



1 Ordinary Least Squares

1.1 Introduction

Linear regression begins with the simple but profound idea that some observed output or {response} variable, $Y \in \mathbb{R}$, is a function of p input or *regressor* variables x_1, \dots, x_p with the addition of some unknown noise variable ε . Namely,

$$Y = f(x_1, \dots, x_p) + \varepsilon \sim N(0, \text{Var})$$

where the noise is generally assumed to have mean zero and finite variance. The function f is unknown and relates the inputs x_i to the output Y . Our goal is to ascertain what f could be.

In this setting, Y is usually considered to be a random variable while the x_i are considered fixed. Hence, the expected value of Y is in terms of the unknown function f and the regressors:

$$\mathbb{E}(Y|x_1, \dots, x_p) = f(x_1, \dots, x_p).$$

While f can be considered to be in some very general classes of functions, we begin with the standard linear setting. Let $\beta_0, \beta_1, \dots, \beta_p \in \mathbb{R}$. Then, the *multiple regression model* is

$$Y = \beta_0 + \beta_1 x_1 + \dots + \beta_p x_p + \varepsilon = \beta^T X + \varepsilon$$

where $\beta = (\beta_0, \dots, \beta_p)^T$ and $X = (1, x_1, \dots, x_p)^T$. The *simple regression model* is a submodel of the above where $p = 1$, which is

$$Y = \beta_0 + \beta_1 x_1 + \varepsilon,$$

and will be treated concurrently with multiple regression.

In the statistics setting, the parameter vector $\beta \in \mathbb{R}^p$ is unknown. The analyst observes multiple replications of regressor and response pairs, $(X_1, Y_1), \dots, (X_n, Y_n)$ where n is the *sample size*, and wishes to choose a *best* estimate for β based on these n observations. This setup can be concisely written in a vector-matrix form as

$$Y = X\beta + \varepsilon \tag{1.1}$$

where

$$Y = \begin{pmatrix} Y_1 \\ \vdots \\ Y_n \end{pmatrix}, \quad X = \begin{pmatrix} 1 & x_{1,1} & \dots & x_{1,p} \\ 1 & x_{2,1} & \dots & x_{2,p} \\ \vdots & \vdots & \ddots & \vdots \\ 1 & x_{n,1} & \dots & x_{n,p} \end{pmatrix}, \quad \beta = \begin{pmatrix} \beta_0 \\ \vdots \\ \beta_p \end{pmatrix}, \quad \varepsilon = \begin{pmatrix} \varepsilon_1 \\ \vdots \\ \varepsilon_n \end{pmatrix}.$$

Note that $Y, \varepsilon \in \mathbb{R}^n$, $\beta \in \mathbb{R}^{p+1}$, and $X \in \mathbb{R}^{n \times (p+1)}$.

As Y is a random variable, we can compute its mean vector and covariance matrix as follows:

$$\mathbb{E}Y = \mathbb{E}(X\beta + \varepsilon) = X\beta$$

and

$$\text{Var}(Y) = E \left((Y - X\beta)(Y - X\beta)^T \right) = E \left(\varepsilon \varepsilon^T \right) = \text{Var}(\varepsilon) = \sigma^2 I_n.$$

$$\text{Var}(\varepsilon) = \begin{bmatrix} \sigma^2 & 0 & 0 & \dots & 0 \\ 0 & \sigma^2 & 0 & \dots & 0 \\ 0 & 0 & \sigma^2 & \dots & 0 \\ \dots & \dots & \dots & \dots & \dots \\ 0 & 0 & 0 & \dots & \sigma^2 \end{bmatrix}$$

An example of a linear regression is this following study from the New England Journal of Medicine¹ can be found in the code below. This study highlights the correlation between chocolate consumption and Nobel prizes received in 16 different countries.

