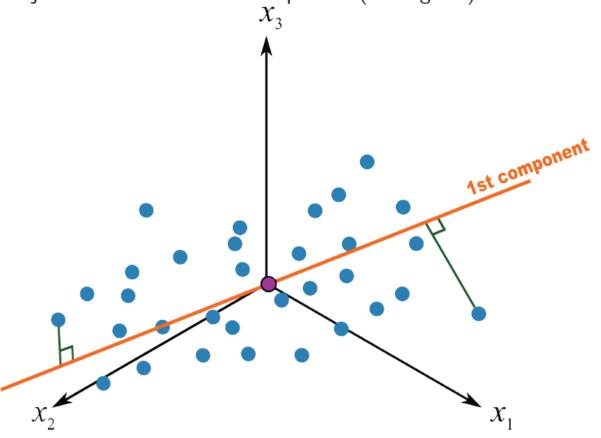
• In many cases, the swarm of points has a specific shape in some direction.



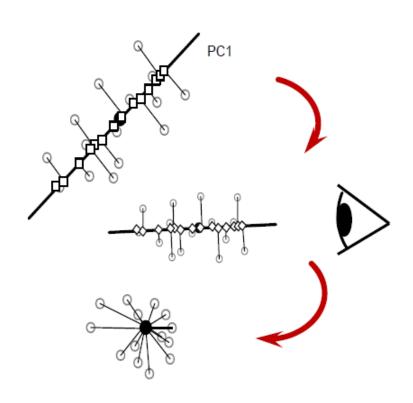
- By picking the direction of largest elongation, this direction will pass through the center of the swarm.
- This line is called the first principal component (PC1) and points in the direction of the maximum data variation.

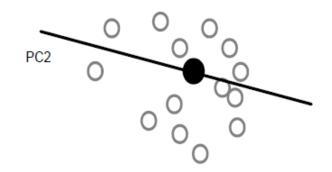


Project observations onto component (90 degrees)

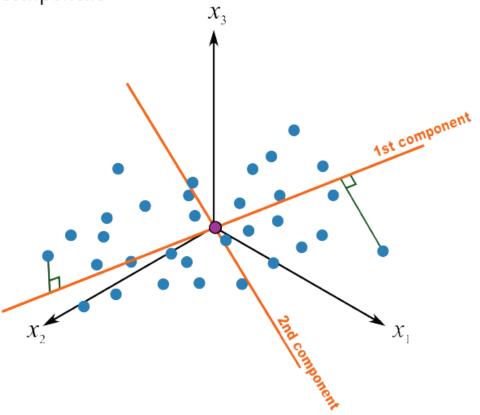


- The first principal component may not be enough to describe the data variation.
- By projecting the samples onto the new coordinate system, there is still unexplained variance (combination of the individual residuals)
- If we repeat the projection process in the remaining part of the space, we will find the second principal component (PC2).

