

# R Programming For Natural Resource Professionals

Week 14/15

Simulations: resampling/bootstrapping

# Learning objectives for this “week”

1. Understand use cases for multivariate statistics in natural resource management/research
2. Learn to run common multivariate analyses
3. Learn to interpret the results of common multivariate analyses

# What is multivariate statistics?

- We've been working with univariate stats
  - $\text{Response} \sim \text{predictor1} + \text{predictor2} \dots$
- Multivariate stats involve many response variables
  - Without predictors: Interdependence methods
  - With predictors: Dependence methods

# Interdependence methods

## **Dimension reducing methods**

- Principal component analysis
- Correspondence analysis
- Factor analysis

## **Cluster analysis**

- K-means clustering
- Hierarchical clustering

## **Multidimensional scaling**

- Non-metric multidimensional scaling

# Dependence methods

## **Discrimination and classification methods**

- Canonical variate analysis
- Neural networks
- Random forest

## **Constrained ordination**

- Canonical correlation analysis
- Redundancy analysis
- Hierarchical clustering

## **Multivariate regression**

# Principal component analysis (PCA)

Would take awhile to explain the math behind PCA..

- Linear algebra and eigensystems. Interested? Take a multivariate stats course!

Goal: simplify large and complex data sets

- “Reduce dimensionality”

All data can be perfectly summarized (all variance explained) if the number of dimensions used to explain it equals number of observations

- Not really in practice though

# Principal component analysis (PCA)

- Explaining data variation in “dimensions”
- Consider a data set with three variables:  $x$ ,  $y$ ,  $z$
- PCA calculates a covariation matrix
  - In what ways are  $x$ ,  $y$ , and  $z$  correlated/related?

$$\begin{bmatrix} Cov(x, x) & Cov(x, y) & Cov(x, z) \\ Cov(y, x) & Cov(y, y) & Cov(y, z) \\ Cov(z, x) & Cov(z, y) & Cov(z, z) \end{bmatrix}$$

# Principal component analysis (PCA)

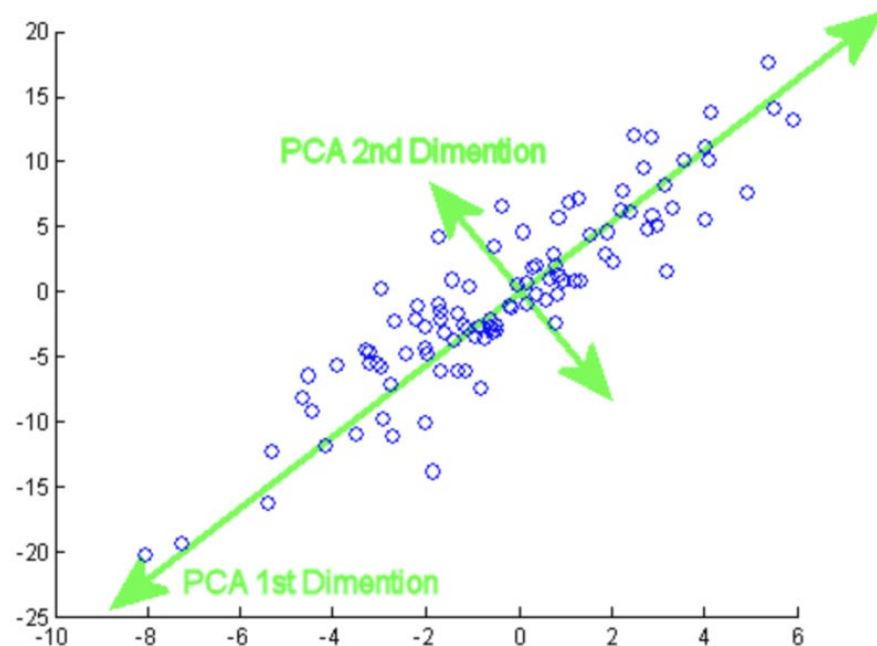
$$\begin{bmatrix} Cov(x, x) & Cov(x, y) & Cov(x, z) \\ Cov(y, x) & Cov(y, y) & Cov(y, z) \\ Cov(z, x) & Cov(z, y) & Cov(z, z) \end{bmatrix}$$

