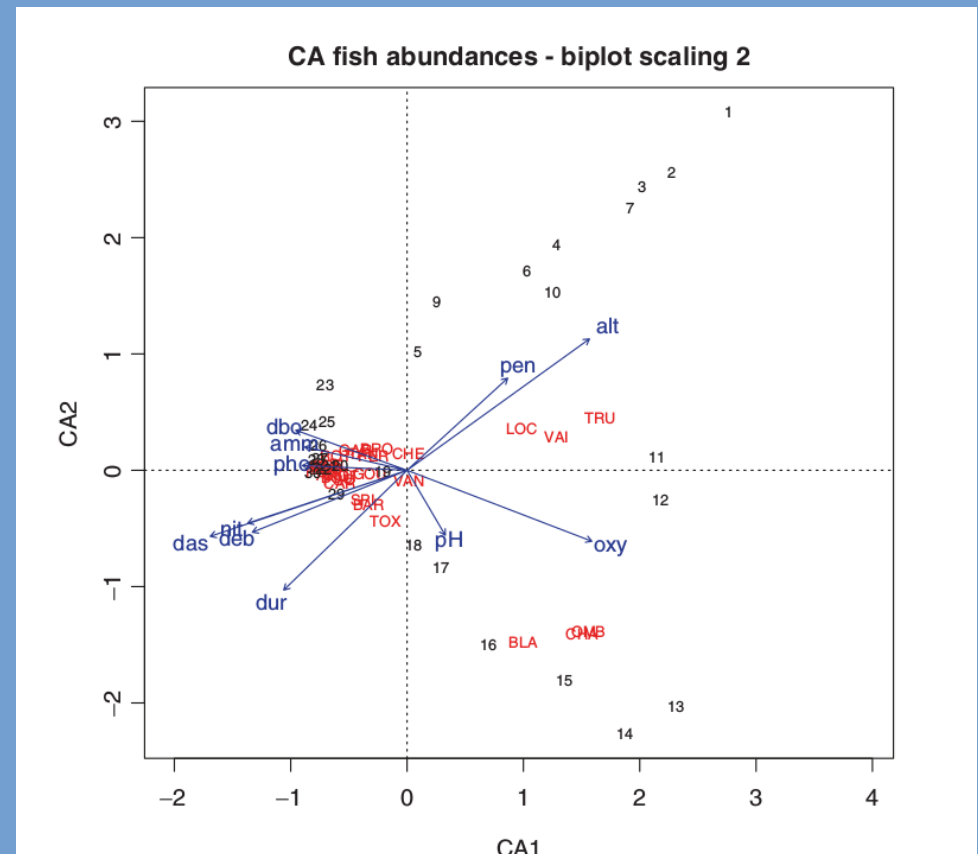
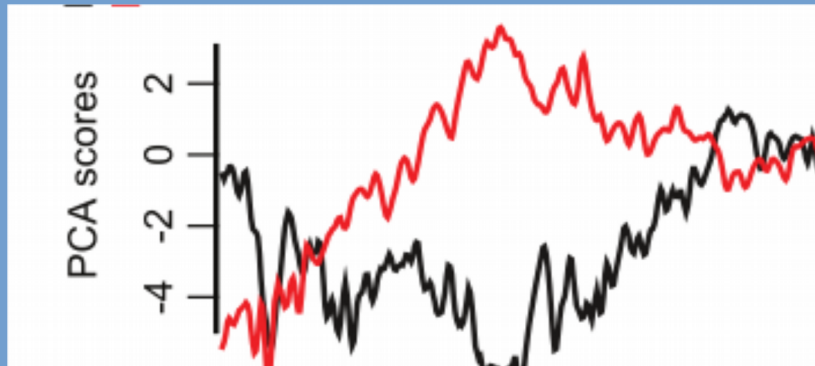


# Introduction to ordination: principal components analysis and related methods



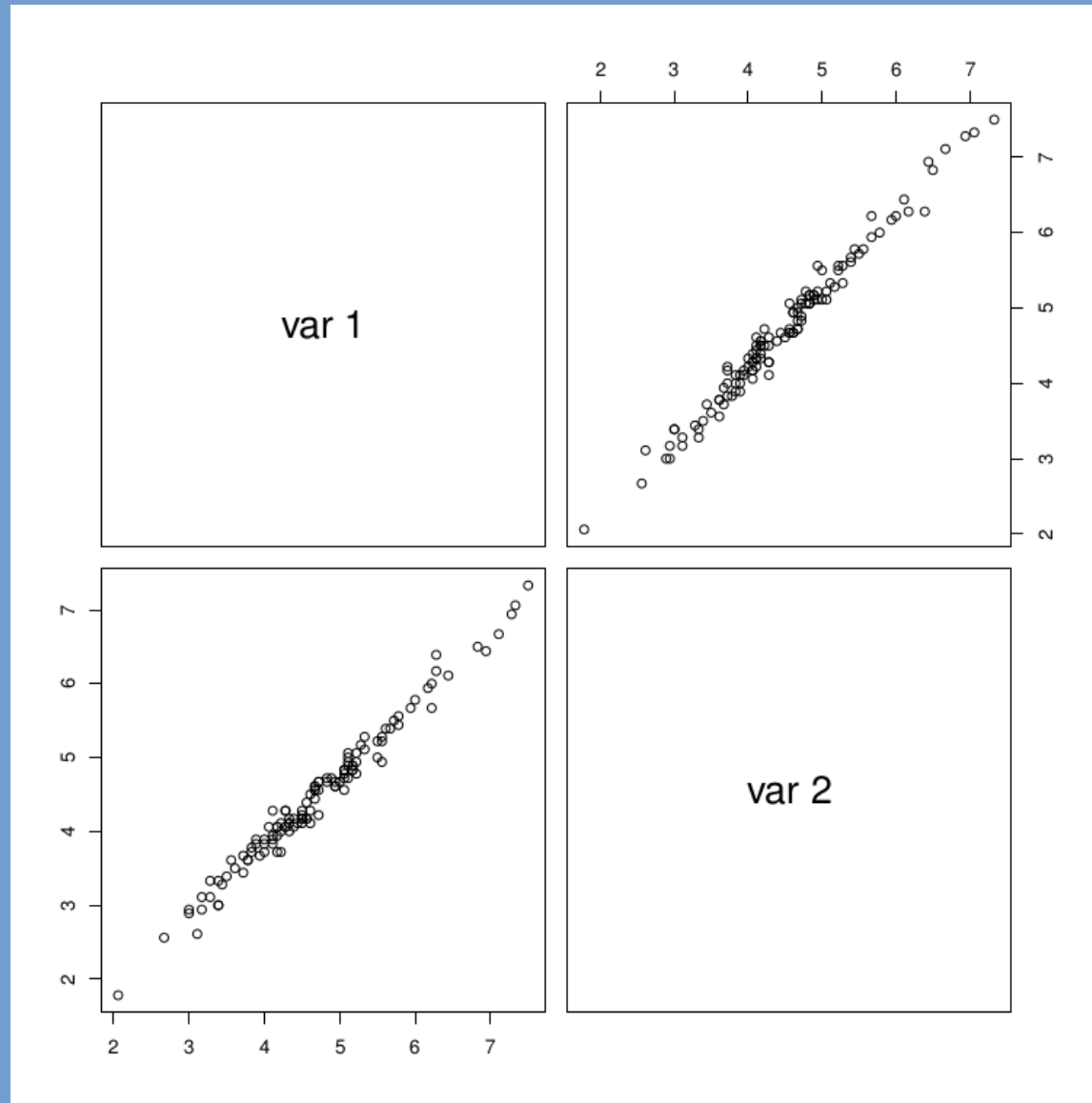
# Aims of ordinations

- Simplify complex datasets
  - Find most relevant patterns in a dataset
  - Summarize a dataset in a simple 2-dimensional scatterplot
    - Relation among variables
    - Distance among samples

