Preprocessing with Principal Components Analysis (PCA)

Jeffrey Leek

Correlated predictors

Correlated predictors

```
names(spam)[c(34,32)]
[1] "num415" "num857"
plot(spam[,34],spam[,32])
```

Basic PCA idea

- We might not need every predictor
- A weighted combination of predictors might be better
- We should pick this combination to capture the "most information" possible
- Benefits
- Reduced number of predictors
- Reduced noise (due to averaging)

We could rotate the plot

$$X = 0.71 \times \text{num}415 + 0.71 \times \text{num}857$$

$$Y = 0.71 \times \text{num}415 - 0.71 \times \text{num}857$$

```
X <- 0.71*training$num415 + 0.71*training$num857
Y <- 0.71*training$num415 - 0.71*training$num857
plot(X,Y)</pre>
```

Related problems

You have multivariate variables X_1, \ldots, X_n so $X_1 = (X_{11}, \ldots, X_{1m})$

- Find a new set of multivariate variables that are uncorrelated and explain as much variance as possible.
- If you put all the variables together in one matrix, find the best matrix created with fewer variables (lower rank) that explains the original data.

The first goal is statistical and the second goal is data compression.

Related solutions - PCA/SVD

SVD

If X is a matrix with each variable in a column and each observation in a row then the SVD is a "matrix decomposition"

$$X = UDV^T$$

where the columns of U are orthogonal (left singular vectors), the columns of V are orthogonal (right singular vectors) and D is a diagonal matrix (singular values).

PCA

The principal components are equal to the right singular values if you first scale (subtract the mean, divide by the standard deviation) the variables.

Principal components in R - prcomp

```
smallSpam <- spam[,c(34,32)]
prComp <- prcomp(smallSpam)
plot(prComp$x[,1],prComp$x[,2])</pre>
```

Principal components in R - prcomp

prComp\$rotation

```
PC1 PC2
num415 0.7081 0.7061
num857 0.7061 -0.7081
```

PCA on SPAM data

```
typeColor <- ((spam$type=="spam")*1 + 1)
prComp <- prcomp(log10(spam[,-58]+1))
plot(prComp$x[,1],prComp$x[,2],col=typeColor,xlab="PC1",ylab="PC2")</pre>
```

PCA with caret

```
preProc <- preProcess(log10(spam[,-58]+1),method="pca",pcaComp=2)
spamPC <- predict(preProc,log10(spam[,-58]+1))
plot(spamPC[,1],spamPC[,2],col=typeColor)</pre>
```

Preprocessing with PCA

```
preProc <- preProcess(log10(training[,-58]+1),method="pca",pcaComp=2)
trainPC <- predict(preProc,log10(training[,-58]+1))
modelFit <- train(training$type ~ .,method="glm",data=trainPC)</pre>
```

Preprocessing with PCA

```
testPC <- predict(preProc,log10(testing[,-58]+1))
confusionMatrix(testing$type,predict(modelFit,testPC))</pre>
```

Confusion Matrix and Statistics

Reference

Prediction nonspam spam nonspam 646 51 spam 64 389

Accuracy: 0.9

95% CI : (0.881, 0.917)

No Information Rate : 0.617 P-Value [Acc > NIR] : <2e-16

Kappa : 0.79

Mcnemar's Test P-Value : 0.263

Sensitivity: 0.910
Specificity: 0.884
Pos Pred Value: 0.927
Neg Pred Value: 0.859
Prevalence: 0.617
Detection Rate: 0.562
Detection Prevalence: 0.606

'Positive' Class : nonspam

Alternative (sets # of PCs)

```
modelFit <- train(training$type ~ .,method="glm",preProcess="pca",data=training)
confusionMatrix(testing$type,predict(modelFit,testing))</pre>
```

Confusion Matrix and Statistics

Reference

Prediction nonspam spam nonspam 660 37 spam 54 399

Accuracy: 0.921

95% CI: (0.904, 0.936)

No Information Rate : 0.621 P-Value [Acc > NIR] : <2e-16

Kappa: 0.833

Mcnemar's Test P-Value : 0.0935

Sensitivity : 0.924 Specificity : 0.915 Pos Pred Value : 0.947 Neg Pred Value : 0.881 Prevalence : 0.621 Detection Rate : 0.574 Detection Prevalence : 0.606

'Positive' Class : nonspam

Final thoughts on PCs

- Most useful for linear-type models
- Can make it harder to interpret predictors
- Watch out for outliers!
- Plot predictors to identify problems
- For more info see
- Exploratory Data Analysis
- Elements of Statistical Learning