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Class Activity 12

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```
library(tidyverse)
```

```
## — Attaching core tidyverse packages -
                                                            - tidyverse 2.0.0 —
## √ dplyr
            1.1.3
                       √ readr
                                    2.1.4
## √ forcats
              1.0.0

√ stringr

                                    1.5.0
## √ ggplot2 3.4.4
                       √ tibble
                                    3.2.1
## ✓ lubridate 1.9.3
                       √ tidyr
                                    1.3.0
## √ purrr
              1.0.2
## -- Conflicts --
                                                      - tidyverse_conflicts() —
## X dplyr::filter() masks stats::filter()
## X dplyr::lag()
                    masks stats::lag()
### i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to becom
e errors
```

library(pls)

```
##
## Attaching package: 'pls'
##
## The following object is masked from 'package:stats':
##
## loadings
```

```
baseball <- read.csv( "C:\\Users\\hodge\\Downloads\\Baseball.csv")</pre>
```

```
baseball <- na.omit(baseball)</pre>
```

1a

in comparison to OLS, PCR lowers the risk of multicolinearity within our model and it shrinks the the # of columns to containt the same information as the *p* numerical random variable, but explained in a more convenient way.

1_b

```
colnames(baseball)
```

"Runs"

"RBI"

##

[1] "X"

summary(pca)

"AtBat"

"Hits"

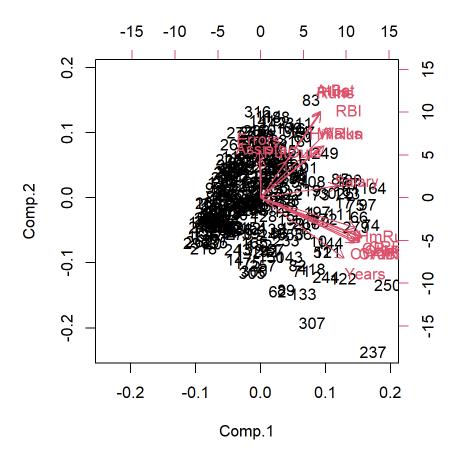
```
"Years"
                                                "CHits"
                                                             "CHmRun"
                                                                          "CRuns"
##
    [7] "Walks"
                                   "CAtBat"
## [13] "CRBI"
                      "CWalks"
                                   "League"
                                                "Division"
                                                             "PutOuts"
                                                                          "Assists"
## [19] "Errors"
                      "Salary"
                                   "NewLeague"
baseball_subset <- subset(baseball, select = -c(X, League, Division, NewLeague))</pre>
pca <- princomp(baseball subset, fix sign = T, cor = T)</pre>
```

"HmRun"

```
## Importance of components:
##
                                                 Comp.3
                             Comp.1
                                       Comp.2
                                                           Comp.4
## Standard deviation
                          2.7733967 2.0302601 1.3148557 0.9575410 0.84109683
## Proportion of Variance 0.4524547 0.2424680 0.1016968 0.0539344 0.04161435
## Cumulative Proportion 0.4524547 0.6949227 0.7966195 0.8505539 0.89216822
##
                              Comp.6
                                         Comp.7
                                                    Comp.8
                                                               Comp.9
                          0.72374220 0.69841796 0.50090065 0.42525940 0.363901982
## Standard deviation
## Proportion of Variance 0.03081193 0.02869339 0.01475891 0.01063797 0.007789685
## Cumulative Proportion 0.92298014 0.95167354 0.96643244 0.97707042 0.984860104
##
                              Comp.11
                                          Comp.12
                                                      Comp.13
## Standard deviation
                          0.312011679 0.243641510 0.232044829 0.163510472
## Proportion of Variance 0.005726546 0.003491834 0.003167341 0.001572687
## Cumulative Proportion 0.990586651 0.994078485 0.997245826 0.998818513
##
                               Comp.15
                                            Comp.16
## Standard deviation
                          0.1186398422 0.0693395039 3.466841e-02
## Proportion of Variance 0.0008279654 0.0002828216 7.069994e-05
## Cumulative Proportion 0.9996464785 0.9999293001 1.000000e+00
```

```
biplot(pca)
```

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1c

PCRRRRR

pcareg <-pcr(Salary~., data = baseball_subset, scale = T,) #does principal component regression</pre>

1d

based on the model summary below, the R^2 if we used the first 3 principal components looks to be 41.93, and the R^2 if we used the first 10 principal components is 45.67.

summary(pcareg)

```
## Data:
           X dimension: 263 16
## Y dimension: 263 1
## Fit method: svdpc
## Number of components considered: 16
## TRAINING: % variance explained
##
           1 comps 2 comps 3 comps 4 comps 5 comps 6 comps 7 comps
                      71.00
                              81.80
                                       87.24
## X
             45.31
                                                 91.60
                                                          94.80
                                                                  96.37
                                                                            97.53
## Salary
             40.67
                      41.87
                              41.93
                                       43.96
                                                 44.36
                                                          44.43
                                                                  44.53
                                                                           45.36
##
           9 comps 10 comps 11 comps 12 comps 13 comps 14 comps 15 comps
             98.36
                      98.97
                                99.35
                                           99.70
                                                    99.87
                                                               99.96
                                                                         99.99
## X
            45.36
                      45.67
                                47.59
                                          48.36
## Salary
                                                    50.81
                                                               51.97
                                                                         52.78
          16 comps
##
## X
            100.00
## Salary
              52.79
```

1e

The salary prediction for the first two rows of the dataset using the first 3 principal components is 509.4619 for the 2nd row and 634.2457 for the third row.

```
new_dat = baseball_subset[1:2, ]
predict(pcareg, new_dat, ncomp = 3)
```

```
## , , 3 comps

##

## Salary

## 2 509.4619

## 3 634.2457
```

and with the first 10 principal components, we have salary predictions of 507.9765 for the 2nd row and 763.0485 for the 3rd row.

```
new_dat = baseball_subset[1:2, ]
predict(pcareg, new_dat, ncomp = 10)
```

```
## , , 10 comps
##
## Salary
## 2 507.9765
## 3 763.0485
```

2

some advantages for using a Lasso regression instead of PCR are the feature selection that automatically selects the most important features in a dataset. PCR considers all input predictors.