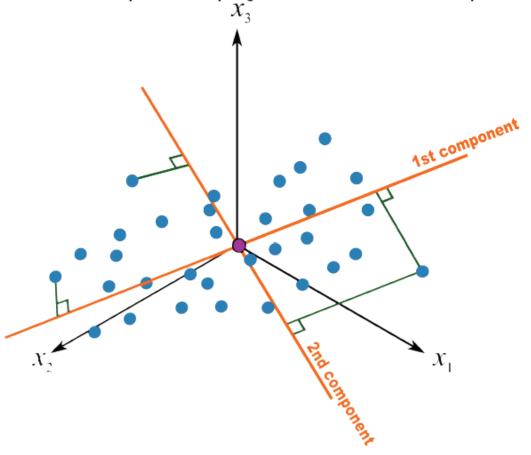




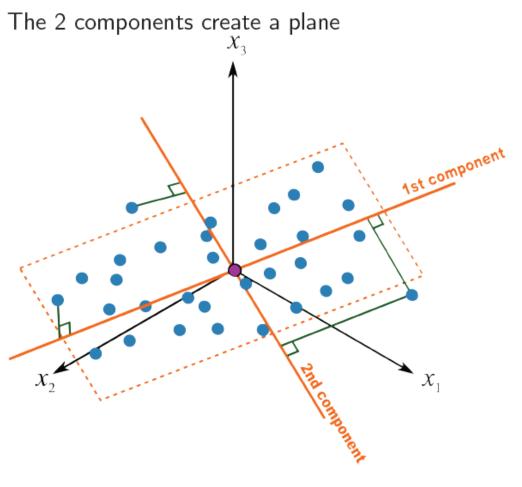
Second component: best-fit line; perpendicular to 1st component



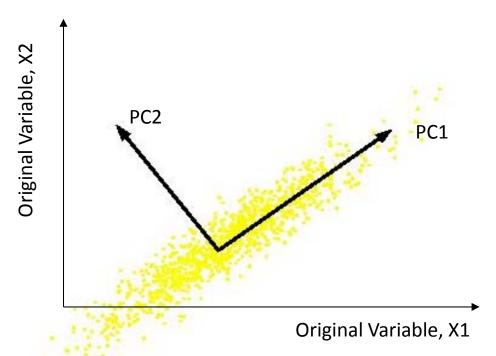
Second component: project onto second component



- With 2 principal components we can build a plane onto which the projections of the swarm lie closer yet to the original variable space.
- We have found the 2dimensional 'window' which best describes the data.
- The process can be continued to find more PCs.



- In summary,
  - PCA finds a few orthogonal axes of greatest variance in data.



 New latent variables are linear combinations of the original variables.

$$PC1 = a_1 X1 + a_2 X2 + a_3 X3$$
  
  $X = Mean + b_1 PC1 + b_2 PC2 + Error$ 

#### Constraints:

- Maximise the dispersion of samples along the latent variables (the variance)
- Orthogonality

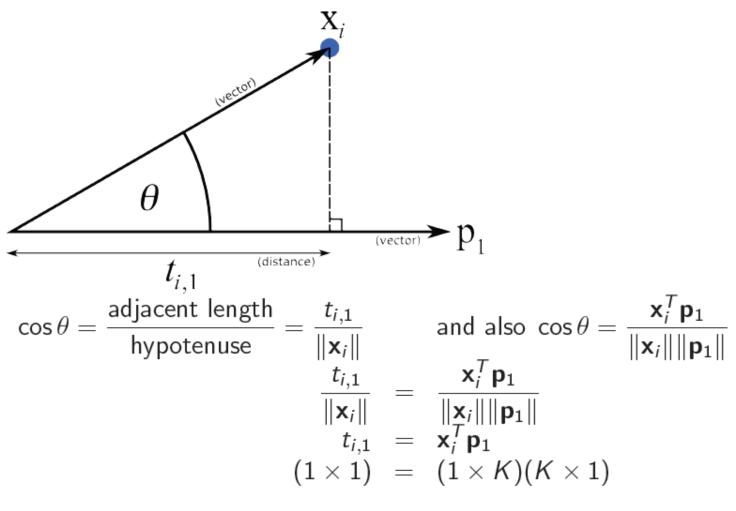
### **Mathematical Derivation**

What has this done?
Break **X** down into 2 parts:

- projected points " on the plane"
- ▶ residual distance " off the plane"

#### **Mathematical Derivation**

• From linear algebra (or engineering mathematics),



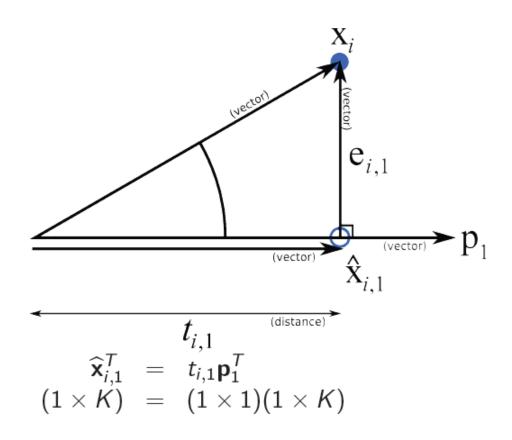
#### **Mathematical Derivation**

$$t_{i,1} = \mathbf{x}_i^T \mathbf{p}_1$$
  
=  $x_{i,1}p_{1,1} + x_{i,2}p_{2,1} + \dots + x_{i,k}p_{k,1} + \dots + x_{i,K}p_{K,1}$ 