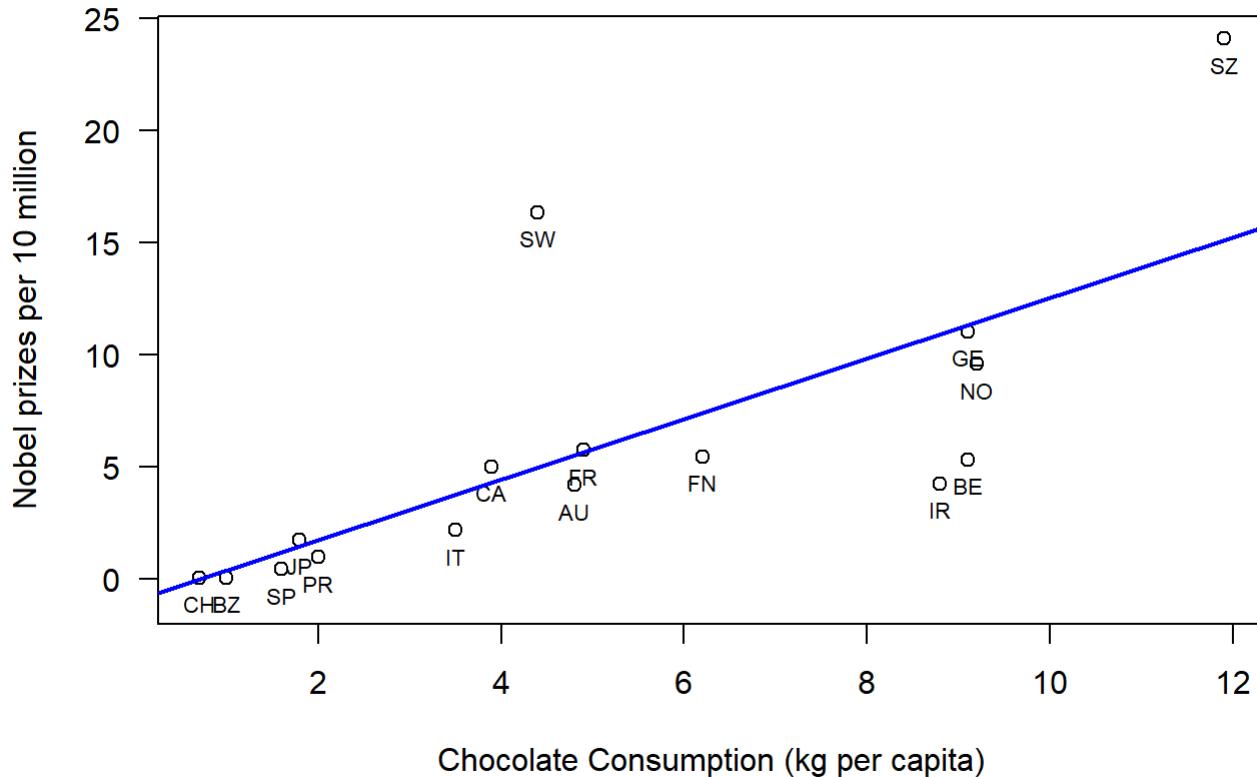
```
# Read in Table
dat = read.table("data/chocoTable.r");

# Plot data
plot(
  dat$choco, dat$nobel,
  xlab="Chocolate Consumption (kg per capita)",
  ylab="Nobel prizes per 10 million",
  las=1,#xlim=c(0,max(dat$choco)+1),
  ylim=c(-1,max(dat$nobel))
);

# Label data
text(
  x=dat$choco, y=dat$nobel,
  labels=dat$abbrev,
  pos=1,cex=0.7
)

# Construct a linear model
lmDat = lm( nobel ~ choco, data = dat );

# plot regression line
abline(lmDat,col='blue',lwd=2)
```



1.1.1 Definitions

Before continuing, we require the following collection of terminology.

The *response* Y and the *regressors* X were already introduced above. These elements comprise the *observed data* in our regression. The *noise* or *error variable* is ε . The entries in this vector are usually considered to be *independent* and identically distributed (iid) random variables with mean zero and finite variance $\sigma^2 < \infty$. Very often, this vector will be assumed to have a multivariate normal distribution: $\varepsilon \sim \mathcal{N}(0, \sigma^2 I_n)$ where I_n is the $n \times n$ identity matrix. The variance σ^2 is also generally considered to be unknown to the analyst.

The unknown vector β is our *parameter vector*. Eventually, we will construct an *estimator* $\hat{\beta}$ from the observed data. Given such an estimator, the *fitted values* are $\hat{Y} := X\hat{\beta}$. These values are what the model believes are the expected values at each regressor.

Given the fitted values, the *residuals* are $r = Y - \hat{Y}$ which is a vector with entries $r_i = Y_i - \hat{Y}_i$. This is the difference between the observed response and the expected response of our model. The residuals are of critical importance to testing how good our model is and will reappear in most subsequent sections.

