Dimension Reduction

Principal components analysis (PCA)

July 10th, 2023

What is the goal of dimension reduction?

We have p variables (columns) for n observations (rows) **BUT** which variables are **interesting**?

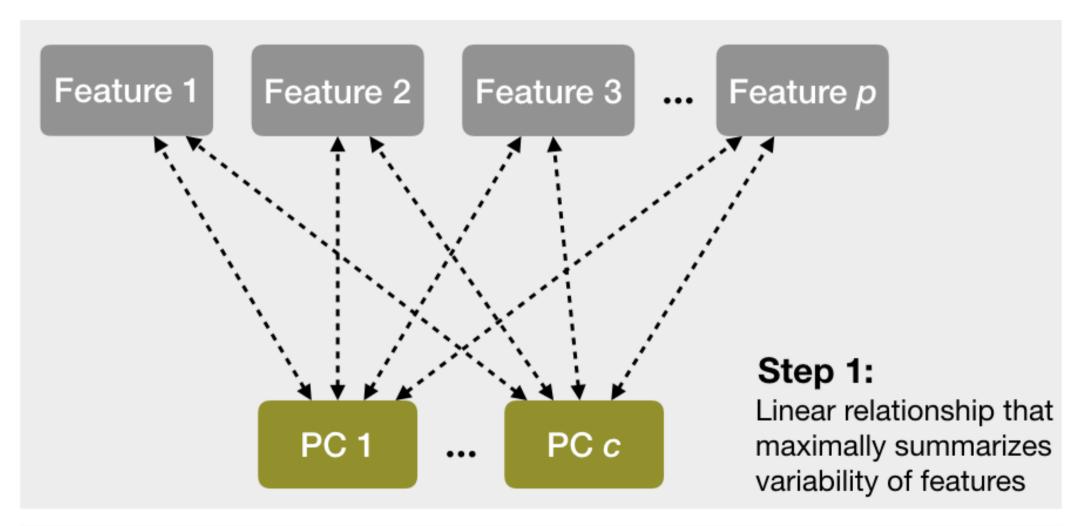
Can we find a smaller number of dimensions that captures the **interesting** structure in the data?

- Could examine all pairwise scatterplots of each variable tedious, manual process
- Tuesday: clustered variables based on correlation
- Can we find a combination of the original *p* variables?

Dimension reduction:

- Focus on reducing the dimensionality of the feature space (i.e., number of columns),
- While **retaining** most of the information / **variability** in a lower dimensional space (i.e., reducing the number of columns)

Principal components analysis (PCA)



Principal components analysis (PCA)

- PCA explores the **covariance** between variables, and combines variables into a smaller set of **uncorrelated** variables called **principal components (PCs)**
- PCs are weighted, linear combinations of the original variables
 - Weights reveal how different variables are *loaded* into the PCs
- We want a **small number of PCs** to explain most of the information / variance in the data

First principal component:

$$Z_1 = \phi_{11} X_1 + \phi_{21} X_2 + \dots + \phi_{p1} X_p$$

- ullet ϕ_{j1} are the weights indicating the contributions of each variable $j \in 1, \dots, p$
- ullet Weights are normalized $\sum_{j=1}^p \phi_{j1}^2 = 1$
- $\phi_1=(\phi_{11},\phi_{21},\ldots,\phi_{p1})$ is the **loading vector** for PC1
- Z_1 is a linear combination of the p variables that has the **largest variance**

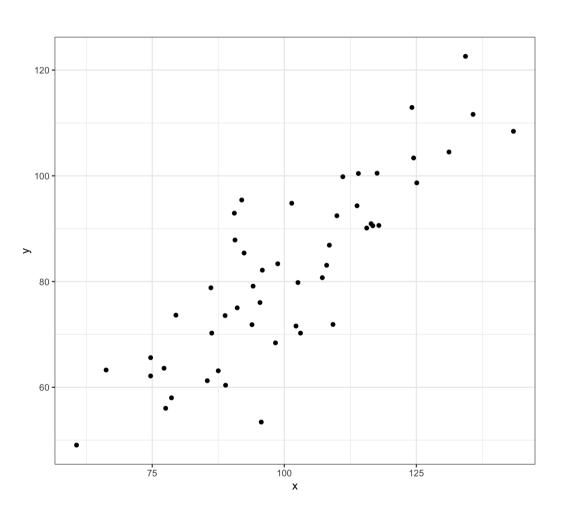
Principal components analysis (PCA)