- ▶ K separate terms: added up (i.e. linear combination) to give  $t_1$
- ► Entire data set: **T** = **XP**

## Predicted value for each observations

•  $\hat{\mathbf{X}}_i$ :projected version of  $\mathbf{X}_i$ 



#### Predicted value for each observations

Residual vector:

$$\mathbf{e}_{i,A}^{T} = \mathbf{x}_{i}^{T} - \hat{\mathbf{x}}_{i,A}^{T}$$

$$(1 \times K) = (1 \times K) - (1 \times K)$$

Residual distance:

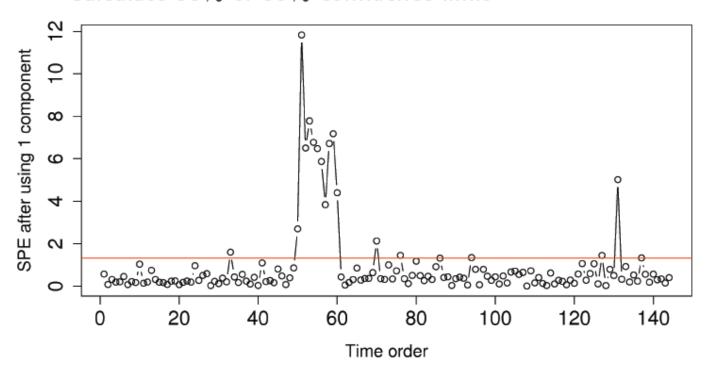
$$SPE_{i} = \sqrt{\mathbf{e}_{i,A}^{T}\mathbf{e}_{i,A}}$$
$$(1 \times 1) = (1 \times K)(K \times 1)$$

## **Square Prediction Error**

$$\mathbf{e}'_{i,A} = \mathbf{x}'_i - \widehat{\mathbf{x}}'_{i,A}$$

▶ 
$$\mathsf{SPE}_i = \sqrt{e_{i,1}^2 + e_{i,2}^2 + \ldots + e_{i,K}^2}$$

- ▶ Smallest SPE:  $SPE_i = 0$
- ► Calculate 95% or 99% confidence limit



## **Square Prediction Error**

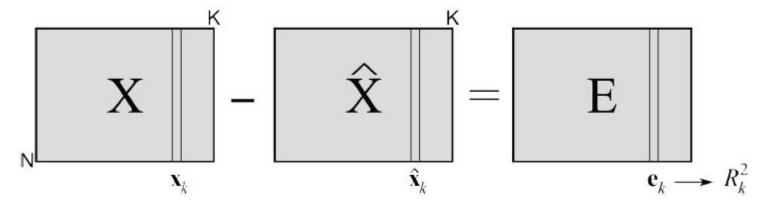
Distance from each observation to the model's plane:

- ▶ Does model explain that point well? SPE=0
- ▶ If SPE > 95% limit:
  - poorly explained by the model
  - something new in this observation
  - new phenomenon?

