Math 308: Principal Component Analysis

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Task:

Perform a principal component analysis of this data using your preferred function. As part of this analysis, please be sure complete the following tasks: - Report the eigenvalues for all 11 principal components. - For the first two principal components, plot and interpret components in terms of the original variables. In particular, explain which variables are most highly correlated with each of these two components and how these components are different from each other. - Choose the smallest number of principal components that you believe can be used to summarize the information from the data and justify your choice.

Solution:

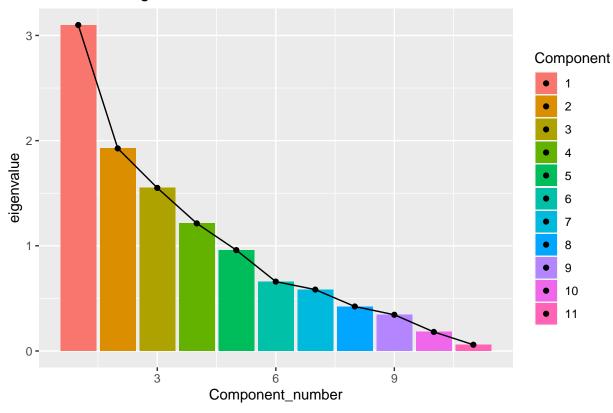
```
library(tinytex)
library(tidyverse)
## -- Attaching core tidyverse packages ------ tidyverse 2.0.0 --
## v dplyr
             1.1.2
                       v readr
                                   2.1.4
## v forcats
              1.0.0
                       v stringr
                                   1.5.0
## v ggplot2
              3.4.2
                       v tibble
                                   3.2.1
## v lubridate 1.9.2
                       v tidyr
                                   1.3.0
## v purrr
              1.0.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                   masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
library(factoextra)
## Welcome! Want to learn more? See two factoextra-related books at https://goo.gl/ve3WBa
library(FactoMineR)
library(GGally)
## Registered S3 method overwritten by 'GGally':
    method from
##
    +.gg
           ggplot2
## Rows: 1599 Columns: 12
## -- Column specification -------
## Delimiter: ","
## dbl (12): fixed.acidity, volatile.acidity, citric.acid, residual.sugar, chlo...
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

```
## # A tibble: 6 x 12
     fixed.acidity volatile.acidity citric.acid residual.sugar chlorides
             <dbl>
                                           <dbl>
                                                           <dbl>
##
                               <dbl>
               7.4
                                0.7
                                                             1.9
                                                                     0.076
## 1
                                            Λ
## 2
               7.8
                                0.88
                                            0
                                                             2.6
                                                                     0.098
## 3
               7.8
                                0.76
                                            0.04
                                                             2.3
                                                                     0.092
## 4
              11.2
                                0.28
                                            0.56
                                                             1.9
                                                                     0.075
## 5
               7.4
                                0.7
                                            Λ
                                                             1.9
                                                                     0.076
## 6
               7.4
                                0.66
                                            0
                                                             1.8
                                                                     0.075
## # i 7 more variables: free.sulfur.dioxide <dbl>, total.sulfur.dioxide <dbl>,
       density <dbl>, pH <dbl>, sulphates <dbl>, alcohol <dbl>, quality <dbl>
## Rows: 1,599
## Columns: 12
## $ fixed.acidity
                           <dbl> 7.4, 7.8, 7.8, 11.2, 7.4, 7.4, 7.9, 7.3, 7.8, 7.5~
## $ volatile.acidity
                           <dbl> 0.700, 0.880, 0.760, 0.280, 0.700, 0.660, 0.600, ~
## $ citric.acid
                           <dbl> 0.00, 0.00, 0.04, 0.56, 0.00, 0.00, 0.06, 0.00, 0~
## $ residual.sugar
                           <dbl> 1.9, 2.6, 2.3, 1.9, 1.9, 1.8, 1.6, 1.2, 2.0, 6.1,~
## $ chlorides
                           <dbl> 0.076, 0.098, 0.092, 0.075, 0.076, 0.075, 0.069, ~
## $ free.sulfur.dioxide <dbl> 11, 25, 15, 17, 11, 13, 15, 15, 9, 17, 15, 17, 16~
## $ total.sulfur.dioxide <dbl> 34, 67, 54, 60, 34, 40, 59, 21, 18, 102, 65, 102,~
## $ density
                           <dbl> 0.9978, 0.9968, 0.9970, 0.9980, 0.9978, 0.9978, 0~
## $ pH
                           <dbl> 3.51, 3.20, 3.26, 3.16, 3.51, 3.51, 3.30, 3.39, 3~
## $ sulphates
                           <dbl> 0.56, 0.68, 0.65, 0.58, 0.56, 0.56, 0.46, 0.47, 0~
## $ alcohol
                           <dbl> 9.4, 9.8, 9.8, 9.8, 9.4, 9.4, 9.4, 10.0, 9.5, 10.~
                          <dbl> 5, 5, 5, 6, 5, 5, 5, 7, 7, 5, 5, 5, 5, 5, 5, 5, 7~
## $ quality
wine_chem<-wine%>%select(-quality)
head(wine_chem)
## # A tibble: 6 x 11
     fixed.acidity volatile.acidity citric.acid residual.sugar chlorides
##
             <dbl>
                               <dbl>
                                           <dbl>
                                                          <dbl>
                                                                     <dbl>
## 1
               7.4
                                0.7
                                            0
                                                             1.9
                                                                     0.076
## 2
               7.8
                                0.88
                                            0
                                                             2.6
                                                                     0.098
## 3
               7.8
                                0.76
                                            0.04
                                                             2.3
                                                                     0.092
## 4
              11.2
                                0.28
                                            0.56
                                                             1.9
                                                                     0.075
## 5
               7.4
                                            0
                                                             1.9
                                                                     0.076
                                0.7
               7.4
                                0.66
                                            0
                                                             1.8
                                                                     0.075
## # i 6 more variables: free.sulfur.dioxide <dbl>, total.sulfur.dioxide <dbl>,
       density <dbl>, pH <dbl>, sulphates <dbl>, alcohol <dbl>
Eigenvalues:
wine_PCA<-PCA(wine_chem,graph=FALSE)</pre>
wine_PCA$eig[,"eigenvalue"]
                                                                comp 6
       comp 1
                  comp 2
                              comp 3
                                         comp 4
                                                    comp 5
## 3.09913244 1.92590969 1.55054349 1.21323253 0.95929207 0.65960826 0.58379122
                  comp 9
                            comp 10
                                        comp 11
## 0.42295670 0.34464212 0.18133317 0.05955831
```