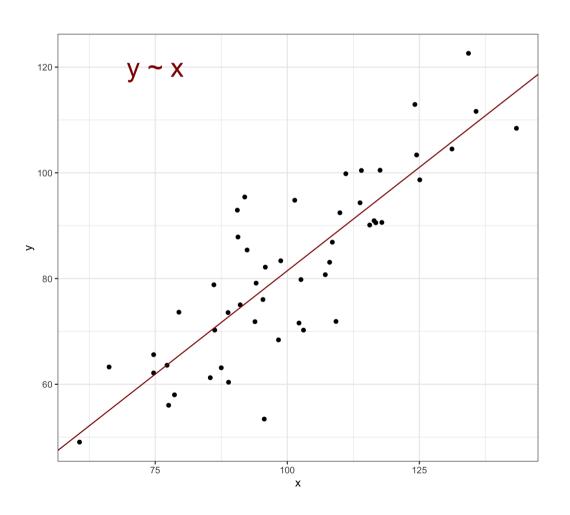
Second principal component:

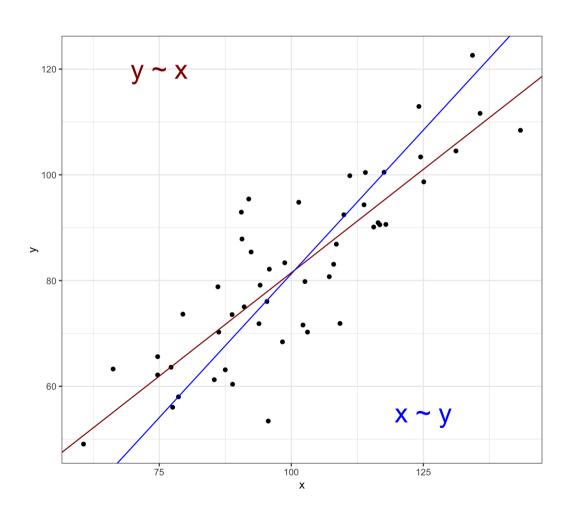
$$Z_2 = \phi_{12} X_1 + \phi_{22} X_2 + \cdots + \phi_{p2} X_p$$

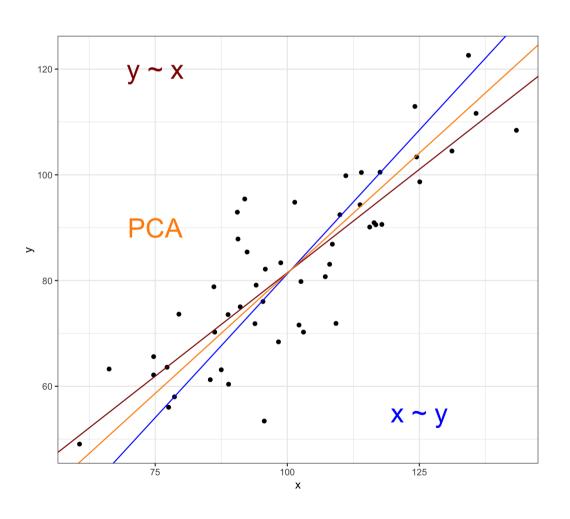
- ullet ϕ_{j2} are the weights indicating the contributions of each variable $j \in 1, \dots, p$
- Weights are normalized $\sum_{j=1}^p \phi_{j1}^2 = 1$
- $\phi_2=(\phi_{12},\phi_{22},\ldots,\phi_{p2})$ is the **loading vector** for PC2
- Z_2 is a linear combination of the p variables that has the **largest variance**
 - $\circ~$ Subject to constraint it is uncorrelated with Z_1