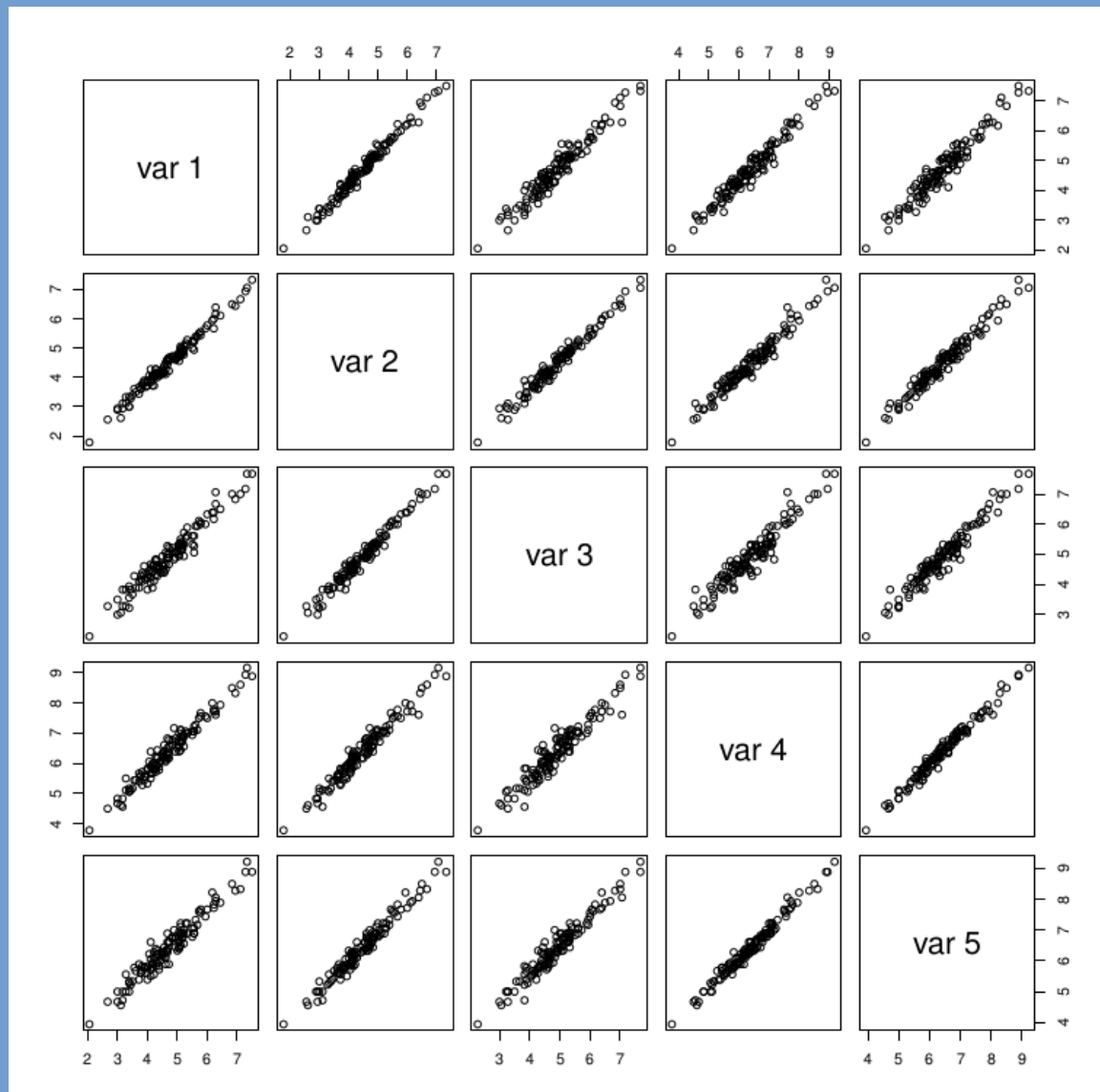
# Simplify a dataset

- Dataset with 9 variables
  - Methods to find main patterns:
    - Visually?

# Temperatures in Wisconsin

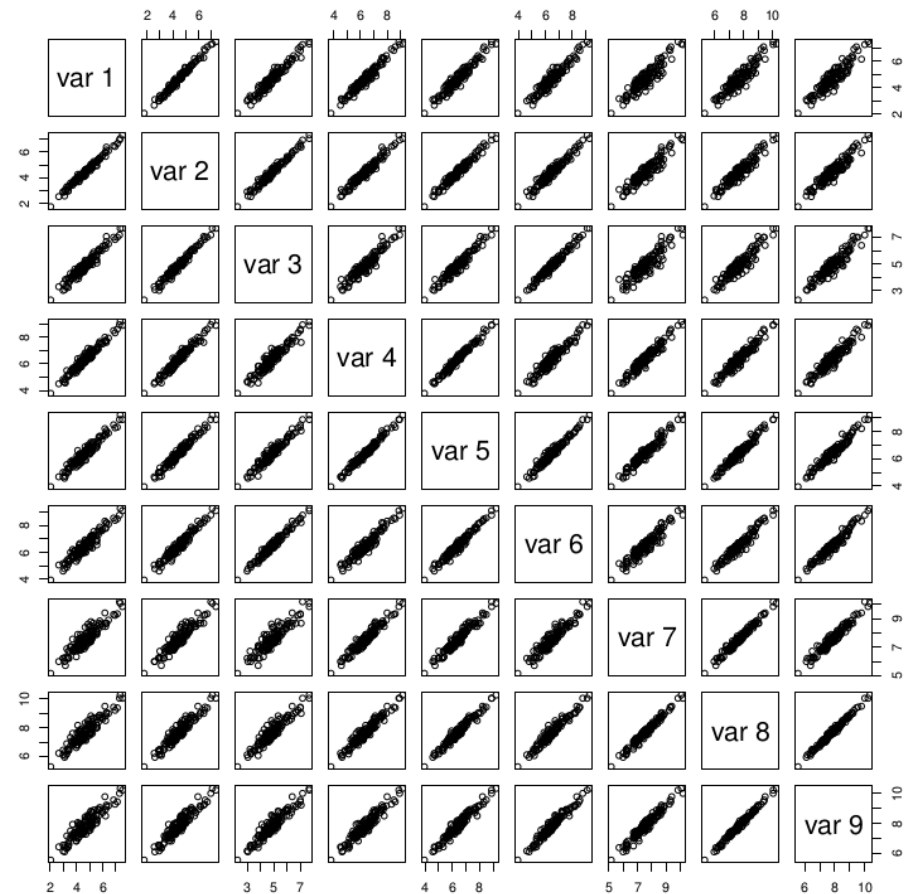


# Temperatures in Wisconsin



# Simplify a dataset

- Dataset with many variables
  - Methods to find main patterns:
    - Visually?
    - Numerically?

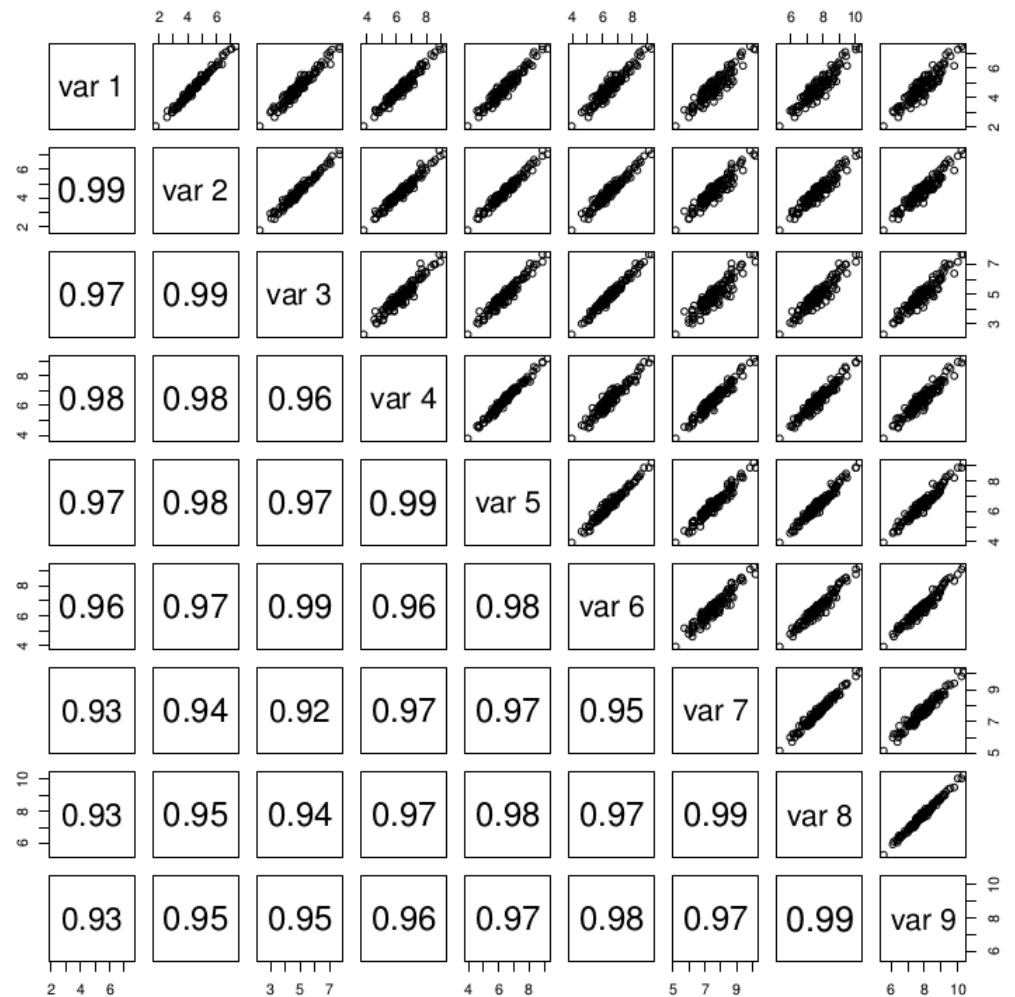


# Simplify a dataset

Methods to find main patterns:

Numerically?