- Covariation is summarized into eigenvectors
- Patterns of covariation are assigned to “principal component axes” or “principal components” or “axes” or “PCs”



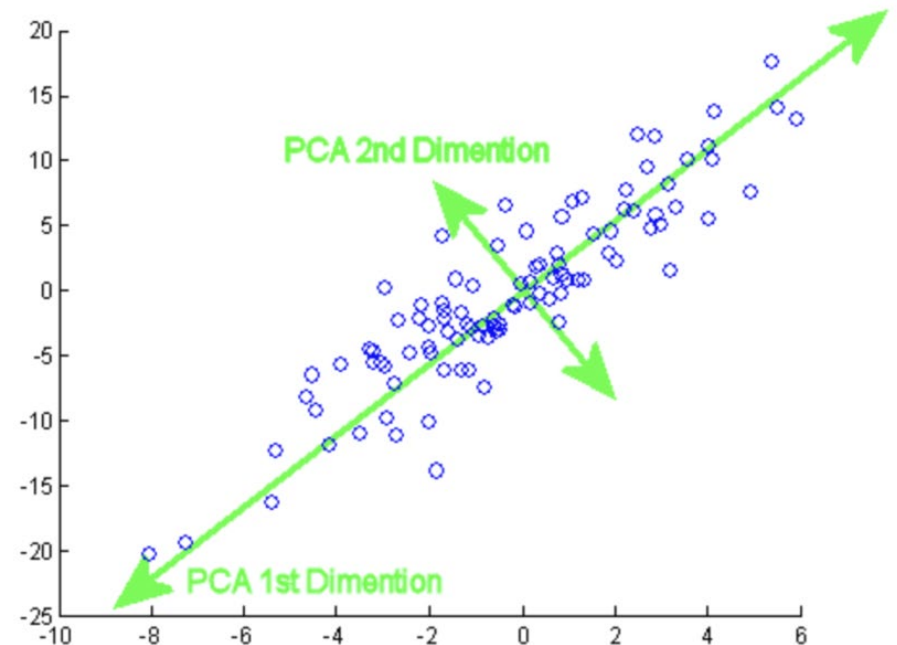
# Principal component analysis (PCA)

- Principal component axes are orthogonal to each other
  - They characterize different “dimensions” of the variation
    - E.g., 10 principal components = 10 dimensions



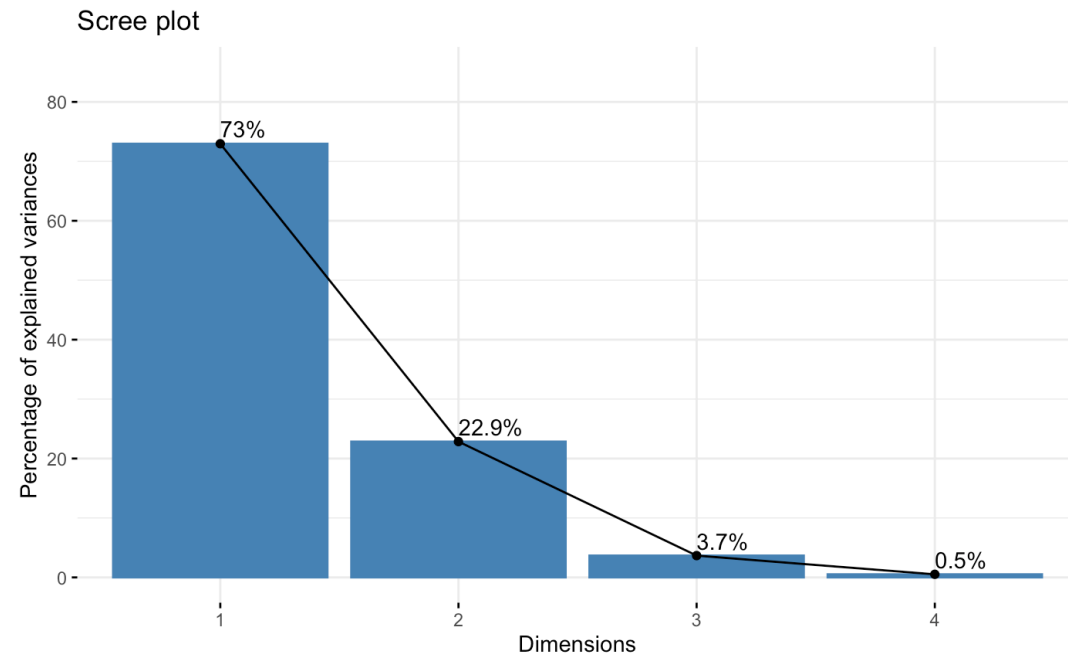
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  - Rest in descending order