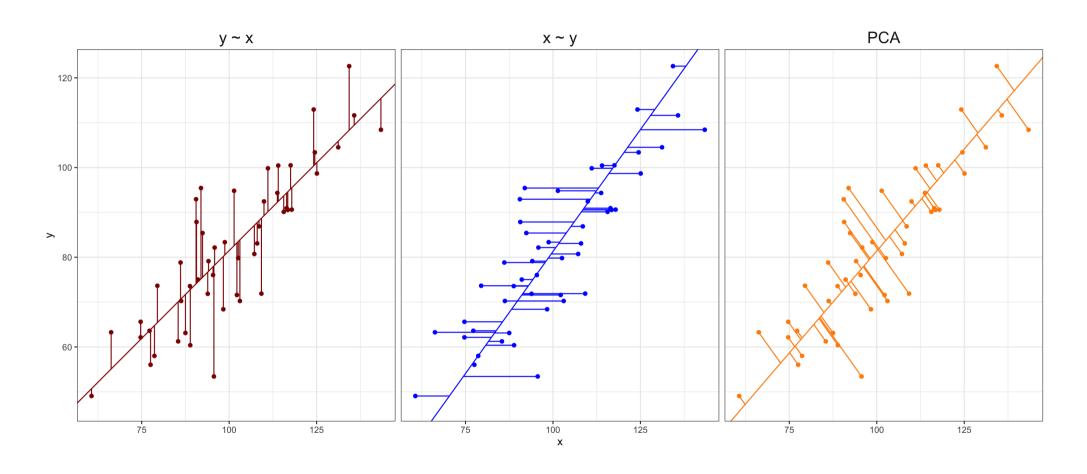
We repeat this process to create p principal components



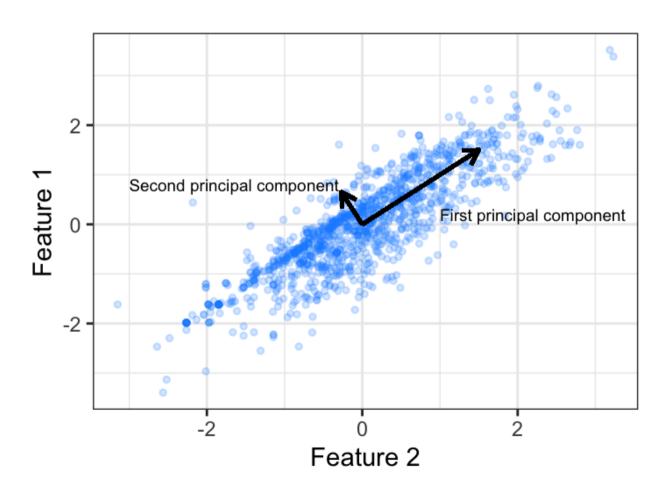








Searching for variance in orthogonal directions



PCA: singular value decomposition (SVD)

$$X = UDV^T$$

- ullet Matrices U and V contain the left and right (respectively) **singular vectors of scaled matrix** X
- ullet D is the diagonal matrix of the **singular values**
- SVD simplifies matrix-vector multiplication as rotate, scale, and rotate again

V is called the **loading matrix** for X with ϕ_i as columns,

• Z = XV is the PC matrix

BONUS eigenvalue decomposition (aka spectral decomposition)

- V are the **eigenvectors** of X^TX (covariance matrix, T means transpose)
- U are the **eigenvectors** of XX^T
- The singular values (diagonal of D) are square roots of the **eigenvalues** of X^TX or XX^T
- Meaning that Z=UD

Eigenvalues solve time travel?



Probably not... but they guide dimension reduction

We want to choose $p^* < p$ such that we are explaining variation in the data

Eigenvalues λ_j for $j \in 1, \ldots, p$ indicate the variance explained by each component

- $\sum_{j=1}^{p} \lambda_j = p$, meaning $\lambda_j \geq 1$ indicates $\mathrm{PC} j$ contains at least one variable's worth in variability
- ullet λ_j/p equals proportion of variance explained by $\mathrm{PC} j$
- ullet Arranged in descending order so that λ_1 is largest eigenvalue and corresponds to PC1
- Can compute the cumulative proportion of variance explained (CVE) with p^* components:

$$ext{CVE}_{p^*} = rac{\sum_{j}^{p^*} \lambda_j}{p}$$

Can use scree plot to plot eigenvalues and guide choice for $p^* < p$ by looking for "elbow" (rapid to slow change)