Covariance or correlation matrix

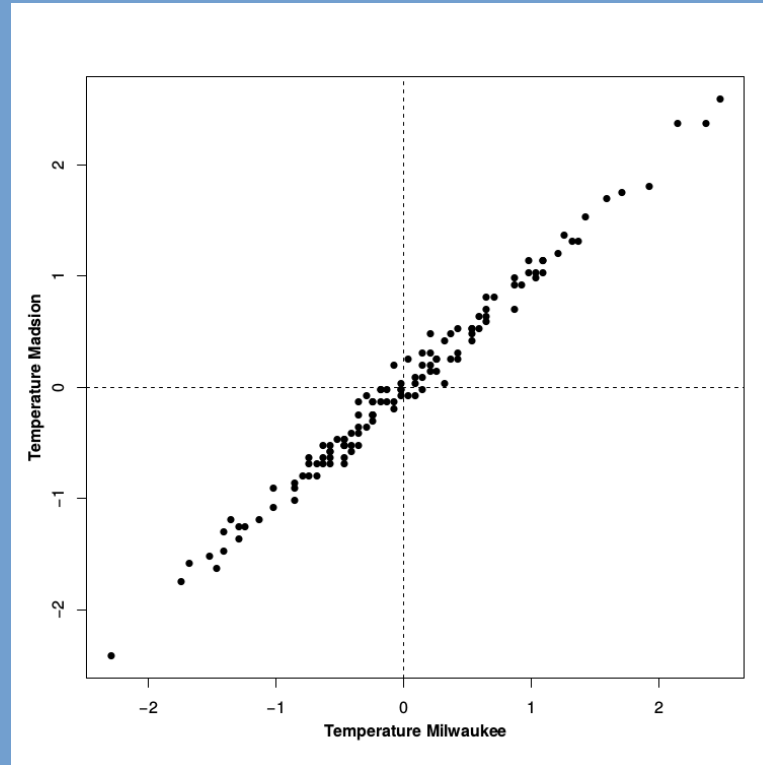


# Simplify a dataset

- Dataset with 20 variables
  - Methods to find main patterns:
    - Visually?
      - Show a scatterplot of all variables
    - Numerically?
      - Covariance or correlation matrix

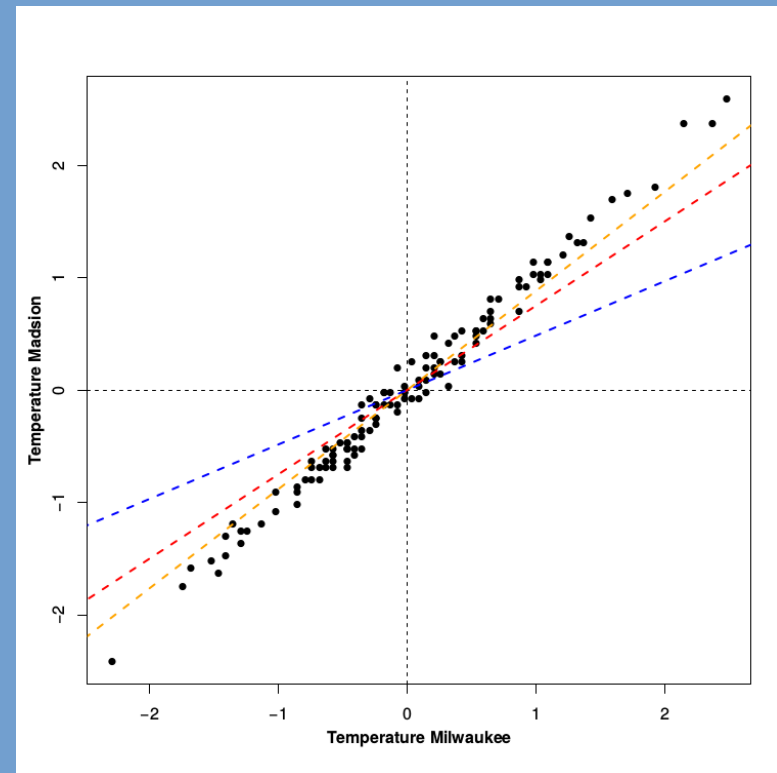
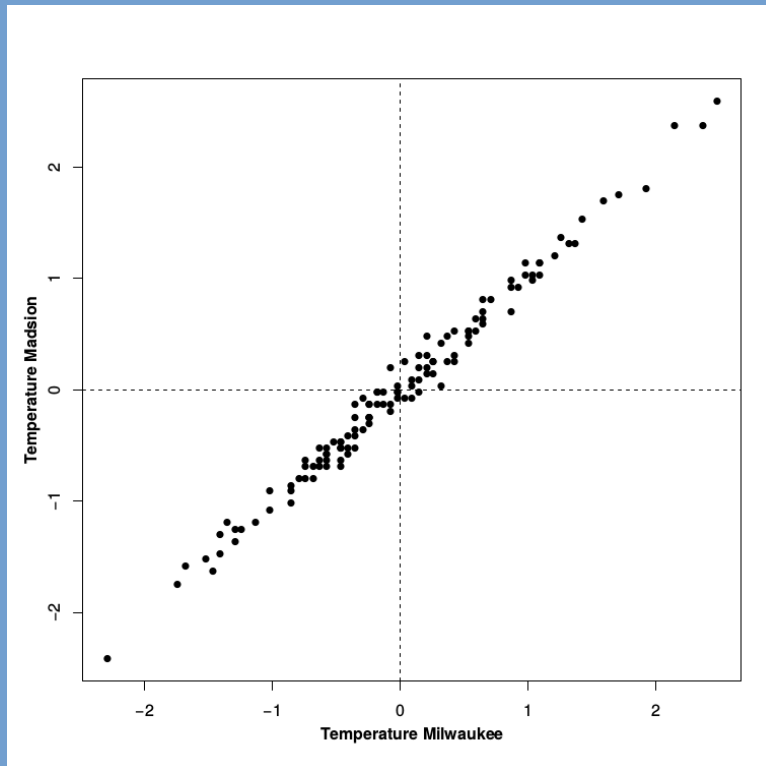


# Two dimensional example



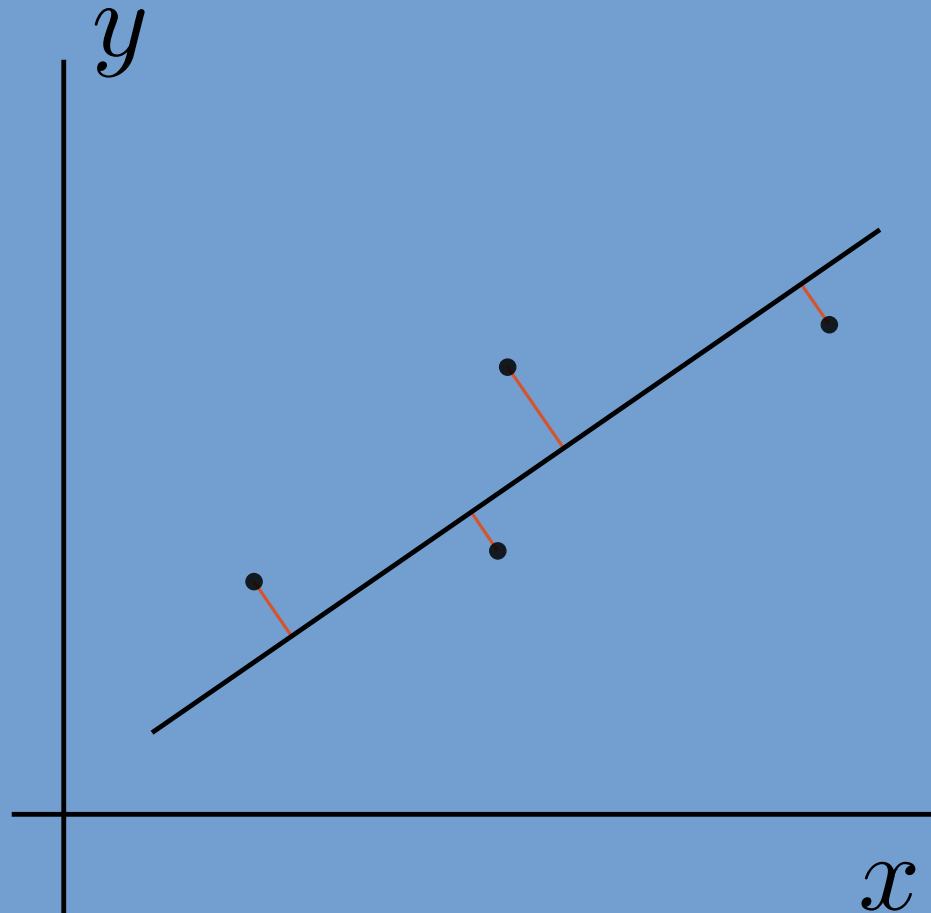
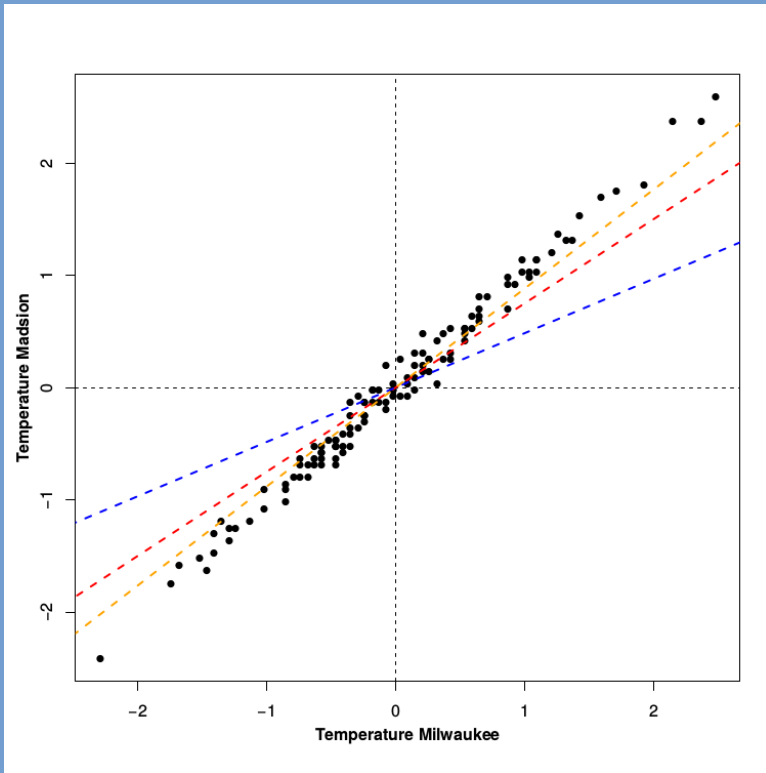
- Data represented by a cloud of points in variable space
- Usually our data has structure = correlation between variables = cloud of points is elongated

# Geometric representation



- Draw a line through the direction of maximum elongation that also passes through the center of the swarm:
  - First principal component is the line that explains most variance
  - Minimizes the distance to the observations
  - Does this remind you of another mathematical operation?