## **Square Prediction Error**

## Column Residual

- ▶ SPE is the row residual for X
- Residuals also calculated for each column



▶ How well each column is explained by the model

### Column Residual

- $R_k^2 = \frac{\operatorname{Var}(\widehat{\mathbf{x}}_k)}{\operatorname{Var}(\mathbf{x}_k)}$
- ▶ The  $R_k^2$  value:
  - ▶ is 0.0 when there are no components
  - increases for every every component added

### Whole Matrix Residual

$$\mathbf{X} = \mathbf{TP}' + \mathbf{E} = \widehat{\mathbf{X}} + \mathbf{E}$$

▶ How well does the model fit the data?

$$R^2 = \frac{\mathsf{Var}(\widehat{\mathbf{X}})}{\mathsf{Var}(\mathbf{X})}$$

- $ightharpoonup R^2 = 0.0$  when there are no components
- $ightharpoonup R^2$  increases with every component added

$$R_{a=0}^2 > R_{a=1}^2 > R_{a=2}^2 > \dots > R_{a=A}^2 = 1.0$$

#### More about direction vectors

- "Direction vectors" = "Loadings"
- ▶ Link between the real-world and the latent-variable world

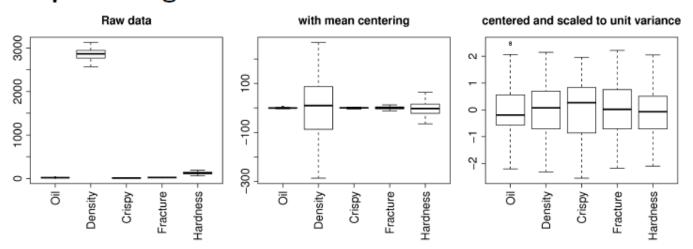
$$T = XP$$
  
 $(N \times A) = (N \times K)(K \times A)$ 

★ Statistically, loading vectors are eigenvectors of X<sup>T</sup>X.

Then how about eigenvalues?

## Preprocessing

#### Pre-processing the data: center and scale



▶ Centering:  $\mathbf{x}_{k,\text{center}} = \mathbf{x}_{k,\text{raw}} - \text{mean}(\mathbf{x}_{k,\text{raw}})$ 

Remember

► Scaling:  $\mathbf{x}_k = \frac{\mathbf{x}_{k,\text{center}}}{\text{standard deviation}(\mathbf{x}_{k,\text{center}})}$ 

$$Z = \frac{X - \mu}{\sigma}$$
 ?

▶ Does not change relationships between variables.

## More on preprocessing Data

- Modifies the columns of X before building the model
- Center
- Scale
- Add transformations:
  - use log(T) instead of temperature, T
  - ▶ use 1/P instead of pressure, P
  - use sqrt(F) instead of flow, F

#### Add extra columns to X:

- heat balance
- dimensionless numbers
- ▶ square terms:  $x_1^2$ ,  $x_2^2$ ,...
- ▶ interaction terms:  $x_1x_2$ ,  $x_1x_3$ ,  $x_2x_3$  ...

#### How is PCA calculated?

- ► Eigenvalue decomposition
  - loadings are the eigenvectors of X'X.
  - once you have the eigenvectors, then T = XP
  - eigenvalues are the variances of the scores,  $s_a^2$
- ► Singular value decomposition
  - $\mathbf{X} = \mathbf{U}\mathbf{\Sigma}\mathbf{V}' = \mathbf{T}\mathbf{P}'$
  - ightharpoonup scores,  $T = U\Sigma$  and the loadings, P = V

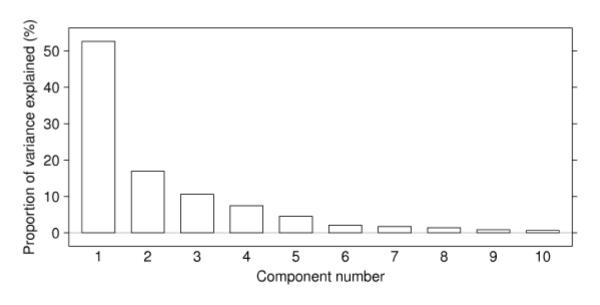
#### How is PCA calculated?

- Non-linear iterative partial least-squares (NIPALS) algorithm
  - One component at a time
  - Handles missing data
  - Iterative; it always converges, but slow sometimes
  - Also called the Power algorithm
  - Excellent on large data sets
  - Google used this algorithm for their first search engine (called PageRank)

More details next.

