Principal Component Regression (PCR)

Contents

- 1. Multiple Linear Regression-Quick Recap
- 2. The Problem of Multicollinearity
- 3. Principal Component Analysis General Approach
- 4. Principal Component Regression (PCR)
 - i. Introduction
 - ii. Statistical Model
- 5. PCR in R

Multiple Linear Regression: Statistical Model

$$Y = b_0 + b_1 X_1 + b_2 X_2 + ... + b_p X_p + e$$

Where,

: Dependent Variable

 $X_1, X_2, ..., X_p$

: Independent Variables

 $b_0, b_1, ..., b_n$: Parameters of Model

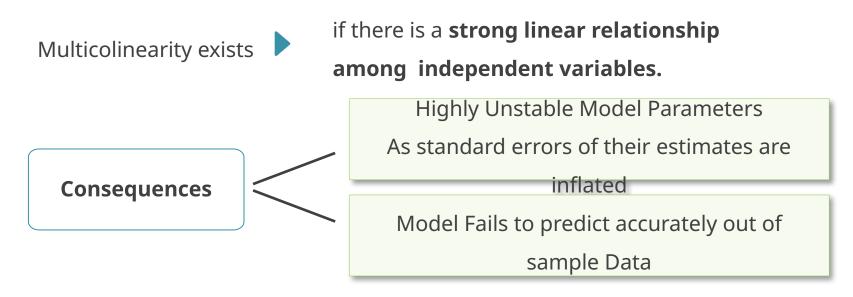
e

: Random Error

Component

Independent variables can either be Continuous or Categorical

Problem of Multicolinearity



Multicolinearity is detected using Variance Inflation Factor, VIF

Tolerance = 1- R_i²

VIF = 1/Tolerance

where R_i² (R Squared) is obtained using regression of Xi on other independent variables

Any VIF > 5, indicates presence of multicollinearity

Multicollinearity - Remedial Measures

The problem of Multicollinearity can be solved by different approaches:

Drop one of the independent variables, which is explained by others

Use Principal Component Regression in case of severe Multicollinearity

Use Ridge Regression

Principal Component Regression

In Principal Component Regression,

First k principal components are used as independent variables instead of original X variables

- Each PC is a linear combination of all X variables
- Final model is expressed in terms of original independent variables for ease of interpretation

Principal Component Regression

Transformatio n into PCs

The original **p** variables are transformed into a new set of orthogonal or uncorrelated variables called "Principal Components "



Regression Analysis



In the second step, after elimination of the least important principal components, a multiple regression analysis of the response variable against the reduced set of principal components is performed using the OLS estimation

Back Transformatio n

In the third step, model equation is back transformed in terms of original variables.

PCR-Statistical Model

Model in terms of original X variables:

$$Y = b_0 + b_1 x_1 + b_2 x_2 + \dots + b_p x_p + e$$

Model in terms of Principal Components:

$$Y = a_0 + a_1 PC_1 + a_2 PC_2 + \dots + a_k PC_k + e'$$

Case Study

Background

A company periodically records data for sales and expenses.
 The company wishes to model the relationship between its sales and sales related expenses and obtain predictions

Objective

 To predict incremental sales based on planned sales related expenses

Available Information

- Data available for 143 micro business zones
- Sales is the Dependent Variable
- Expenditure towards advertisements and promotions in the current and previous months are Predictors

Data Snapshot

		Dependent variable]	Independent variables				
		SRNO	SALES	AD	PRO)	SALEXP	ADPRE	PROPR	E
		1	20.11	1.98	0.9		0.31	2.02	0	
	Columns		Description			Туре		Measurem ent		Possible values
	SRNO		Serial Number			-		-		Intergers
	SALES		Incremental Sales			Numerical INR M		illion	positive value	
	AD		Current Advertising Expenses			Nu	Numerical INR Mi		illion	positive value
	PRO		Current Promotional Expenses			Numerical		INR M	illion	positive value
	SALEXP		Misc. Sales Expenses			Numerical		INR Million		positive value
	ADPRE		Previous Period's Advertising Expenses			Numerical		INR Million p		positive values
	PROPRE		Previous Period's Promotional Expenses			Numerical IN		INR M	illion	Positive value

```
# Import csv file "pcrdata"
salesdata<-read.csv("pcrdata.csv",header=T)

# Fitting a Linear Model :
predsales<-lm(SALES~AD+PRO+SALEXP+ADPRE+PROPRE,data=salesdata)

summary(predsales)

Im() fits a linear regression model.

summary() generates model

# Output of summary</pre>
```

```
Coefficients:
          Estimate Std. Error t value Pr(>|t|)
(Intercept) -10.8147 6.5314 -1.656 0.10005
        4.6762 1.4100 3.316 0.00117 **
ΑD
PRO
         7.7886 1.2628 6.168 7.3e-09 ***
SALEXP 22.4089 0.7704 29.089 < 2e-16 ***
         3.1856 1.2442 2.560 0.01154 *
ADPRE
                     1.3697 2.553 0.01177 *
PROPRE
       3.4970
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.201 on 137 degrees of freedom
Multiple R-squared: 0.9089, Adjusted R-squared: 0.9055
F-statistic: 273.2 on 5 and 137 DF, p-value: < 2.2e-16
```

Interpretation:

Multiple R-Squared is 0.9089, showing model to be a good fit.