# PCA as extension of regression

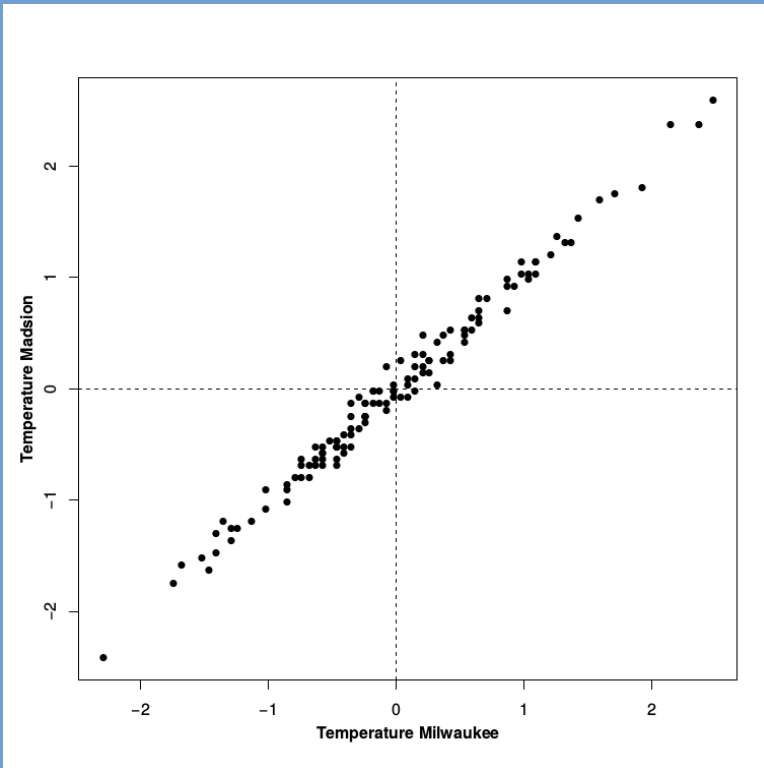


Type II regression: minimizing distance perpendicular to the regression line

# Formal solution

- Eigenvalue decomposition of **correlation or covariance matrix**
  - Eigenvalue decomposition (a little magic, no need to know what that is, *eigen* in R)
  - **Eigenvalue** indicating how important a component is (as many eigenvalues as variables)
  - **Eigenvector** describing the new coordinate system (as many eigenvectors as variables)

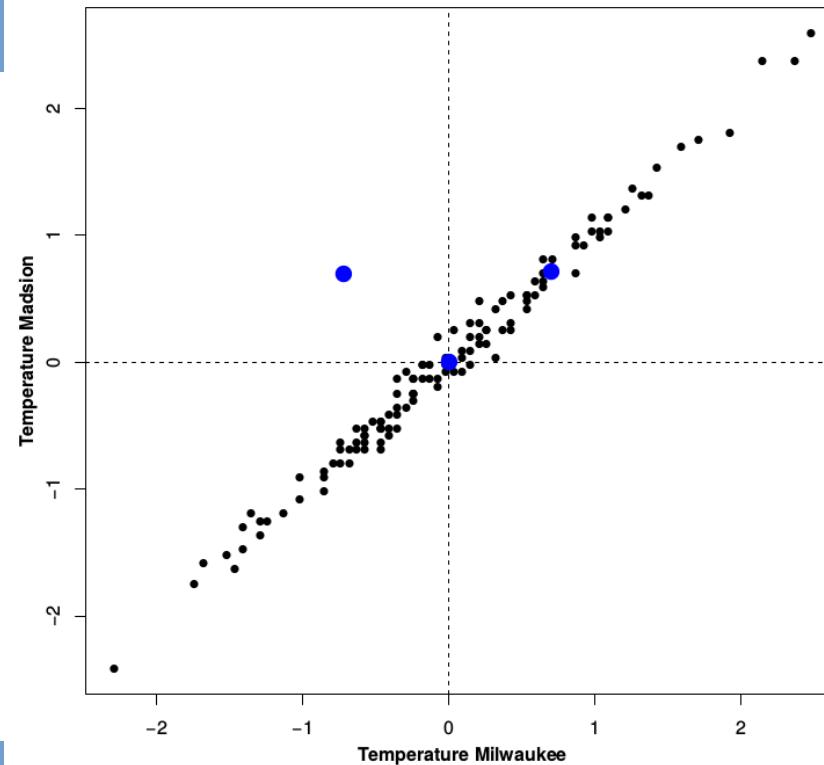
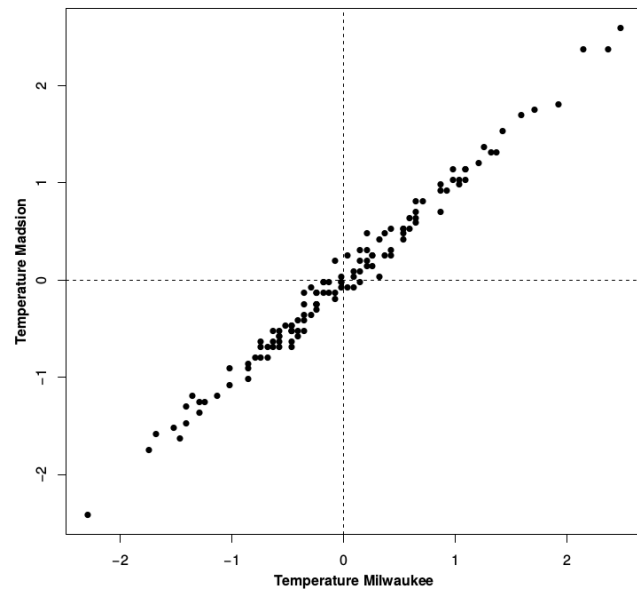
# Eigenvalue decomposition 2 dimensions



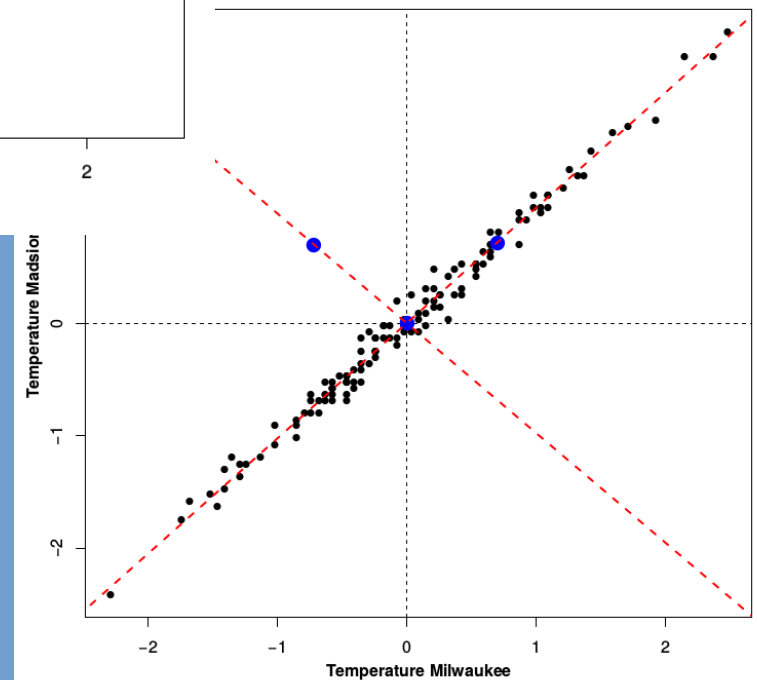
- Eigenvalue (importance):
  - Value 1: 1.985 (99.25%)
  - Value 2: 0.015 (0.75%)
- Eigenvectors (new coordinate system)

|           | Vector 1 | Vector 2 |
|-----------|----------|----------|
| Milwaukee | 0.698    | -0.715   |
| Madison   | 0.715    | 0.698    |