# How many components?

- ▶ Eigenvalues
  - sum of the eigenvalues =  $\sum_{a=K}^{a=K} \lambda_a = K$
  - keep adding components as long as \( \lambda\_a > 1 \)
- ▶ Plot  $R^2$  for each component

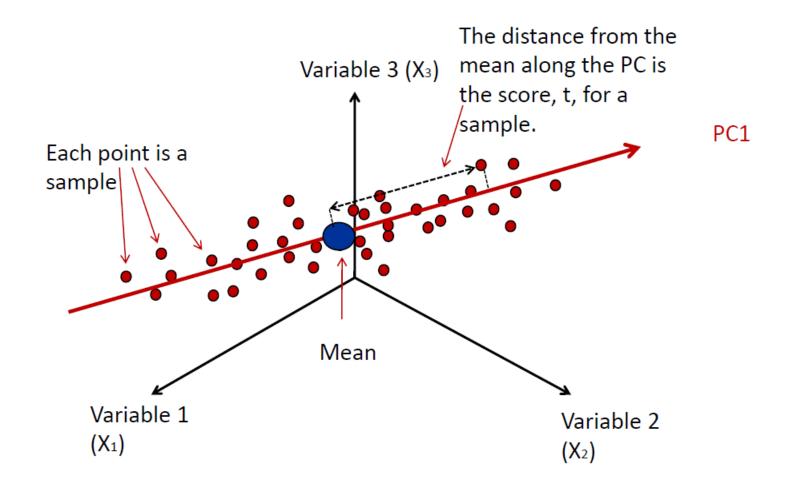


▶ Use cross-validation

Called scree plot

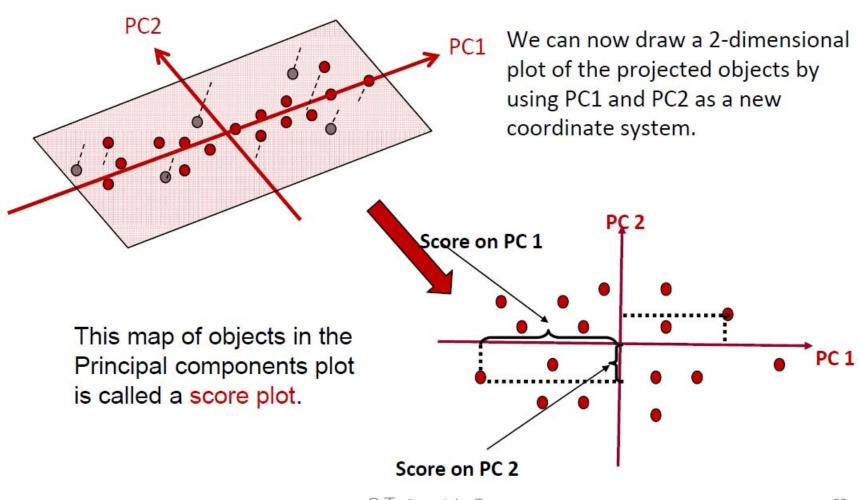
## Review of PCA

• What is score?



#### Review of PCA

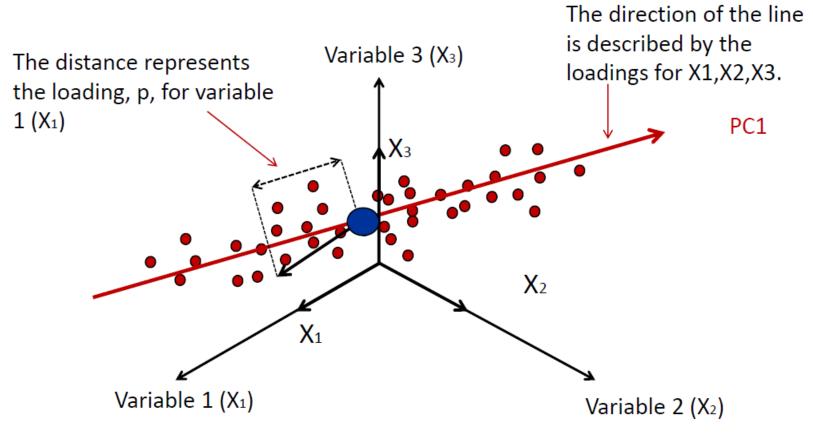
Score plot – low dimensional summary of samples