Checking for Multicolinearity

```
install.packages("car")
library(car)

vif(predsales)
 vif() in package car calculates VIFs.
```

Output of VIF

```
AD PRO SALEXP ADPRE PROPRE 36.159771 31.846727 1.076284 24.781948 42.346468
```

Interpretation:

VIF values are very high (>5, except for SALEEXP) indicating severe multicolinearity problem.

```
# PCA in R
# Subsetting data for getting Principal components and performing PCA
salesdatapca<-subset(salesdata, select=c(-SRNO, -SALES))</pre>
pc<-princomp(formula=~.,data=salesdatapca, cor=T)
summary(pc)
                   princomp()from base R performs PCA on
                   the
                      given numeric data matrix
                     formula= contains the numeric variables.
                      ensures all variables are taken
                      cor=T indicates that calculations should
                   be done
                      using the Correlation Matrix.
                      summary() generates the summary of
                   PCA
```

Output

Interpretation:

The first three principal components explain 82% of the variation in the data. Therefore, we can use
 3 PC's in Regression Model

PCR in R

```
install.packages("pls")
                             Install and load package pls (Partial
library(pls)
                             Least Squares).
pcmodel<-
pcr(SALES~AD+PRO+SALEXP+ADPRE+PROPRE, ncomp=3, aata=salesaata, scale=IROE)
     pcr() in package pls performs Principal Component Regression
     □ ncomp=3 is the number of components to be included in the model
        scale=TRUE indicates X is scaled by dividing each variable by its
     standard
        deviation. This ensures data is standardised before running the PCR
sales algorithm.
head(salesdata)
                          predict() is used to get predictions by
                          PCR.
```

Output

```
AD PRO SALEXP ADPRE PROPRE pred_pcr
 SRNO SALES
    1 20.11 1.98 0.9
                      0.31 2.02
                                   0.0 21.29053
2
    2 15.10 1.94 0.0 0.30 1.99
                                  1.0 18.16976
3
    3 18.68 2.20 0.8 0.35 1.93 0.0 21.27149
    4 16.05 2.00 0.0 0.35 2.20 0.8 17.62114
5
    5 21.30 1.69 1.3 0.30 2.00 0.0 22.97930
    6 17.85 1.74 0.3
                    0.32 1.69
                                  1.3 20.57217
```

pred_pcr column gives predicted values of SALES using PCR.

Comparing Linear Regression Model and PCR model on Test data

Importing Test Data

```
salesdata_test<-read.csv("pcrdata_test.csv",header=TRUE)</pre>
```

Getting RMSE of linear regression model

```
salesdata_test$Impredict<-predict(predsales,salesdata_test) ←
salesdata_test$Imres<-(salesdata_test$SALES-salesdata_test$Impredict)
RMSE_lm<-sqrt(mean(salesdata_test$lmres**2))</pre>
```

Getting RMSE of PCR model

predict () will give the predicted value for the model.

```
salesdata_test$pcrpredict<-predict(pcmodel,salesdata_test,ncomp=3) 
salesdata_test$pcrres<-(salesdata_test$SALES-salesdata_test$pcrpredict)
RMSE_pcr<-sqrt(mean(salesdata_test$pcrres**2))</pre>
```

Comparing Linear Regression Model and PCR model on Test data

Viewing data after adding predicted & residual variables

```
head(salesdata_test)
 # Output
               PRO SALEXP ADPRE PROPRE
                                        lmpredict
                                                        lmres pcrpredict
SRNO SALES
            AD
                                                                            pcrres
                                        32.31368 -3.3836776
  1 28.93 2.75 1.00
                      0.72 1.97
                                   0.02
                                                               23, 23291
                                                                         5.6970943
  2 25.96 1.73 1.06
                      0.89 2.77
                                   0.02
                                        34.36925 -8.4092464
                                                               22,26693
                                                                         3.6930660
                                   0.42 32.29821
  3 31.25 2.19 1.26
                      0.79
                           1.22
                                                  -1.0482117
                                                               27.61578
                                                                         3.6342207
  4 25.05 1.82 1.45
                      0.83 2.23
                                  0.15 35.21751 -10.1675083
                                                               25.21307 -0.1630736
  5 27.32 2.38 1.01
                                  0.07
                                        28.22616 -0.9061594
                                                               27.05439
                      0.74 1.01
                                                                         0.2656139
  6 23.23 2.97 0.46
                      0.96 2.36
                                   0.12 36.10681 -12.8768143
                                                               20.92296
                                                                         2.3070370
```

```
RMSE_lm
[1] 9.111682
RMSE_pcr
[1] 2.851245
```

Interpretation:

RMSE using PCR is less than RMSE using linear regression, we may conclude that PCR model predicts SALES better than linear regression model when multicolinearity exists.

Quick Recap

Multiple Linear Regression and Multicollinearity

• Highly correlated predictor variables is a very frequent phenomenon in real world analytics.

Principal Component Regression

 PCR is a three way process where the variables are first transformed to principal components, regression is run by considering these components as regressors and finally, they are transformed back to their original forms.

PCR in R

• pcr() function in package pls performs PCR.

THANK YOU!