# Two dimensional example



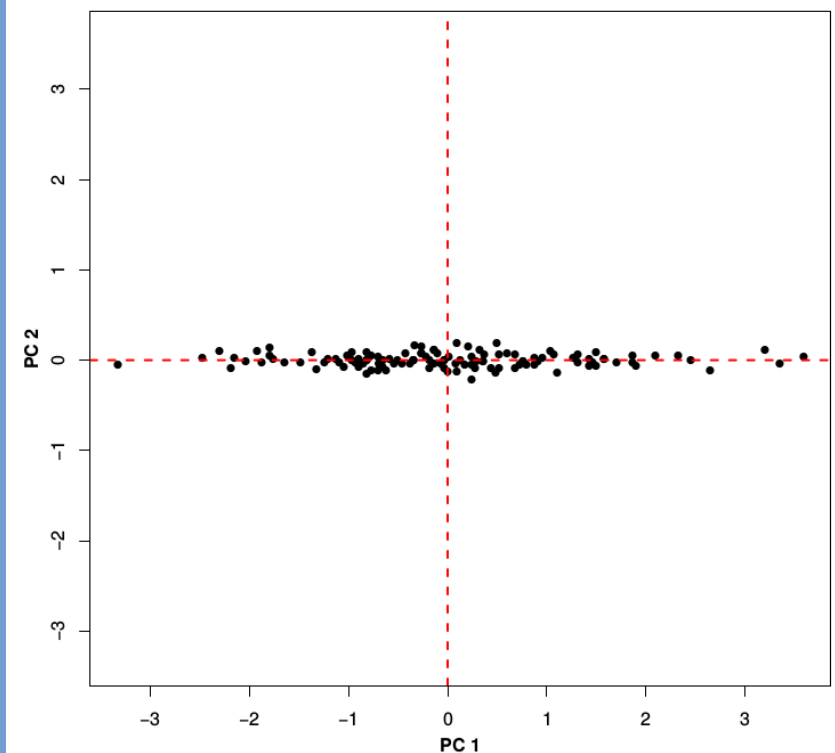
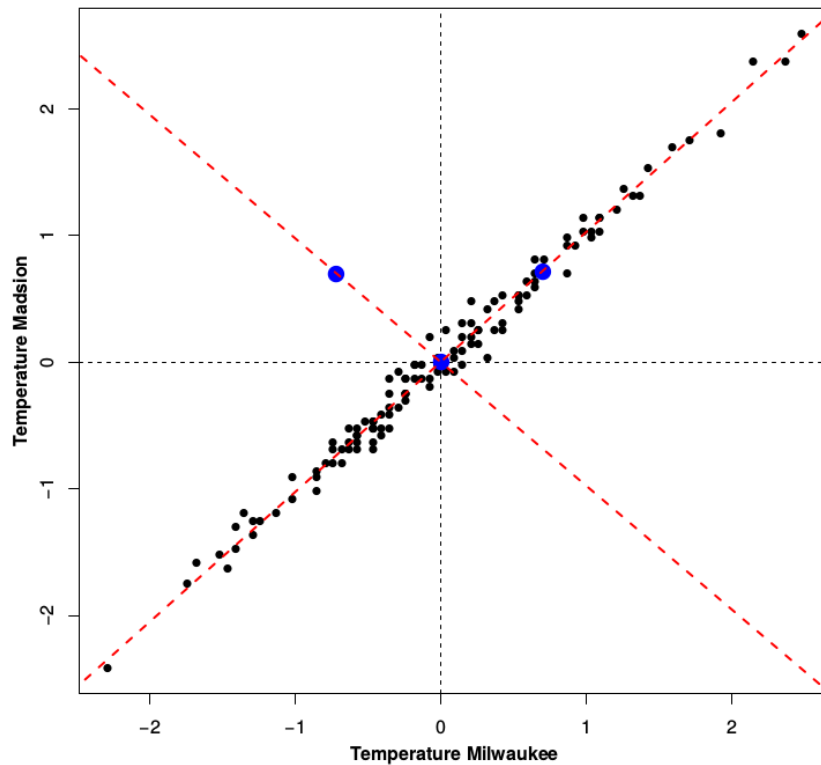
|           | Vector 1 | Vector 2 |
|-----------|----------|----------|
| Milwaukee | 0.698    | -0.715   |
| Madison   | 0.715    | 0.698    |



# PC1 and PC2

- $PC1 = (Vec1_{Mil} * T_{Mil} + Vec1_{Ma} * T_{Ma})$
- $PC2 = (Vec2_{Mil} * T_{Mil} + Vec2_{Ma} * T_{Ma})$

|           | Vector 1 | Vector 2 |
|-----------|----------|----------|
| Milwaukee | 0.698    | -0.715   |
| Madison   | 0.715    | 0.698    |



# PC1 and PC2

- $PC1 = (Vec1_{Mil} * T_{Mil} + Vec1_{Ma} * T_{Ma})$
- $PC2 = (Vec2_{Mil} * T_{Mil} + Vec2_{Ma} * T_{ma})$

|           | Vector 1 | Vector 2 |
|-----------|----------|----------|
| Milwaukee | 0.698    | -0.715   |
| Madison   | 0.715    | 0.698    |

- PCs are linear combinations of initial variables with eigenvectors as coefficients: **PC Scores** (site score)
- Eigenvectors also called **PC loadings** (species scores)



# Variance explained

- Variance PC1 = Eigenvalue 1 =  $\lambda_1$
- Variance PC2 = Eigenvalue 2 =  $\lambda_2$

Variance PC1 = 1.539019 Eigenvalue 1 = 1.539018890

Variance PC2 = 0.005447696 Eigenvalue 2 = 0.005447696

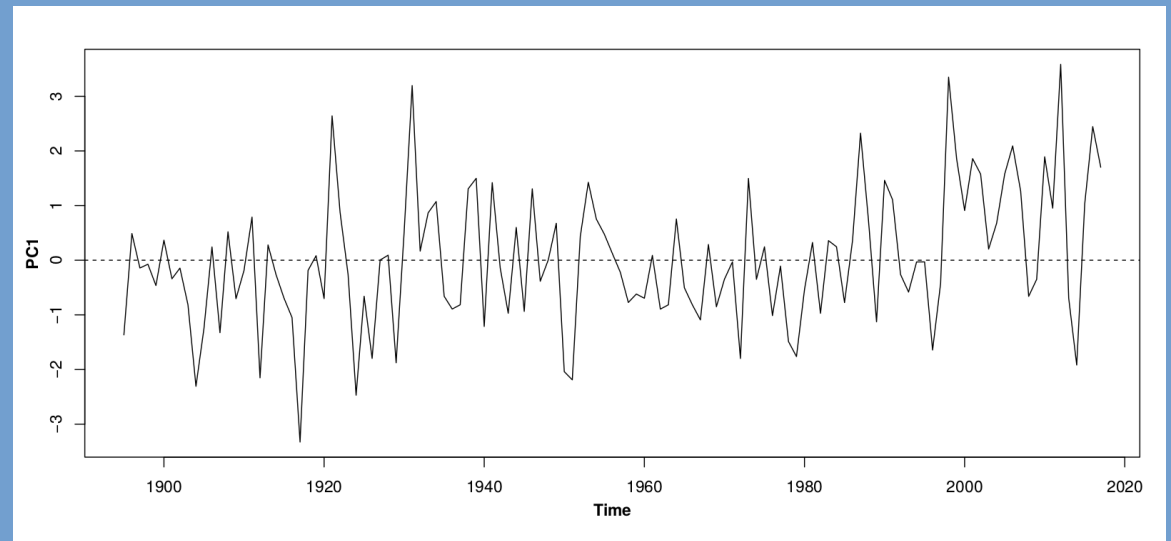
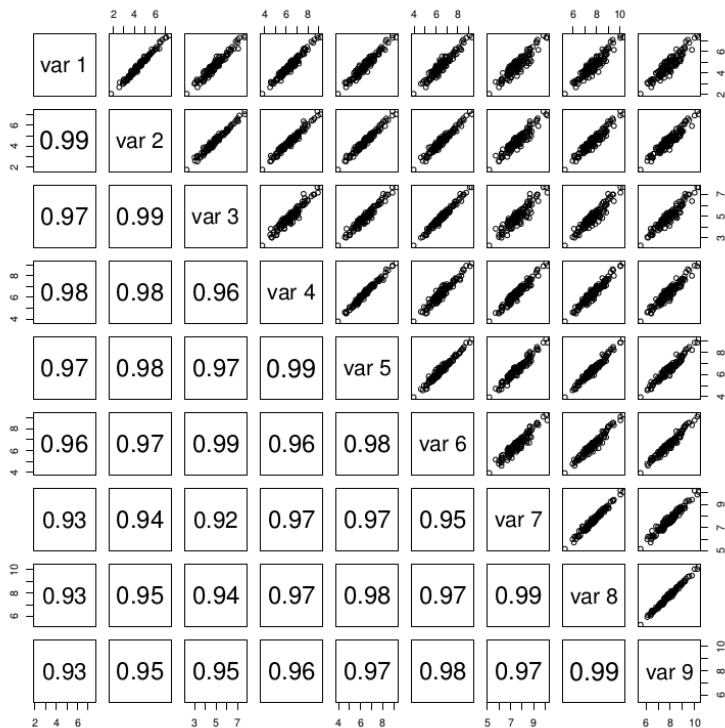
- Total Variance = Var TMil + Var TMad = Var PC1 + Var PC2 = Eig val 1 + Eig val 2
- **Variance explained** = Eig val 1 (Var PC1) / Total Variance

Example:  $1.539/1.544 = 0.996$

**PC1 explains 99.6% of the total variance**

# Summary

- Simplify dataset:
  - Correlation or covariance matrix
  - A little magic



# High dimensional dataset

- Same methodology
- No longer possible to visualize

# Visualization of PCA results

- PCA Biplot:
  - Joint representation of variables and observations (samples, sites) on the same diagram

Usual to represent observations/samples/sites by points and variables by arrows

Arrows point in direction of maximum rate of change of that variable across the diagram

Length of arrows indicate relative rate of change in that direction

# Example: Climate data Minnesota

- North American Modern Pollen Database (NAMPD)
- Modern climate data at sites with pollen data
- Choose:
  - P Ann
  - P Nov
  - P Dec
  - T ave
- PCA with correlation matrix (standardized variables)

