

# Dimension reduction and Principal components regression

---

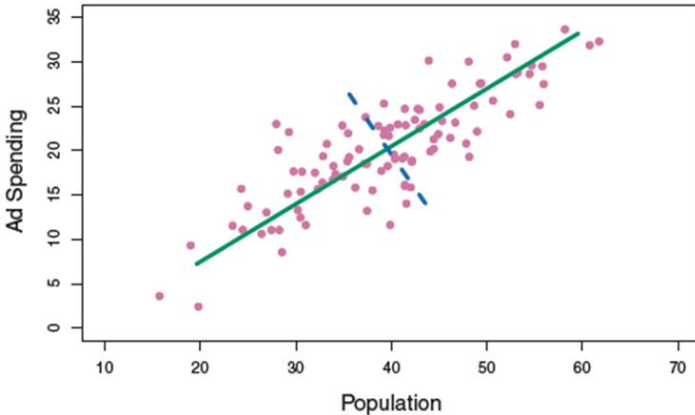
# Outline

- Dimension reduction methods
- Principal components analysis
- Principal components regression
- Considerations in high dimensions

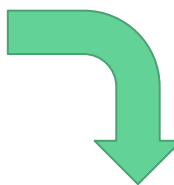
# Dimensions reduction methods

... make new variables by combining existing variables.

$$Z_m = \sum_{j=1}^p \phi_{jm} X_j$$



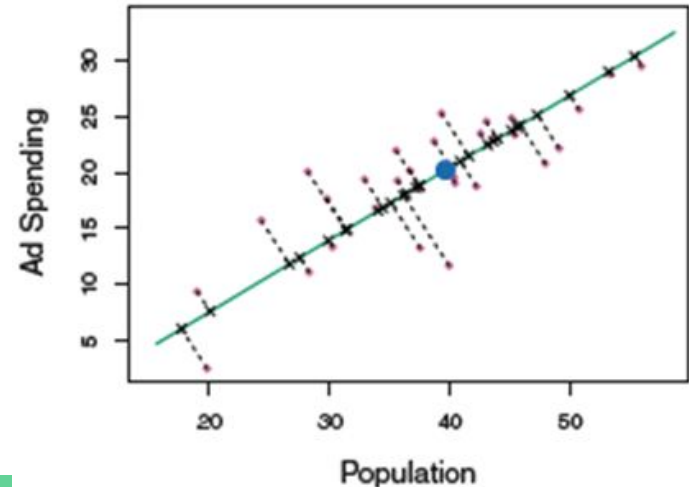
| Income  | Age | Gender | Utility | Childcare | Groceries | Leisure | Health |
|---------|-----|--------|---------|-----------|-----------|---------|--------|
| 14.891  | 34  | Male   | 0.54    | 0.62      | 0.57      | 0.18    | 0.61   |
| 106.025 | 82  | Female | 1.45    | 0         | 0.95      | 0.58    | 0.15   |
| 104.593 | 71  | Male   | 1.95    | 0         | 0.64      | 0.28    | 0.42   |
| 148.924 | 36  | Female | 0.7     | 0.5       | 0.74      | 0.1     | 0.28   |
| 55.882  | 68  | Male   | 1.32    | 0         | 0.56      | 0.43    | 0.15   |
| 80.18   | 77  | Male   | 1.53    | 0         | 0.66      | 0.12    | 0.72   |
| 20.996  | 37  | Female | 1.56    | 0.51      |           |         |        |
| 71.408  | 87  | Male   | 0.56    | 0         |           |         |        |
| 15.125  | 66  | Female | 1.15    | 0         |           |         |        |
| 71.061  | 41  | Female | 0.76    | 0.89      |           |         |        |



| Income  | Age | Gender | Indispensable | Dispensable |
|---------|-----|--------|---------------|-------------|
| 14.891  | 34  | Male   | 1.73          | 0.79        |
| 106.025 | 82  | Female | 2.4           | 0.73        |
| 104.593 | 71  | Male   | 2.59          | 0.7         |
| 148.924 | 36  | Female | 1.94          | 0.38        |
| 55.882  | 68  | Male   | 1.88          | 0.58        |
| 80.18   | 77  | Male   | 2.19          | 0.86        |
| 20.996  | 37  | Female | 2.65          | 0.9         |
| 71.408  | 87  | Male   | 1.32          | 1.21        |
| 15.125  | 66  | Female | 1.71          | 0.92        |
| 71.061  | 41  | Female | 2.43          | 1.01        |