The four variables in the linear regression model of [Equation 1.1](#) split between whether they are fixed or random variables and between whether or not the analyst knows their value.

	Known	Unknown
Fixed	X	β

Known	Unknown
Random	Y

Lastly, there is the concept of sum of squares. Letting $\bar{Y} = n^{-1} \sum_{i=1}^n Y_i$ be the sample mean for Y , the *total sum of squares* is $SS_{\text{tot}} = \sum_{i=1}^n (Y_i - \bar{Y})^2$, which can be thought of as the total variation of the responses. This can be decomposed into a sum of the *explained sum of squares* and the *residual sum of squares* as follows:

$$SS_{\text{tot}} = SS_{\text{exp}} + SS_{\text{res}} = \sum_{i=1}^n (\hat{Y}_i - \bar{Y})^2 + \sum_{i=1}^n (Y_i - \hat{Y}_i)^2.$$

The explained sum of squares can be thought of as the amount of variation explained by the model while the residual sum of squares can be thought of as a measure of the variation that is not yet contained in the model. The sum of squares gives us an expression for the so called *coefficient of determination*, $R^2 = SS_{\text{exp}}/SS_{\text{tot}} = 1 - SS_{\text{res}}/SS_{\text{tot}} \in [0, 1]$, which is treated as a measure of what percentage of the variation is explained by the given model.

1.2 Point Estimation

$$\hat{\beta} = \mathbf{X}(\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top$$

In the *ordinary least squares setting*, the our choice of estimator is

$$\hat{\beta} = \arg \min_{\tilde{\beta} \in \mathbb{R}^p} \sum_{i=1}^n (Y_i - \underbrace{\mathbf{X}_{i,\cdot} \cdot \tilde{\beta}}_{\text{One term}})^2 \quad (1.2)$$

where $\mathbf{X}_{i,\cdot}$ is the i th row of the matrix \mathbf{X} . In the simple regression setting, this reduces to

$$(\hat{\beta}_0, \hat{\beta}_1) = \arg \min_{(\tilde{\beta}_0, \tilde{\beta}_1) \in \mathbb{R}^2} \sum_{i=1}^n (Y_i - (\tilde{\beta}_0 + \tilde{\beta}_1 x_i))^2.$$

Note that this is equivalent to choosing a $\hat{\beta}$ to minimize the sum of the squared residuals.

It is perfectly reasonable to consider other criterion beyond minimizing the sum of squared residuals. However, this approach results in an estimator with many nice properties. Most notably is the Gauss-Markov theorem:

Theorem 1.1 (Gauss-Markov Theorem) Given the regression setting from [Equation 1.1](#) and that for the errors, $E\varepsilon_i = 0$ for $i = 1, \dots, n$, $\text{Var}(\varepsilon_i) = \sigma^2$ for $i = 1, \dots, n$, and $\text{cov}(\varepsilon_i, \varepsilon_j) = 0$ for $i \neq j$, then the least squares estimator results in the minimal variance over all linear unbiased estimators.
(This is sometimes referred to as the *Best Linear Unbiased Estimator* or BLUE)

Hence, it can be shown that the estimator is unbiased, $E\hat{\beta} = \beta$. Furthermore, as long as the model contains an intercept term β_0 , the constructed least squares line passes through the centre of the data in the sense that the sum of the residuals is zero, $\sum_{i=1}^n r_i = 0$ and that $\bar{Y} = \hat{\beta}\bar{X}$ where $\bar{Y} = n^{-1} \sum_{i=1}^n Y_i$ is the sample mean of the Y_i and where \bar{X} is the vector of column means of the matrix \mathbf{X} .

1.2.1 Derivation of the OLS estimator

The goal is to derive an explicit solution to [Equation 1.2](#). First, consider the following partial derivative:

$$\begin{aligned}\frac{\partial}{\partial \hat{\beta}_k} \sum_{i=1}^n (Y_i - X_{i,\cdot} \cdot \hat{\beta})^2 &= -2 \sum_{i=1}^n (Y_i - X_{i,\cdot} \cdot \hat{\beta}) X_{i,k} \\ &= -2 \sum_{i=1}^n (Y_i - \sum_{j=1}^{p+1} X_{i,j} \hat{\beta}_j) X_{i,k} \\ &= -2 \sum_{i=1}^n Y_i X_{i,k} + 2 \sum_{i=1}^n \sum_{j=1}^{p+1} X_{i,j} X_{i,k} \hat{\beta}_j\end{aligned}$$