# Correlation matrix

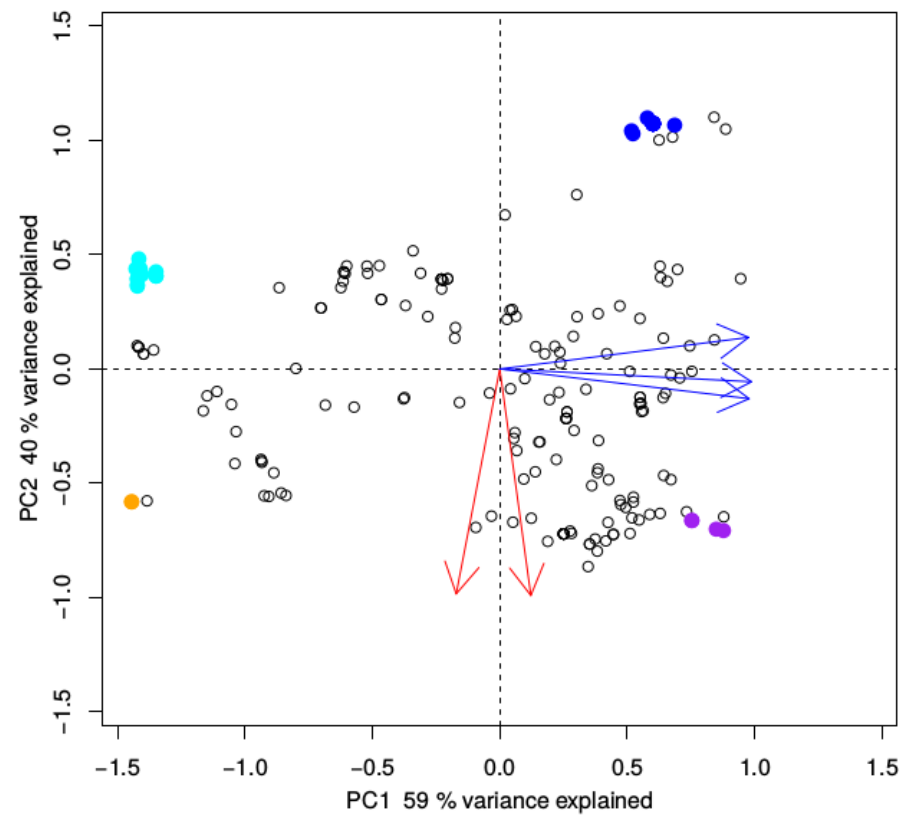
Eigenvector and eigenvalues depend on correlation of variables among each other

|      | tave  | tjul  | annp  | pnov  | pdec  |
|------|-------|-------|-------|-------|-------|
| tave | 1     | 0.95  | 0.24  | 0.18  | -0.01 |
| tjul | 0.95  | 1     | -0.04 | -0.11 | -0.3  |
| annp | 0.24  | -0.04 | 1     | 0.96  | 0.92  |
| pnov | 0.18  | -0.11 | 0.96  | 1     | 0.94  |
| pdec | -0.01 | -0.3  | 0.92  | 0.94  | 1     |

- Two highly correlated groups of variables
- Two groups are unrelated
- Probably two important PCs

# Biplot Example

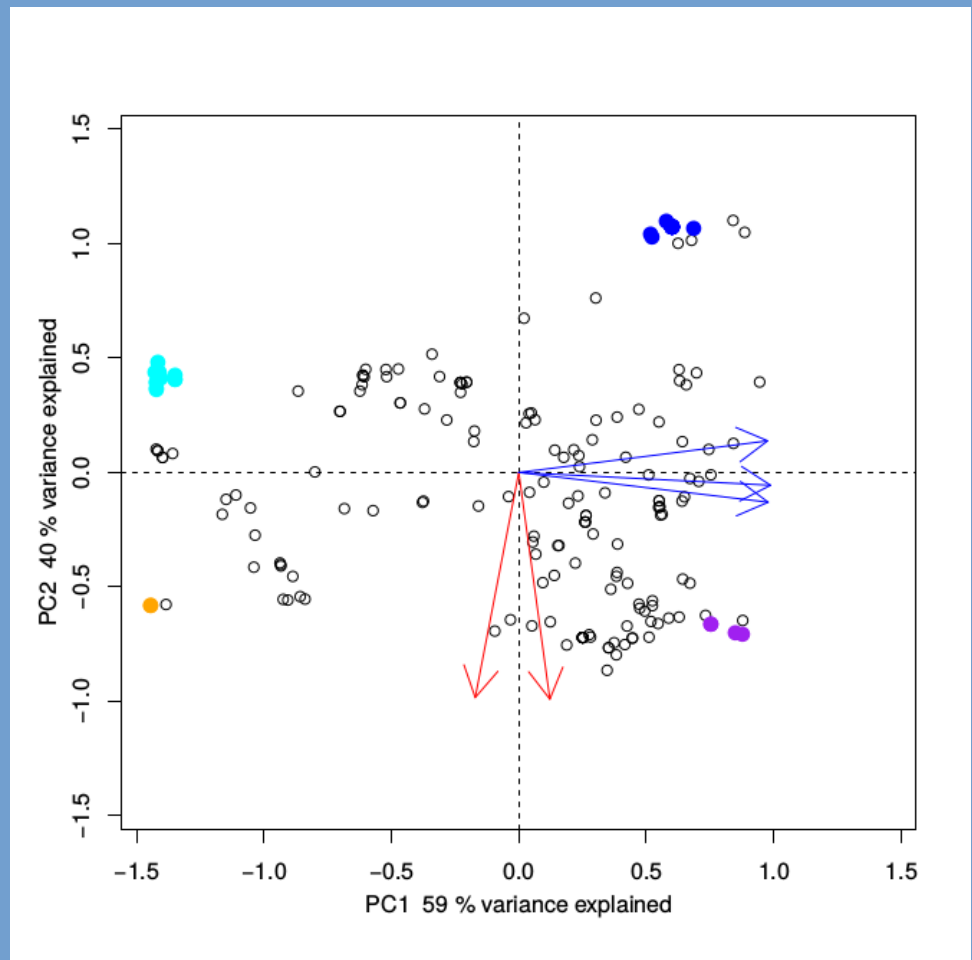
- Dimension reduction 5 to 2



# Biplot Example

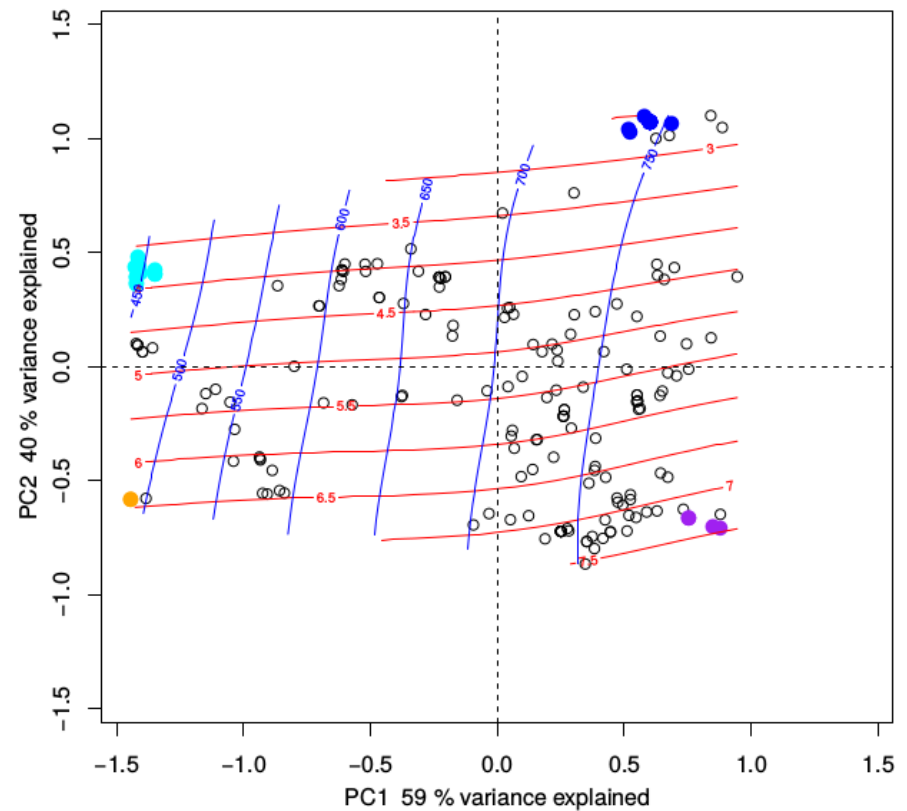
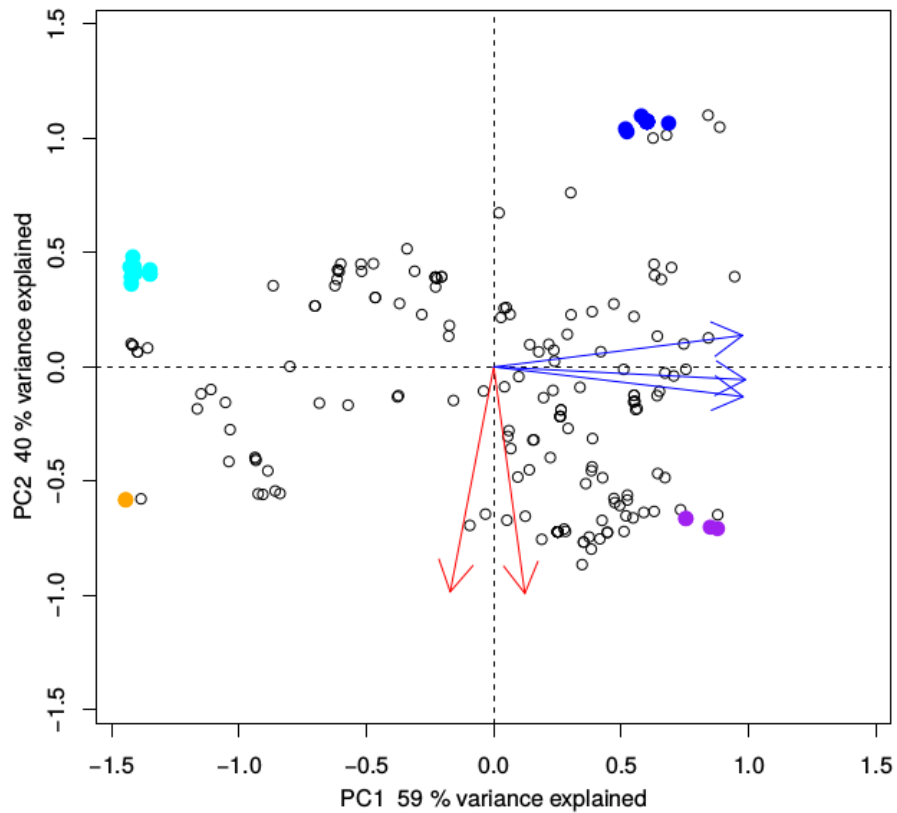
- Angle between arrows:
  - relation between variables
  - Same direction:
    - Positive relation
  - Orthogonal:
    - No relation
  - Opposite direction
    - Negative relation