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  - They characterize different “dimensions” of the variation
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- First axis explains the most variation
  - Rest in descending order
- Variance explained often decays quickly
  - Visualized with a scree plot

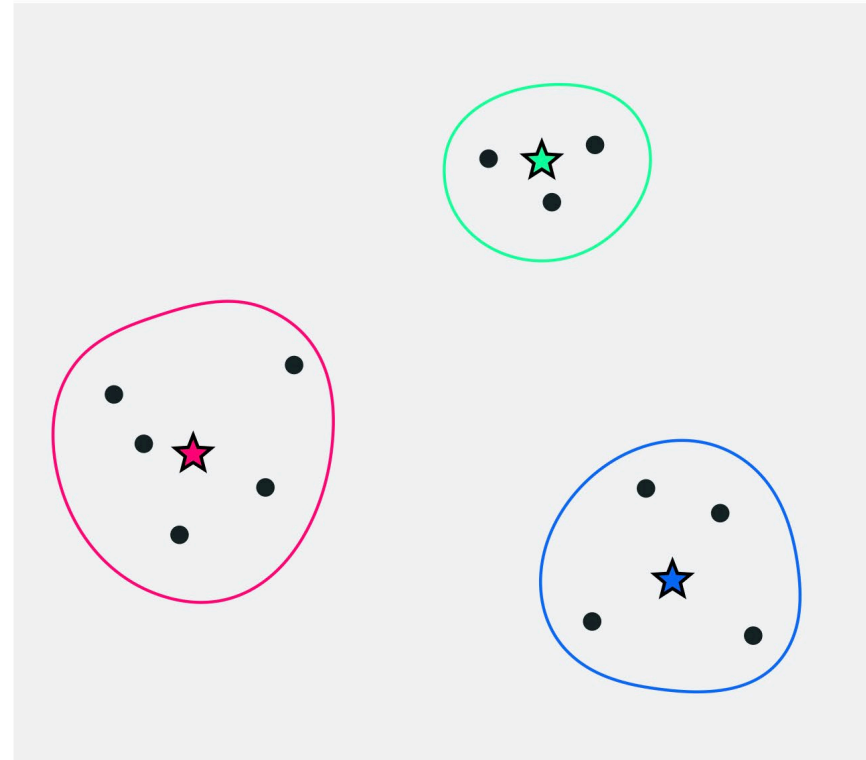
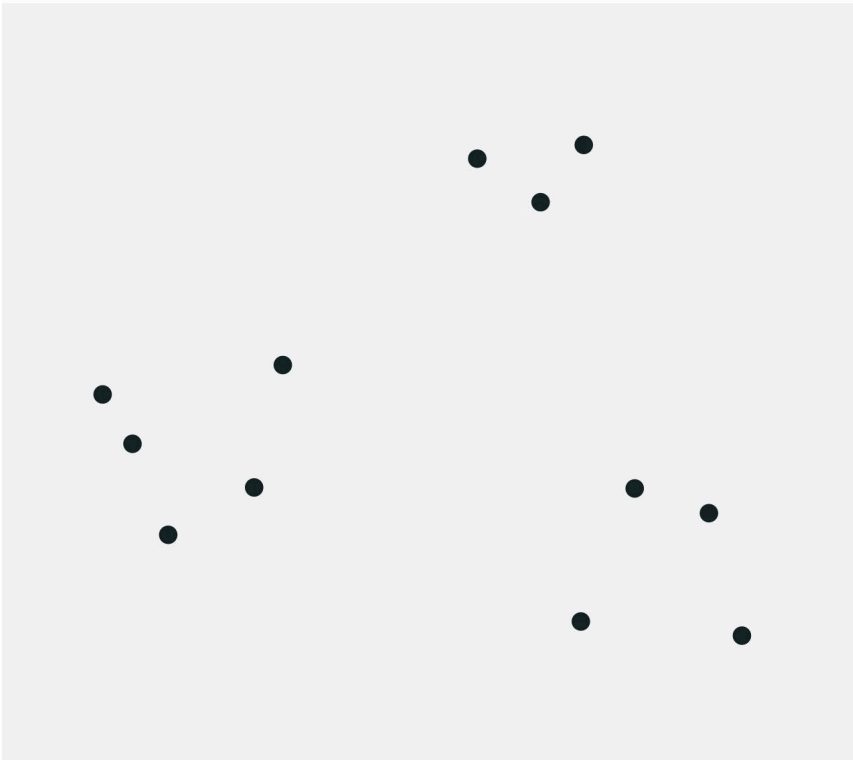