The above is the k th entry in the vector $\nabla \sum_{i=1}^n (Y_i - X_{i,\cdot} \hat{\beta})^2$. Hence,

$$\nabla \sum_{i=1}^n (Y_i - X_{i,\cdot} \hat{\beta})^2 = -2X^T Y + 2X^T X \hat{\beta}.$$

Setting this equal to zero results in a critical point at

$$X^T Y = X^T X \hat{\beta}$$

or $\hat{\beta} = (X^T X)^{-1} X^T Y$ assuming $X^T X$ is invertible. Revisiting the terminology in the above definitions sections gives the following table:

Object	Formula
Least Squares Estimator:	$\hat{\beta} = (X^T X)^{-1} X^T Y$
Fitted Values:	$\hat{Y} = X \hat{\beta} = X(X^T X)^{-1} X^T Y$
Residuals:	$r = Y - \hat{Y} = (I_n - X(X^T X)^{-1} X^T) Y$

$P_X \sim \text{rank } (p+1)$
proj. matrix

In the case that $n > p$ and that the columns of X are linearly independent, the matrix $P_X := X(X^T X)^{-1} X^T$ is a rank $p+1$ projection matrix. Similarly, $I_n - P_X$ is the complementary rank $n-p-1$ projection matrix. Intuitively, this implies that the fitted values are the projection on the observed values onto a p -dimensional subspace while the residuals arise from a projection onto the orthogonal subspace. As a result, it can be shown that $\text{cov}(\hat{Y}, r) = 0$. Since \hat{Y} and r are orthogonal

Now that we have an explicit expression for the least squares estimator $\hat{\beta}$, we can show that it is unbiased.

$$E\hat{\beta} = E((X^T X)^{-1} X^T Y) = (X^T X)^{-1} X^T EY = (X^T X)^{-1} X^T X \beta = \beta.$$

Following that, we can compute its variance.

$$\begin{aligned}
\text{Var}(\hat{\beta}) &= \mathbb{E}\left((\hat{\beta} - \beta)(\hat{\beta} - \beta)^T\right) \\
&= \mathbb{E}(\hat{\beta}\hat{\beta}^T) - \beta\beta^T \\
&= \mathbb{E}\left((X^TX)^{-1}X^TY((X^TX)^{-1}X^TY)^T\right) - \beta\beta^T \\
&= (X^TX)^{-1}X^T\mathbb{E}(YY^T)X(X^TX)^{-1} - \beta\beta^T \\
&= (X^TX)^{-1}X^T(\sigma^2I_n + X\beta\beta^TX^T)X(X^TX)^{-1} - \beta\beta^T \\
&= \sigma^2(X^TX)^{-1}.
\end{aligned}$$

Thus far, we have only assumed that ε is a random vector with iid entries with mean zero and variance σ^2 . If in addition, we assumed that ε has a *normal or Gaussian distribution*, then

$$\varepsilon \sim \mathcal{N}(0, \sigma^2 I_n), \quad Y \sim \mathcal{N}(X\beta, \sigma^2 I_n), \quad \text{and } \hat{\beta} \sim \mathcal{N}(\beta, \sigma^2(X^TX)^{-1}).$$

Furthermore, with a little work, one can show that for the fitted values and residuals also have normal distributions in this setting:

$$\hat{Y} \sim \mathcal{N}(X\hat{\beta}, \sigma^2 P_X), \quad \text{and } r \sim \mathcal{N}(0, \sigma^2(I_n - P_X)).$$

Notice that the two above covariance matrices are not generally of full rank. This assumption that the errors follow a normal distribution is a very common assumption to make in practice.

1.2.2 Maximum likelihood estimate under normality

In the previous section, the OLS estimator is derived by minimizing the sum of the squared errors. Now, given the additional assumption that the errors have a normal distribution, we can compute an alternative estimator for β : the maximum likelihood estimate (MLE). We can also use this to simultaneously compute the MLE for σ^2 .