Describe highlighted points

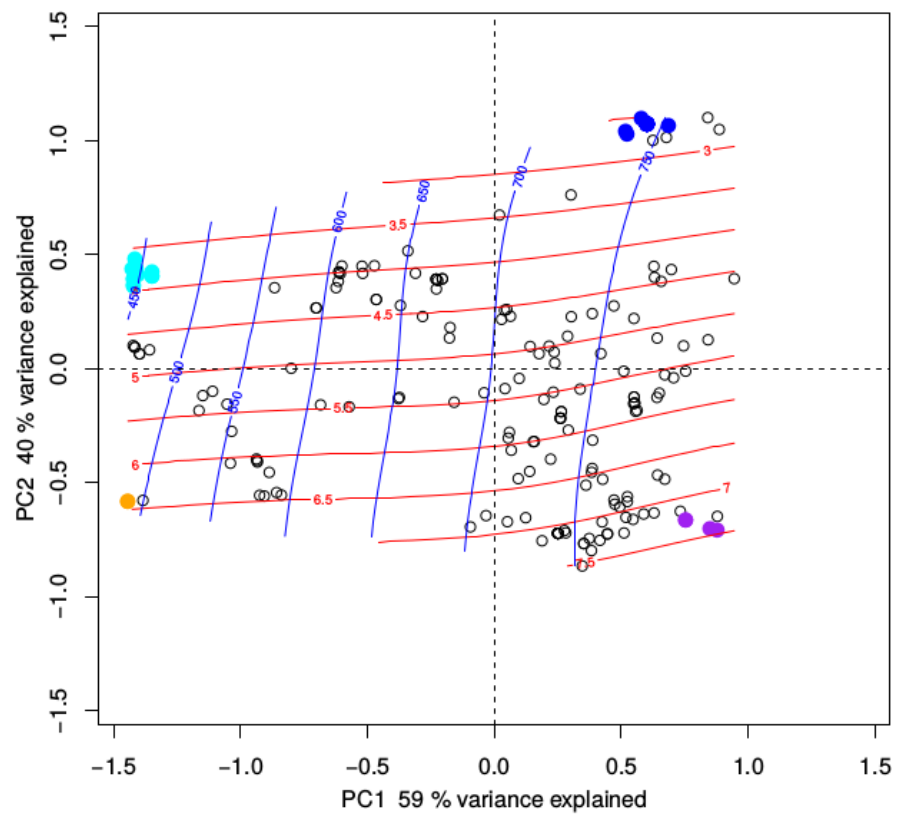




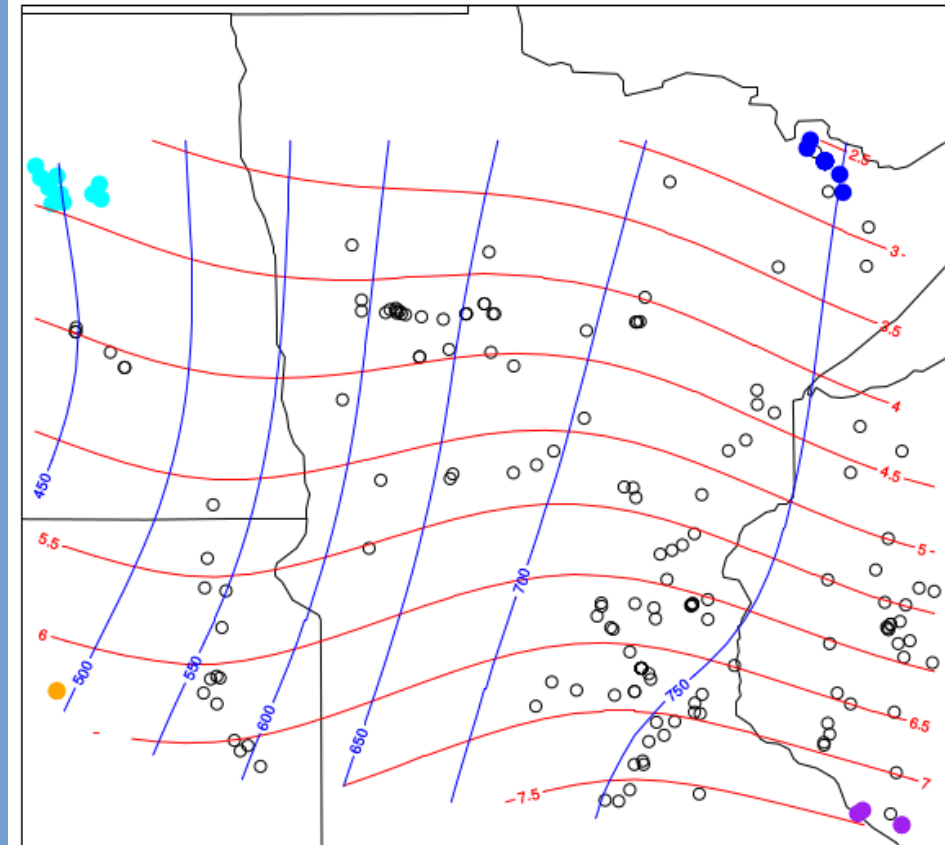
# Ordination surface



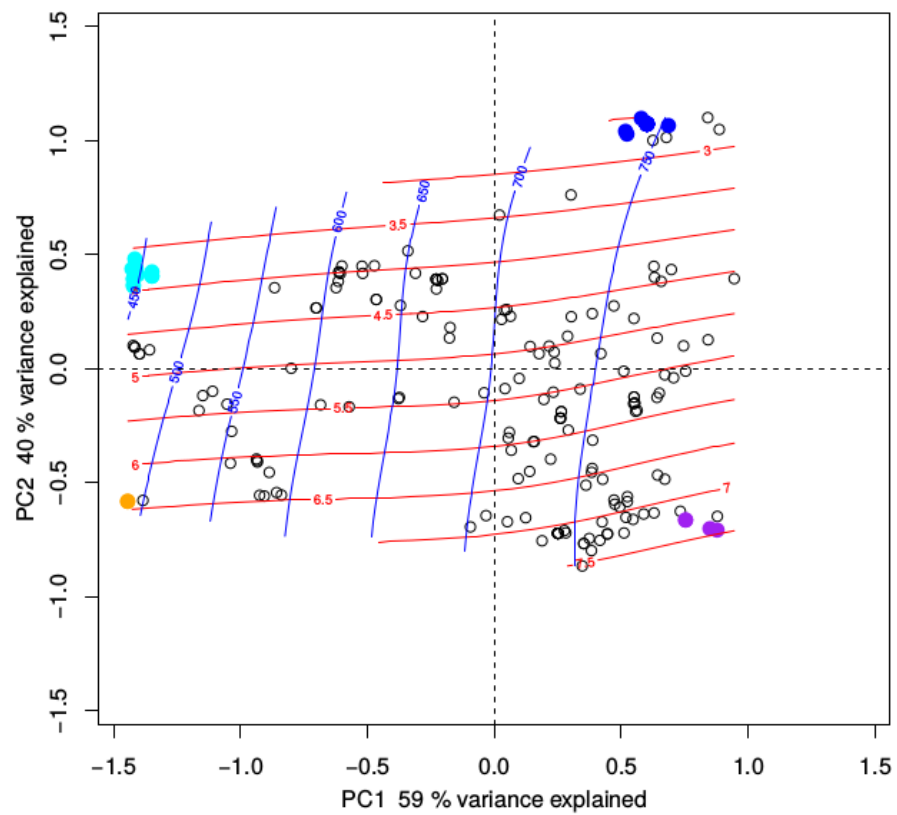
## Climate Space



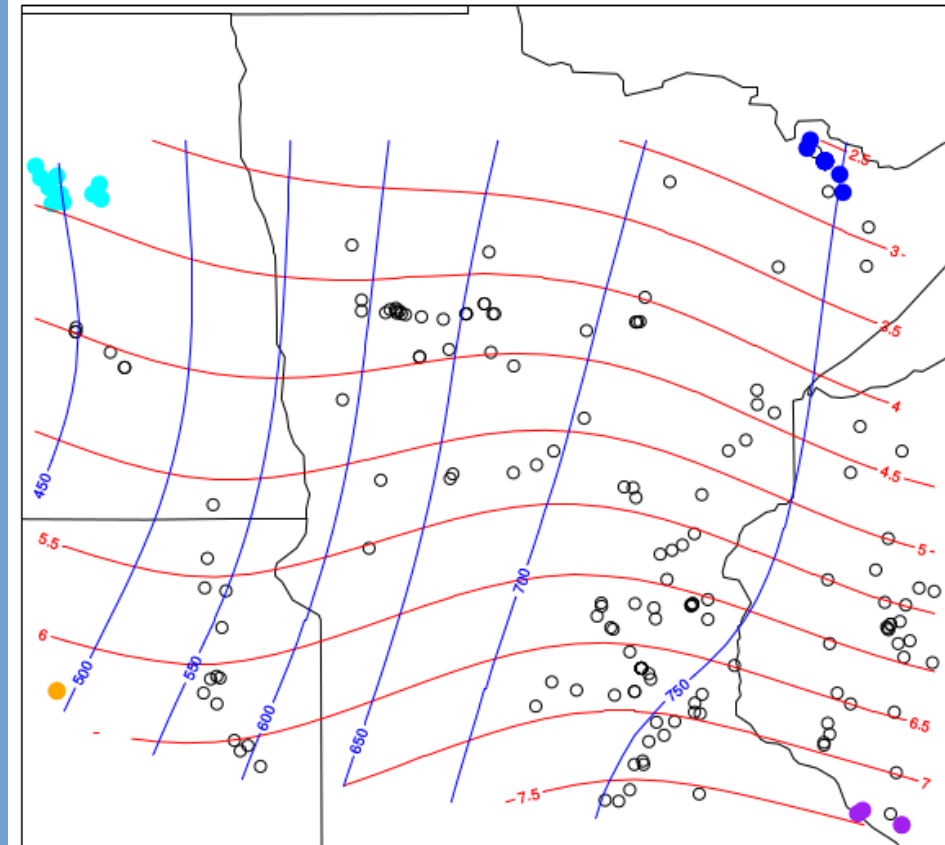
## Real Space



## Climate Space

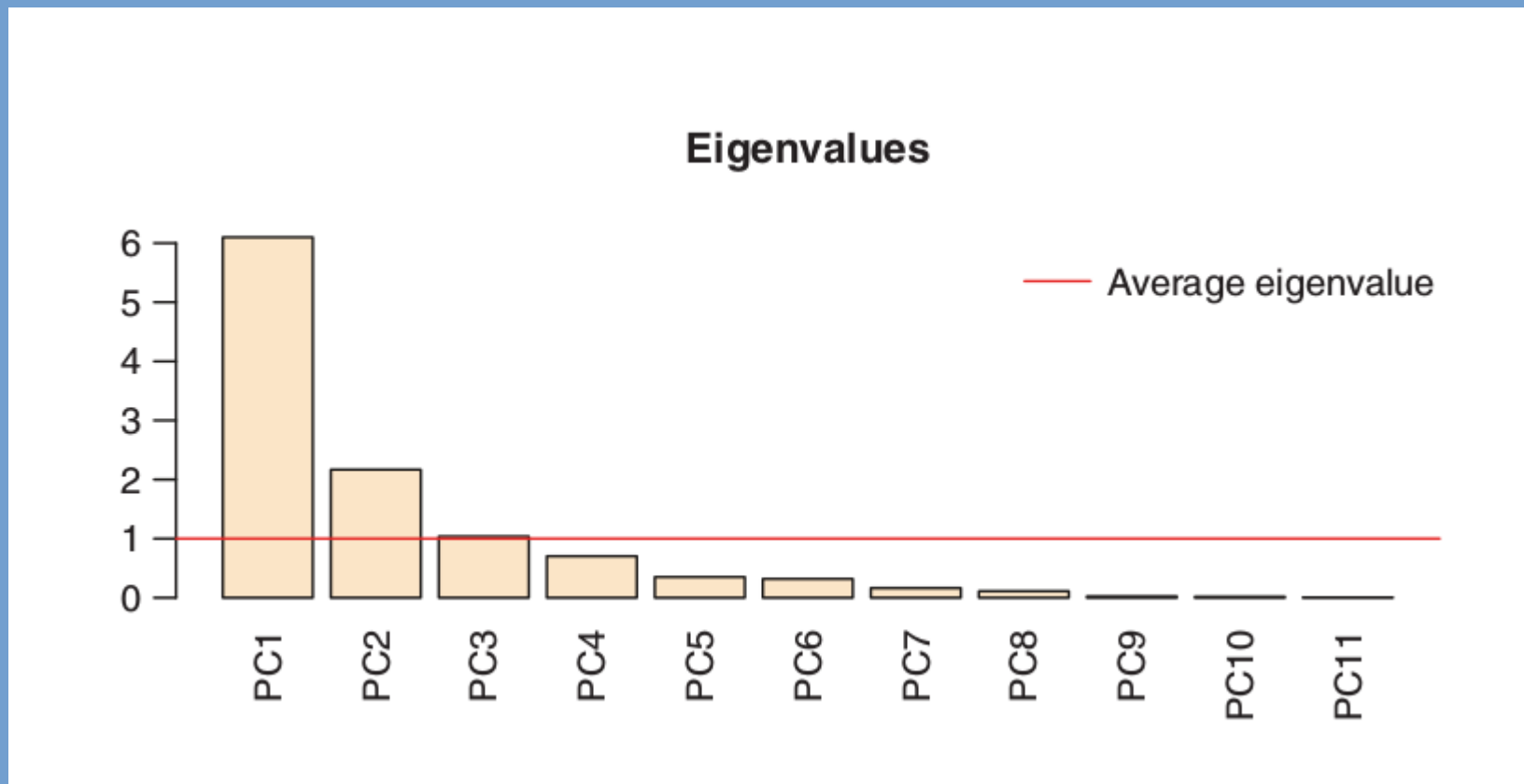


## Real Space

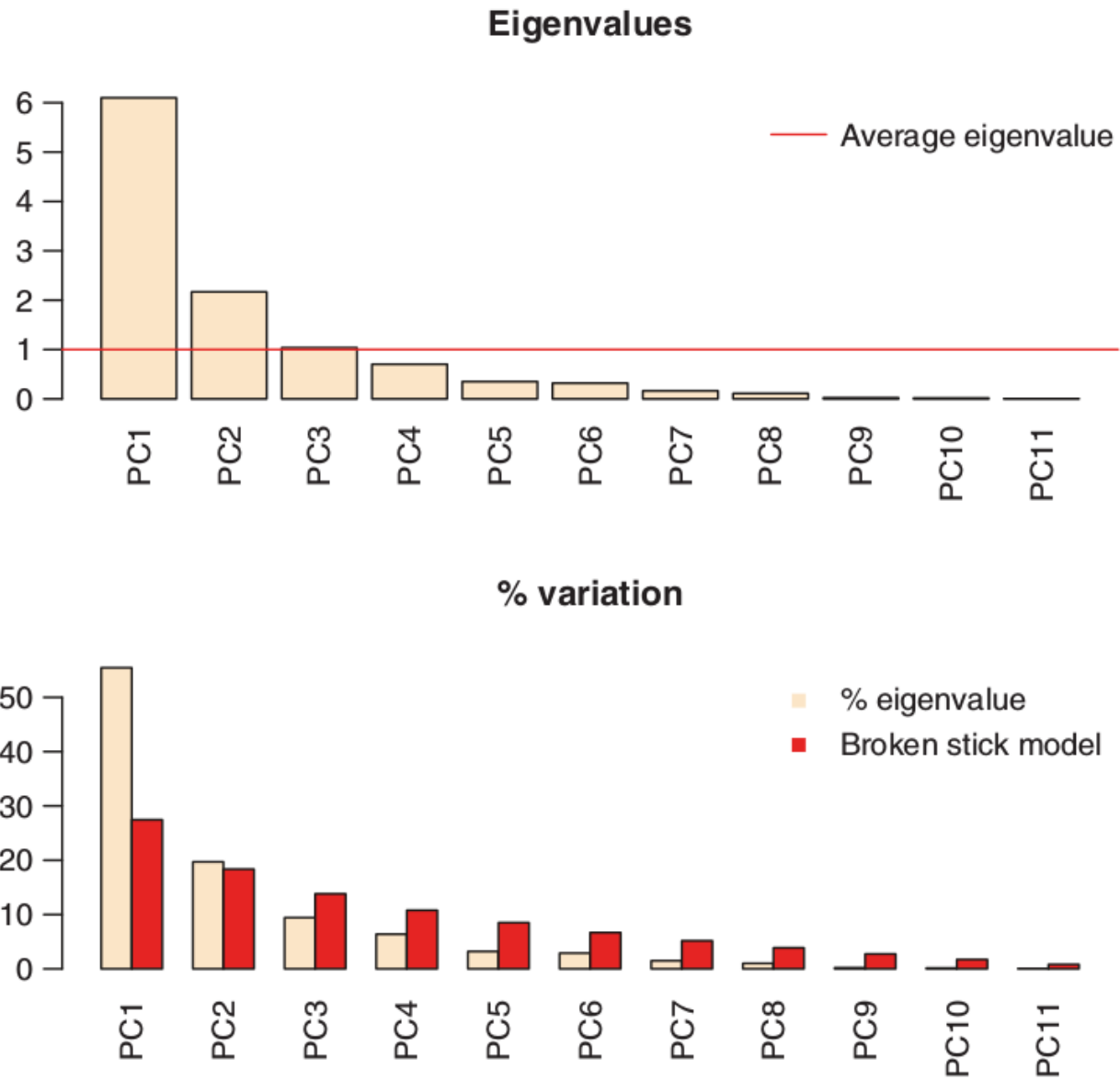


# How many axes?

- Screeplot: Histogram of variance explained by each PC axis



# How many axes?



# Standardized or non-standardized PCA?

- Principal components represent directions of maximum variation through multivariate space
- If original variables are measured on different scales variables with large absolute values will dominate simply because they represent directions of maximum variance
- To avoid this standardize data by centering and dividing by the standard deviation

# Standardized or non-standardized PCA?

- A PCA on the non-standardized data is equivalent to a PCA of a covariance matrix between variables
  - Also known as centered variables
- A PCA on the standardized data is equivalent to a PCA of a correlation matrix between variables

# Data transformations for PCA

- Usual to center data (subtract mean) – implicit in most software. Variables implicitly weighted by their variances
- For non-species data
  - Standardize data – variables have equal weight
- For species data
  - $\text{Log}_{10}(x+1)$  transformation useful for abundance (count) data
  - Square root transformation useful for percentage data



# Dealing with closed data

- Closed data: percentage or as a fraction of a fixed total
- Frequent in geochemical and microfossil studies
  - Log-ratio transformation
  - PCA: log transform and center by sample (site) and variable (species)
- Need to consider this for data with small number of variables (species) but usually makes little difference if number of variables is large