# Principal component analysis (PCA)

- Best practice is usually to scale and center data before PCA
- In `prcomp()` use arguments 'scale' and 'center'
  - Scale: standardize range of variable values
  - Center: center variable values on zero

# K-means clustering

- Goal: group observations based on their overall similarity



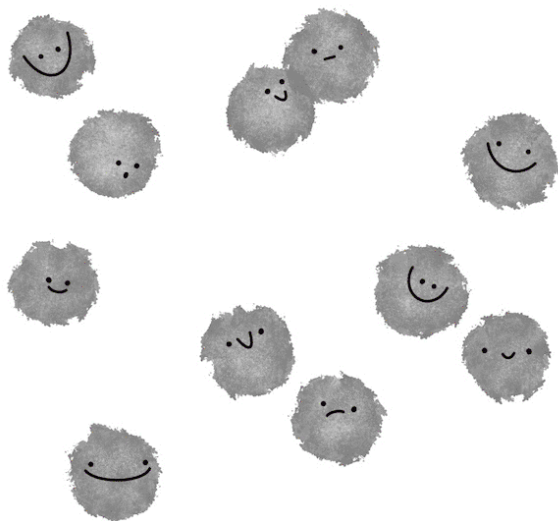
# K-means clustering

- Goal: group observations based on their overall similarity
- Algorithm:
  - Place “centroids” among a scatterplot
    - Assessed number of centroids determined by either biological hypothesis or identified statistically
  - Measure distance from each point to each centroid
  - Move centroids to a locations where distances are minimized
  - Repeat until optimal solution found (smallest average distance from a point to its nearest centroid).

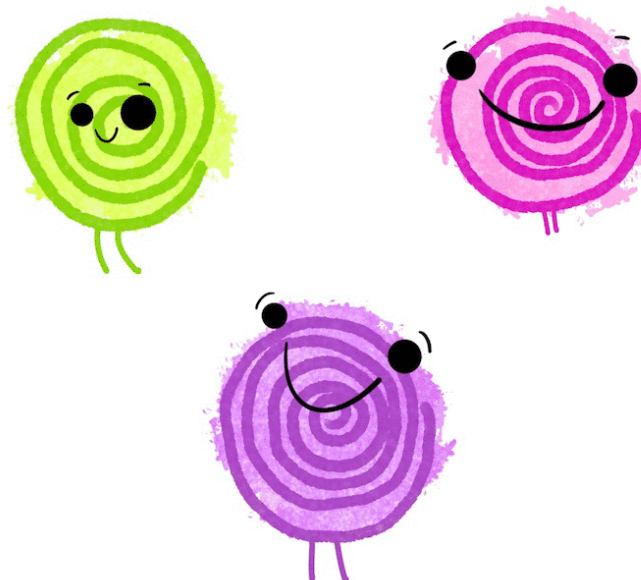
# k-means clustering

- assign each observation to one of  $k$  clusters based on the nearest cluster centroid.

OBSERVATIONS



cluster  
CENTROIDS



# K-means clustering

## What is the value of K?

- Within-cluster sum of squares
- Goal is to minimize within-cluster distance from each point to a centroid
- “Elbow method” for determining best fit
- Lots of other methods for determining best K