Example data: NFL teams summary

Created dataset using nflfastR summarizing NFL team performances from 1999 to 2021

```
library(tidyverse)
nfl_teams_data <- read_csv("https://shorturl.at/cfmpW")
nfl_model_data <- nfl_teams_data %>%
   mutate(score_diff = points_scored - points_allowed) %>%
   # Only use rows with air yards
   filter(season >= 2006) %>%
   dplyr::select(-wins, -losses, -ties, -points_scored, -points_allowed, -season, -team)
```

NFL PCA example

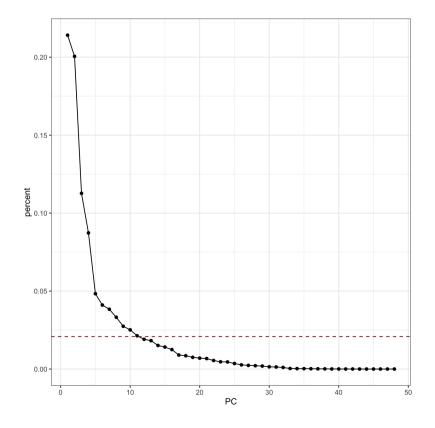
Use the prcomp function (uses SVD) for PCA on centered and scaled data

```
model_x <- as.matrix(dplyr::select(nfl_model_data, -score_diff))</pre>
pca_nfl <- prcomp(model_x, center = TRUE, scale = TRUE)</pre>
summary(pca_nfl)
## Importance of components:
##
                             PC1
                                     PC2
                                            PC3
                                                                             PC7
                                                    PC4
                                                            PC5
                                                                     PC6
## Standard deviation
                          3.2060 3.1026 2.3257 2.04728 1.52301 1.40350 1.35714
## Proportion of Variance 0.2141 0.2006 0.1127 0.08732 0.04832 0.04104 0.03837
## Cumulative Proportion 0.2141 0.4147 0.5274 0.61468 0.66301 0.70405 0.74242
##
                                       PC9
                                              PC10
                                                      PC11
                                                              PC12
                                                                       PC13
                              PC8
                                                                               PC14
                          1.26250 1.14773 1.09881 1.01200 0.95689 0.93513 0.85233
## Standard deviation
## Proportion of Variance 0.03321 0.02744 0.02515 0.02134 0.01908 0.01822 0.01513
## Cumulative Proportion
                          0.77562 0.80307 0.82822 0.84956 0.86863 0.88685 0.90199
                                              PC17
                                                      PC18
                                                                      PC20
##
                             PC15
                                      PC16
                                                              PC19
                                                                              PC21
## Standard deviation
                          0.82315 0.77434 0.65692 0.64016 0.60076 0.5796 0.56756
## Proportion of Variance 0.01412 0.01249 0.00899 0.00854 0.00752 0.0070 0.00671
## Cumulative Proportion 0.91610 0.92859 0.93758 0.94612 0.95364 0.9606 0.96735
##
                             PC22
                                      PC23
                                              PC24
                                                      PC25
                                                              PC26
                                                                       PC27
                                                                               PC28
## Standard deviation
                          0.51349 0.47233 0.46768 0.41284 0.35810 0.33597 0.32018
## Proportion of Variance 0.00549 0.00465 0.00456 0.00355 0.00267 0.00235 0.00214
## Cumulative Proportion 0.97284 0.97749 0.98205 0.98560 0.98827 0.99062 0.99276
```

Proportion of variance explained

prcomp\$sdev corresponds to the singular values, i.e., $\sqrt{\lambda_j}$, what is pca_nfl\$sdev^2 / ncol(model_x)? Can use the broom package easily tidy prcomp summary for plotting

• Add reference line at 1/p, why?

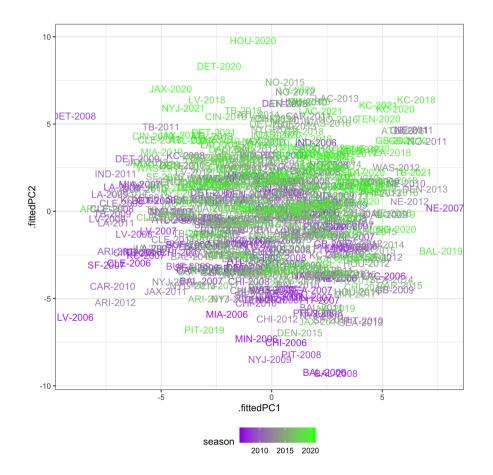


Display data in lower dimensions

prcompx corresponds to the matrix of **principal component scores**, i.e., Z = XV

