Chapter 3: Regression

- ► Simple linear regression
- ► Multiple linear regression
- Understanding regression results
- ► Non-linear effects in linear regression
- ► Regression tree

Let us first recall simple linear regression. Simple linear regression is a very straightforward simple linear approach for predicting a quantitative response Y on the basis of a single predictor variable X. Mathematically, we can write this linear relationship as

$$Y = \beta_0 + \beta_1 X + \epsilon.$$

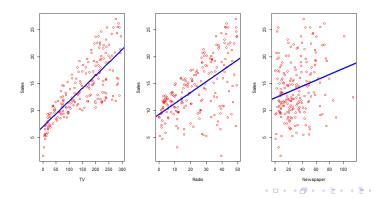
Once we have used our training data to produce estimates $\hat{\beta}_0$ and $\hat{\beta}_1$ for the model coefficients, we can predict future values of \hat{y} on the basis of a particular x by computing

$$\hat{y} = \hat{\beta}_0 + \hat{\beta}_1 x,$$

where \hat{y} indicates a prediction of Y on the basis of X = x.

We will now plot Sales against TV, Radio and Newspaper respectively along with least squares regression line using the plot() and abline() functions.

```
par(mfrow = c(1, 3))
plot(Advertising$TV, Advertising$Sales, col = "red")
abline(Im(formula = Sales~TV , data = Advertising), lwd = 3, col = "
blue") # Write other two plots by yourself.
```



We will start by using the Im() function to fit a simple linear regression model, with TV as the predictor and Sales as the response. The basic syntac is $Im(y\sim x, data)$, where y is the response, x is the predictor, and data is the data set in which these two variables are kept.

```
> library(ISLR)
> attach(Advertising)
> adSLR = Im(formula = Sales"TV)

Error in eval(predvars, data, env) : object 'Sales' not found

> adSLR = Im(formula = Sales"TV, data = Advertising)
> attach(Advertising)
> adSLR = Im(formula = Sales"TV)
```

The command causes an error because R does not know where to find the variables TV and Sales. The next line tells that the variables are in Advertising. If we attach Advertising, R will recognize the variables.

If we type adSLR, some basic information about the model is output.

```
> adSLR

Call:
Im(formula = Sales ~ TV)

Coefficients:
(Intercept) TV
7.03259 0.04754
```

We can use the names() function in order to find out what other pieces of information are stored in adSLR.

```
> names(adSLR)
2 [1] "coefficients" "residuals" "effects" "rank"
    "fitted.values" "assign" "qr" "df.residual"
[9] "xlevels" "call" "terms" "model"
4
```

Detailed information about the model:

```
> summary(adSLR)
1
3
    Call:
    Im(formula = Sales ~TV, data = Advertising)
5
    Residuals:
7
    Min 10 Median 30 Max
    -8.3860 -1.9545 -0.1913 2.0671 7.2124
9
    Coefficients:
11
    Estimate Std. Error t value Pr(>|t|)
    (Intercept) 7.032594 0.457843 15.36 <2e-16 ***
13
    TV 0.047537 0.002691 17.67 <2e-16 ***
    Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 0.1 1
15
17
    Residual standard error: 3.259 on 198 degrees of freedom
    Multiple R-squared: 0.6119, Adjusted R-squared: 0.6099
19
    F-statistic: 312.1 on 1 and 198 DF, p-value: < 2.2e-16
```

Simple linear regression is a useful approach for predicting a response on the basis of a single predictor variable. However, in practice we often have more than one predictor.

Instead of fitting a separate simple linear regression model for each predictor, a better approach is to extend the simple linear regression model so that it can directly accommodate multiple predictors. We can do this by giving each predictor a separate slope coefficient in a single model. In general, suppose that we have p distinct predictors. Then the multiple linear regression model takes the form

$$Y = \beta_0 + \beta_1 X_1 + \cdots + \beta_p X_p + \epsilon.$$

Once we have used our training data to produce estimates $\hat{\beta}_0, \dots, \hat{\beta}_p$ for the model coefficients, we can predict future values of \hat{y} on the basis of a particular x by computing

$$\hat{y} = \hat{\beta}_0 + \hat{\beta}_1 x_1 + \dots + \hat{\beta}_p x_p,$$

where \hat{y} indicates a prediction of Y on the basis of $X = (x_1, \dots, x_p)$.

```
> multipleLR = Im(formula = Sales~TV+Radio+Newspaper , data =
        Advertising)
   > summary (multipleLR)
   Call:
 5 \mid \text{Im}(\text{formula} = \text{Sales} \text{ }^{\sim} \text{ } \text{TV} + \text{Radio} + \text{Newspaper}, \text{ } \text{data} = \text{Advertising})
   Residuals:
       Min
                 10 Median
                                   3<mark>0</mark>
                                           Max
9 - 8.8277 - 0.8908 0.2418 1.1893
                                       2.8292
11 Coefficients:
                 Estimate Std. Error t value Pr(>|t|)
13 (Intercept) 2.938889
                              0.311908 9.422 <2e-16 ***
   TV
                 0.045765 0.001395 32.809 < 2e-16 ***
15 Radio
            0.188530 0.008611 21.893 < 2e-16 ***
   Newspaper -0.001037 0.005871 -0.177
                                                     0.86
17
   Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 0.1 1
19
   Residual standard error: 1.686 on 196 degrees of freedom
21 Multiple R-squared: 0.8972, Adjusted R-squared: 0.8956
   F-statistic: 570.3 on 3 and 196 DF, p-value: < 2.2e-16
```

```
> cor(Advertising[-1])

TV Radio Newspaper Sales

TV 1.0000000 0.05480866 0.05664787 0.7822244
Radio 0.05480866 1.0000000 0.35410375 0.5762226
Newspaper 0.05664787 0.35410375 1.00000000 0.2282990
Sales 0.78222442 0.57622257 0.22829903 1.0000000
```