From above we have that $Y \sim \mathcal{N}(X\beta, \sigma^2 I_n)$, and hence the likelihood is

$$L(\beta, \sigma^2; X, Y) = (2\pi\sigma^2)^{-n/2} \exp\left(-\frac{1}{2\sigma^2}(Y - X\beta)^T(Y - X\beta)\right).$$

The log likelihood is then

$$\begin{aligned}
\ell(\beta, \sigma^2; X, Y) &= \log L(\beta, \sigma^2; X, Y) = \\
&= -\frac{n}{2}\log 2\pi - \frac{n}{2}\log \sigma^2 - \frac{1}{2\sigma^2}(Y - X\beta)^T(Y - X\beta).
\end{aligned}$$

This implies that the MLE for β comes from solving

$$0 = \frac{\partial \ell}{\partial \beta} = \frac{\partial}{\partial \beta}(Y - X\beta)^T(Y - X\beta),$$

which is solved by the OLS estimator from above. Hence, the MLE under normality is the least squares estimator.

For the variance term σ^2 , the MLE is similarly found by solving

$$0 = \frac{\partial \ell}{\partial \sigma^2} = -\frac{n}{2}(\sigma^2)^{-1} + \frac{(\sigma^2)^{-2}}{2}(Y - X\beta)^T(Y - X\beta).$$

This occurs for $\hat{\sigma}^2 = n^{-1}(Y - X\hat{\beta})^T(Y - X\hat{\beta})$, which is just the average sum of squares of the residuals: $\hat{\sigma}^2 = n^{-1} \sum_{i=1}^n r_i^2$. However, this is a biased estimator of the variance as the residuals are not independent and have a degenerate covariance matrix of rank $n - p - 1$. Intuitively, this implies that the sum of squared residuals has $n - p - 1$ degrees of freedom resulting in

$$\frac{SS_{\text{res}}}{\sigma^2} = \frac{1}{\sigma^2} \sum_{i=1}^n r_i^2 \sim \chi^2(n - p - 1)$$

$$\hat{\sigma}^2 = \frac{\sigma^2 \cdot n}{n-p-1}$$

and the unbiased estimator of σ^2 being $SS_{\text{res}}/(n - p - 1) = \hat{\sigma}^2(n/(n - p - 1))$.

For a more precise explanation of where this comes from, see [Cochran's Theorem](#) which is beyond the scope of this course.

1.2.2.1 Chocolate-Nobel Data

Running a regression in R on the chocolate consumption vs Nobel prize data from above results in a fitted model

$$(\text{Nobel Prizes}) = -0.991 + 1.3545(\text{Chocolate})$$

This indicates that a 1 kg increase in chocolate consumption per capita corresponds to an expected increase in 1.35 Nobel prizes per 10 million people.

```
# Read in Table
dat = read.table("data/chocoTable.r");
# Construct a linear model
lmDat = lm( nobel ~ choco, dat=dat );
# print summary of model
summary(lmDat);
```

Call:

`lm(formula = nobel ~ choco, data = dat)`

Residuals:

Min	1Q	Median	3Q	Max
-6.6876	-1.6504	-0.5288	0.1484	11.3922

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-0.9910	2.1327	-0.465	0.64932
choco	1.3545	0.3446	3.931	0.00151 **

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 4.666 on 14 degrees of freedom
 Multiple R-squared: 0.5246, Adjusted R-squared: 0.4907
 F-statistic: 15.45 on 1 and 14 DF, p-value: 0.001508

$$RSE = \sqrt{\frac{SSR}{n-p-1}}$$

1.2.3 Proof of the Gauss-Markov Theorem

Proof. Any linear estimator can be written as AY for some non-random matrix $A \in \mathbb{R}^{(p+1) \times n}$. We can in turn write $A = (X^T X)^{-1} X^T + D$ for some matrix $D \in \mathbb{R}^{(p+1) \times n}$. Then, as

$$\begin{aligned} E(AY) &= AX\beta \\ &= [(X^T X)^{-1} X^T + D] X\beta \\ &= \beta + DX\beta, \end{aligned}$$

the unbiased condition implies that $DX\beta = 0$ for any $\beta \in \mathbb{R}^{p+1}$ and hence that $DX = 0$.