Can augment dataset with PC scores for plotting

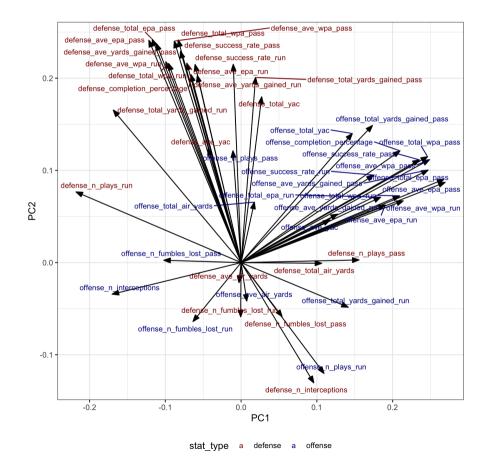
Add team and season for context



What are the loadings of these dimensions?

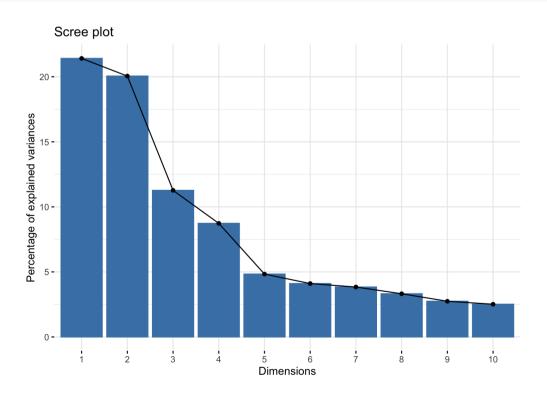
prcomp\$rotation corresponds to the ${f loading\ matrix}$, i.e., V

```
arrow style <- arrow(</pre>
 angle = 20, ends = "first", type = "closed"
 length = grid::unit(8, "pt")
library(ggrepel)
pca nfl %>%
 tidv(matrix = "rotation") %>%
 pivot wider(names from = "PC", names prefix
              values from = "value") %>%
 mutate(stat type = ifelse(str detect(column
                            "offense", "defen
 ggplot(aes(PC1, PC2)) +
 geom segment(xend = 0, yend = 0, arrow = ar
 geom text repel(aes(label = column, color =
                  size = 3) +
 scale_color_manual(values = c("darkred", "d
 theme bw() +
 theme(legend.position = "bottom")
```



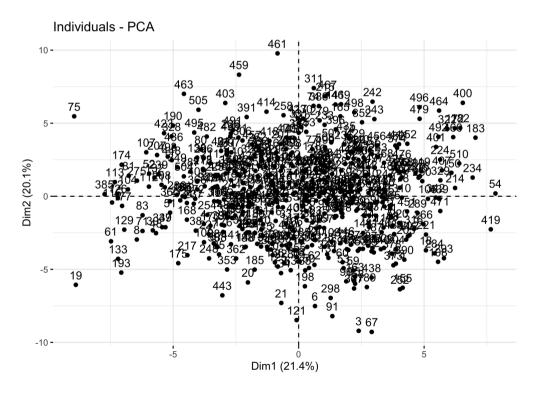
Visualize the proportion of variance explained by each PC with factoextra

```
library(factoextra)
fviz_eig(pca_nfl)
```



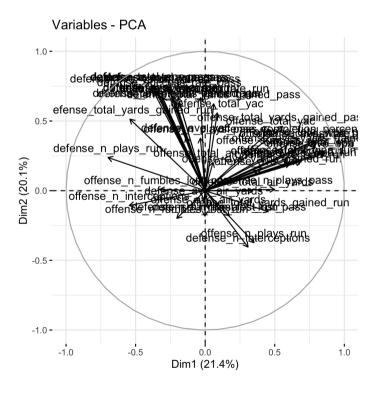
Display observations with first two PC

fviz_pca_ind(pca_nfl)



Projection of variables - angles are interpreted as correlations, where negative correlated values point to opposite sides of graph

fviz_pca_var(pca_nfl)



Biplot displays both the space of observations and the space of variables

• Arrows represent the directions of the original variables

fviz_pca_biplot(pca_nfl)