# Principal components analysis (PCA)

- A technique to reduce dimension of an  $n \times p$  data matrix
- Determining new variables by linearly combining the existing variables
- There will be at most  $p$  principal components, but ...
- The *1st principal component* contains *most variability (information)* of the original data. Its direction represents the *line closest to the original data*.
- All principal components are uncorrelated  $\Rightarrow$  their directions are perpendicular to each other.
- Relying only on the predictors  $\Rightarrow$  an unsupervised analysis method



# Steps to determine the principal components

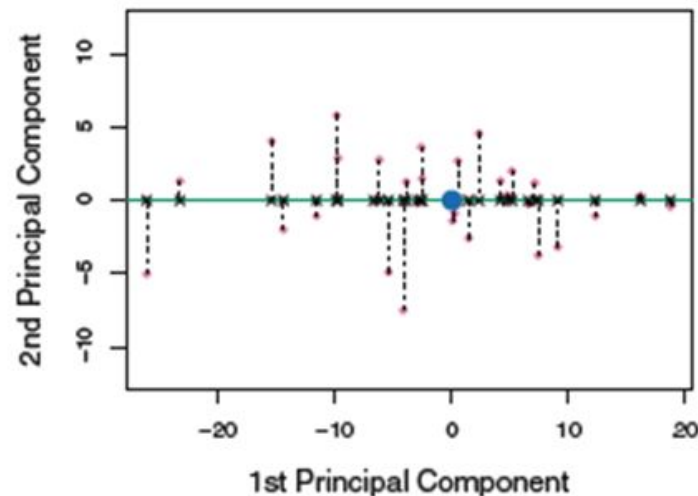
**Step 1.** Standardise the variables (or high variance variable would dominate the principal components)

**Step 2.** Compute covariance matrix

**Step 3.** Calculate the principal components

- Calculate eigenvectors and eigenvalues of the covariance matrix
- Eigenvector of the largest eigenvalue is the 1st PC, and so on

**Step 4.** Project the original variables onto the direction of the (selected) principal components



# Principal components regression (PCR)

PCR is OLS regression of the response on the 1st  $M$  principal components of the original predictors.

PCR assumes that *'the directions in which  $X_1, \dots, X_p$  show the most variation are the directions that are associated with  $Y$ '*

(There's no guarantee that the above assumption is correct in all cases.)

$$y_i = \theta_0 + \sum_{m=1}^M \theta_m z_{im} + \epsilon_i, \quad i = 1, \dots, n,$$

where

$$Z_m = \sum_{j=1}^p \phi_{jm} X_j$$

# Principal components regression (PCR)

PCR is a dimension reduction method (if  $M < p$ ) but not a feature selection method

Interpreting PCR may be challenging, especially when needs to be related to the original predictors

The number of principal components used can be decided by cross-validation

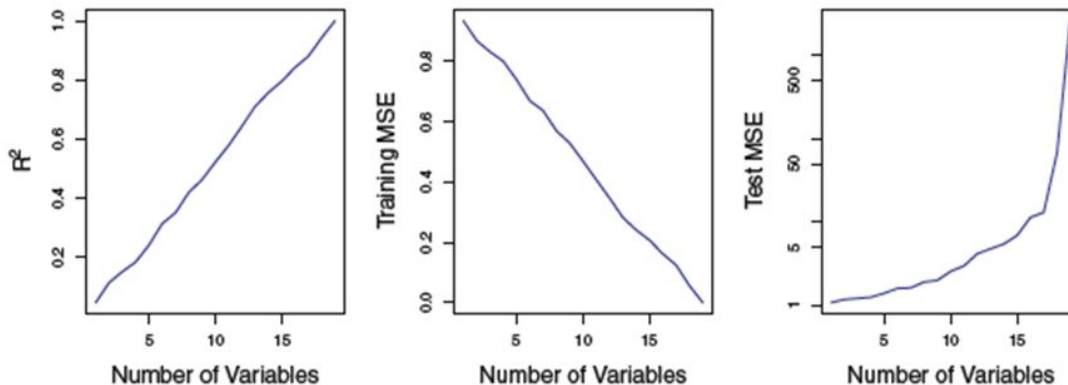
Using a trained PCR model for prediction (e.g. of a test set)