### Review of PCA

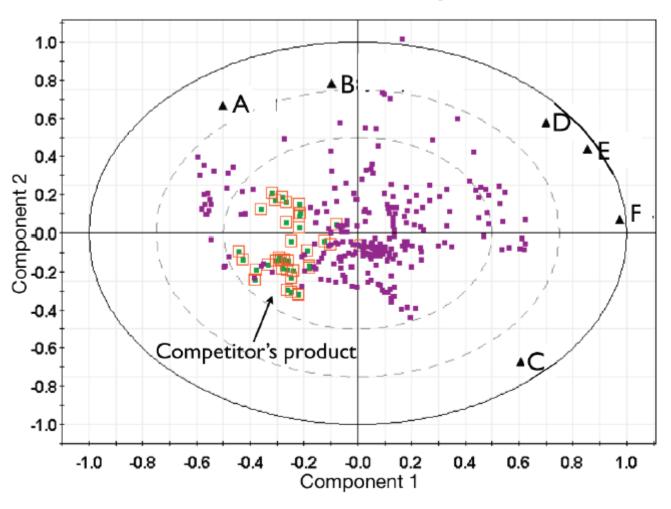
## What is loading?

Coefficients in the linear combination PC1 =  $a_1 X1 + a_2 X2 + a_3 X3$ 



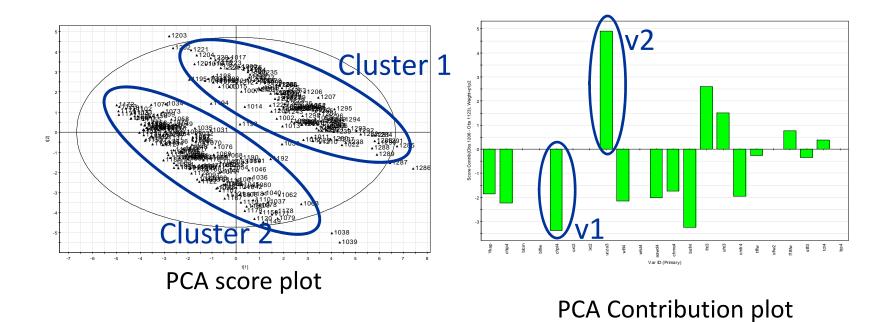
## Use of PCA

Improved Process Understanding



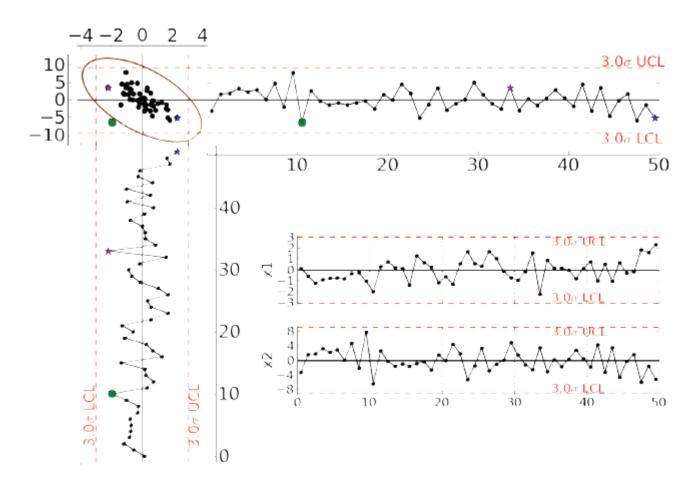
## Use of PCA

Troubleshooting Process Problem



## Use of PCA

Multivariate Stastiscal Process Control (MSPC)



### In the next lecture

- Tutorials
- NIPALS algorithm
- A bit more on PCA
- Assignment #1