```
eigenvalues_wine<-as.data.frame(wine_PCA$eig) %>%
   rownames_to_column(var="Component")
eigenvalues_wine <- eigenvalues_wine %>%
   mutate(Component=map_chr(Component,~str_split(.x," ")[[1]][2]),
   Component_number=as.integer(Component),
   Component=factor(Component_number))

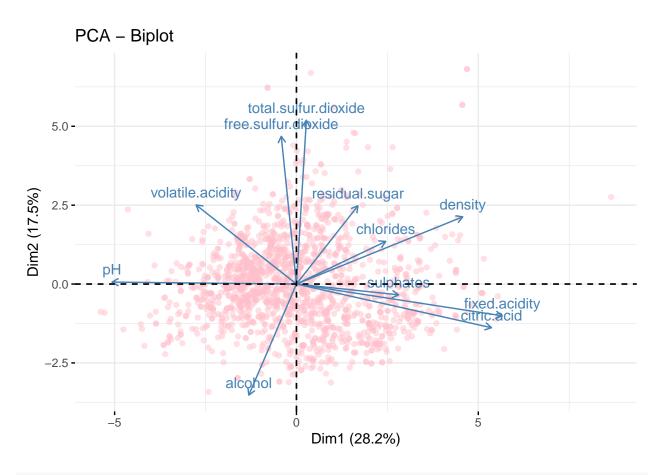
ggplot(eigenvalues_wine,aes(y=eigenvalue,x=Component_number,fill=Component)) +
   geom_bar(stat="identity") + geom_line(aes(fill=NULL)) + geom_point() +ggtitle("Wine Data Eigenvalues")
```

Wine Data Eigenvalues



Component Analysis:

```
fviz_pca_biplot(wine_PCA,col.ind="pink",
    fill.ind="pink",label="var",
    alpha.ind=c(0.5))
```



round(wine_PCA\$var\$coord[,c(1:2)],2)

```
##
                        Dim.1 Dim.2
## fixed.acidity
                         0.86 - 0.15
## volatile.acidity
                        -0.42 0.38
## citric.acid
                         0.82 -0.21
## residual.sugar
                         0.26 0.38
## chlorides
                         0.37
                               0.21
## free.sulfur.dioxide
                        -0.06
                               0.71
## total.sulfur.dioxide
                         0.04
                               0.79
## density
                         0.70 0.32
## pH
                         -0.77 0.01
## sulphates
                         0.43 -0.05
## alcohol
                        -0.20 -0.54
```

From the plot and the table above, we see that the first component mostly is a contrast between the pH level and the fixed acidity, citric acid and density values. These are the variables with the strongest correlations with the first component. The second component mostly depends on the free and total sulfur dioxide measures, which are contrasted mostly with the alcohol content, as these three variables are most strongly associated with the second component.

Number of Components:

wine_PCA\$eig

```
eigenvalue percentage of variance cumulative percentage of variance
          3.09913244
                                  28.1739313
## comp 1
                                                                      28.17393
## comp 2 1.92590969
                                  17.5082699
                                                                      45.68220
## comp 3
          1.55054349
                                  14.0958499
                                                                      59.77805
## comp 4
          1.21323253
                                  11.0293866
                                                                      70.80744
## comp 5 0.95929207
                                   8.7208370
                                                                      79.52827
## comp 6
          0.65960826
                                   5.9964388
                                                                      85.52471
## comp 7
           0.58379122
                                                                      90.83191
                                   5.3071929
## comp 8
          0.42295670
                                   3.8450609
                                                                      94.67697
## comp 9 0.34464212
                                   3.1331102
                                                                      97.81008
## comp 10 0.18133317
                                   1.6484833
                                                                      99.45856
## comp 11 0.05955831
                                   0.5414392
                                                                     100.00000
```

We see that using an 80% cumulative variance rule, we would decide to use either 5 or 6 components depending on how strictly we wanted to use the cutoff.