- Standardise predictors in the test set
- Project the standardised predictors onto the axis of the principal components
- Use the projected predictors for input into the PCR model

# Considerations in high dimensions

What goes wrong in high dimensions

- Most traditional learning methods weren't designed for high dimensions (and when  $n$  is not much larger than  $p$ )
- Below are results from an example of regressing 20 response values against 1 to 20 predictors, all of which *unrelated to the response*.



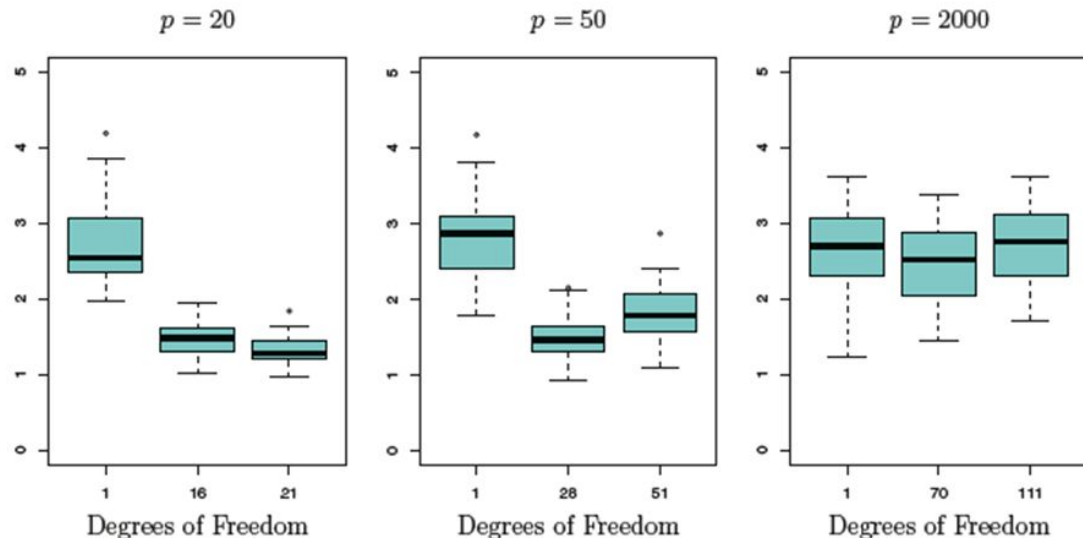


# Considerations in high dimensions

Subset selection methods, regularisation and PCR are useful for regression in high dimensions => *avoiding overfitting* by using *less flexible approach* than OLS

The example on the right (from a lasso) say 3 things

- Lasso helps to reduce dimension
- Correct penalty needed for good predictive
- The curse of dimensionality



# Considerations in high dimensions

More features would do more harm than good

- Deteriorating the (prediction) quality of the fitted model
- More time needed to do feature selection
- Costly data collection and preparation
- Risks of not having the features available in real-life applications

=> choosing variables relevant to the response is critical and must involve subject matter experts in the model building process.

# Considerations in high dimensions

## Interpreting results in high dimensions

- High dimensions present a very good chance of multicollinearity => not sure which variables are predictive of the response
- Large regression coefficients may be assigned to *variables that are correlated to the variables that are truly predictive* of the response
- Be very cautious to conclude a subset of predictors is better than others in predicting the response.
- More reasonable to say the selected subset of predictors forms *one of many possible models* to predict the response.
- *Never* use metrics on the training data (p-value,  $R^2$ ) to report the quality of fit - *always* use independent test sets where possible.