Since newspaper is not significant, we can refine the model using TV and radio only. With the model $sales=2.9211+0.0458\,TV+0.1880\,radio$, the predicted sales for spending 100 on TV and 20 on radio is 11.26, with the 95% confidence interval [10.99, 11.53] and the 95% predictive interval [7.93, 14.58].

Regression - Interaction Terms

Linear regression models are linear in coefficients $\beta_0, \beta_1, \dots, \beta_p$, while regressors X_1, \dots, X_p can be replaced by any known functions of them.

In this example, the p-values associated with TV, radio, and the interaction term all are statistically significant, and so it is obvious that all three variables should be included in the model. The hierarchical principle states that if we include an interaction in a model, we hierarchical should also include the main effects, even if the p-values associated with principle their coefficients are not significant.

Regression - Polynomial Regression

The data set Auto contains various indices for 387 cars. Let us consider the relationship between mpg (gas mileage in miles power gallon) versus horsepower. Looking at scatter plots, the relationship does not appear to be linear. We fit polynomial regression:

$$mpg = \beta_0 + \beta_1 hpower + \cdots + \beta_p hopower^p + \epsilon$$
, for $p \in \mathbb{N}$.

The results for p = 2 are listed below

```
> polynomialLR = Im(formula = Auto$mpg~Auto$horsepower+I(Auto$horsepower^2), data = Auto)

2 > coefficients(summary(polynomialLR))
Estimate Std. Error t value Pr(>|t|)

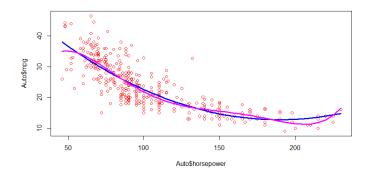
4 (Intercept) 56.9001 1.80042 31.60367 1.74091e-109
Auto$horsepower -0.4662 0.03112 -14.9781 2.2829e-40

6 I(Auto$horsepower^2) 0.00123 0.000122 10.08009 2.1963e-21
```

The function I() is needed since the ^2 has a special meaning in a formula.

Regression - Polynomial Regression

```
> plot(Auto$horsepower, Auto$mpg, col = "red")
2  lines(sort(Auto$horsepower), fitted(polynomialLR)[order(Auto$horsepower)], col='blue', type='l', lwd = 3)
> polynomialLR = lm(formula = Auto$mpg~poly(Auto$horsepower, 5),
    data = Auto)
4  lines(sort(Auto$horsepower), fitted(polynomialLR)[order(Auto$horsepower)], col = 6, type='l', lwd = 3)
```



sort(): sort a vector or factor (partially) into ascending or descending order.
order(): return a permutation which rearranges its first argument into ascending
or descending order. Use ?sort and ?order to learn about those two functions
with more examples.

```
> set.seed(1)
> a = sample(c(1:5))
> a
4 [1] 2 5 4 3 1
> sort(a)
6 [1] 1 2 3 4 5
> order(a)
8 [1] 5 1 4 3 2
```

Question: What is the output of a[order(a)]?

Regression - Regression Tree

Here we fit a regression tree to the Boston data set (part of MASS package). First, we create a training set, and fit the tree to the training data. In the context of a regression tree, the deviance is simply the sum of squared errors for the tree.

```
> library (MASS)
   > library(tree)
    > set . seed (1)
    > train = sample(1:nrow(Boston), nrow(Boston)/2)
    > tree.boston = tree(medv~., Boston, subset = train)
6
    > summary (tree.boston)
8
    Regression tree:
    tree (formula = medv ~ ., data = Boston, subset = train)
10
    Variables actually used in tree construction:
     [1] "Istat" "rm" "dis"
12
    Number of terminal nodes: 8
    Residual mean deviance: 12.65 = 3099 / 245
     Distribution of residuals:
14
    Min. 1st Qu. Median Mean 3rd Qu.
                                                       Max.
16
     -14.10000 \quad -2.04200 \quad -0.05357
                                     0.0000
                                               1.96000 12.60000
```

Regression - Regression Tree

Now we use the cv.tree() to see whether pruning the tree will improve performance. In this case, the most complex tree is selected by cross-validation.

```
> cv.boston = cv.tree(tree.boston)
> plot(cv.boston$size, cv.boston$dev, type = 'b')
```

However, if we wish to prune the tree, we could do so as follows, using prune.tree() function.

```
> prune.boston = prune.tree(tree.boston, best = 5)
> plot(prune.boston)
> text(prune.boston, pretty = 0)
```

Regression - Regression Tree

In keeping with the cross-validaion results, we use the unpruned tree to make predictions on the test set.

```
1 > yhat = predict(tree.boston, newdata = Boston[-train,])
> boston.test = Boston[-train, "medv"]
3 > plot(yhat, boston.test)
> abline(0,1)
5 > mean((yhat-boston.test)^2)
[1] 25.04559
```