Next, we compute the variance of the arbitrary linear unbiased estimator to get

$$\begin{aligned} \text{Var}(AY) &= A\text{Var}(Y)A^T \\ &= \sigma^2 [(X^T X)^{-1} X^T + D] [(X^T X)^{-1} X^T + D]^T \\ &= \sigma^2 [(X^T X)^{-1} + (X^T X)^{-1} X^T D^T + DX(X^T X)^{-1} + DD^T] \\ &= \sigma^2 [(X^T X)^{-1} + DD^T]. \end{aligned}$$

Hence, to minimize the variance, we must minimize DD^T as DD^T is necessarily a positive semi-definite matrix. This is achieved by setting $D = 0$ and arriving at $(X^T X)^{-1} X^T Y$ having minimal variance.

Remark 1.1. Note that DD^T is positive semi-definite for any choice of D as for any $w \in \mathbb{R}^{p+1}$, we have

$$w^T (DD^T) w = (D^T w)^T (Dw) = \|Dw\|_2^2 \geq 0.$$

Remark 1.2. While $\hat{\beta}$ has minimal variance over all unbiased estimators, we can lessen the variance further if we allow for biased estimators. This is considered in many more advanced regression methods such as ridge regression and lasso.

Best Linear Unbiased Estimator

1.3 Hypothesis Testing

1.3.1 Goodness of fit

We now have a model for our data, and in some sense, this model is optimal as it minimizes the squared errors. However, even being optimal, we are still interested in knowing whether or not this is a good model for our data. This is a question of *goodness of fit*. ~ whether our model is 'good'.

The first question to ask is, do any of the regressors provide information about the response in the linear model framework? This can be written mathematically as

$$H_0 : \beta_1 = \dots = \beta_p = 0, \quad H_1 : \exists i \geq 1 \text{ s.t. } \beta_i \neq 0, \quad (1.3)$$

which is asking is there at least one β_i that we can claim is non-zero and hence implies that the regressor x_i has some nontrivial influence over y .

To test this hypothesis, we revisit the explained and residual sums of squares introduced in the Definitions section. Specifically, we already have that $SS_{\text{res}}/\sigma^2 \sim \chi^2(n - p - 1)$ from above. Similarly, $SS_{\text{exp}}/\sigma^2 \sim \chi^2(p)$ under the null hypothesis where $\beta_1 = \dots = \beta_p = 0$, and hence any variation in those terms should be pure noise. Lastly, it can be demonstrated that SS_{res} and SS_{exp} are independent random variables, which intuitively follows from the orthogonality of the fitted values and the errors. Once again, this can be made precise via [Cochran's Theorem](#).

The usual test statistic for the hypothesis in [Equation 1.3](#) is

$$\frac{SS_{\text{exp}}/p}{SS_{\text{res}}/(n - p - 1)} \sim F(p, n - p - 1),$$

which leads to an *F test*. If the test statistic is large, then the explained variation is larger than the noise resulting in a small p-value and a rejection of the null hypothesis.

1.3.1.1 F test on Chocolate-Nobel data

From the final line of the R output from the `summary()` command, we have a test statistic value of 15.45 with degrees of freedom 1 and 14. This results in a very small p-value of 0.001508.

```
summary(lmDat)
```

Call:

```
lm(formula = nobel ~ choco, data = dat)
```

Residuals:

Min	1Q	Median	3Q	Max
-6.6876	-1.6504	-0.5288	0.1484	11.3922

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-0.9910	2.1327	-0.465	0.64932
choco	1.3545	0.3446	3.931	0.00151 **

Signif. codes:	0 ***	0.001 **	0.01 *	0.05 .
	'	'	'	'

Residual standard error: 4.666 on 14 degrees of freedom

Multiple R-squared: 0.5246, Adjusted R-squared: 0.4907

F-statistic: 15.45 on 1 and 14 DF, p-value: 0.001508

If you were to run the regression in R without the intercept term, which is fixing $\beta_0 = 0$, then the result is $\hat{\beta}_1 = 1.22$, a value for the test statistic for the F test of 44.24, now with degrees of freedom 1 and 15, and an even smaller p-value of 7.7×10^{-6} . Typically, regression models always include an intercept term. However, there are some situations where we wish to enforce that an input of zero returns an output of zero.

```
# Construct a linear model
lmDat0 = lm(nobel ~ choco - 1, dat=dat);
# print summary of model
summary(lmDat0);
```

Call:

```
lm(formula = nobel ~ choco - 1, data = dat)
running w/o intercept.
```

Residuals:

Min	1Q	Median	3Q	Max
-6.4991	-1.7891	-1.3237	-0.1955	10.9910

Coefficients:

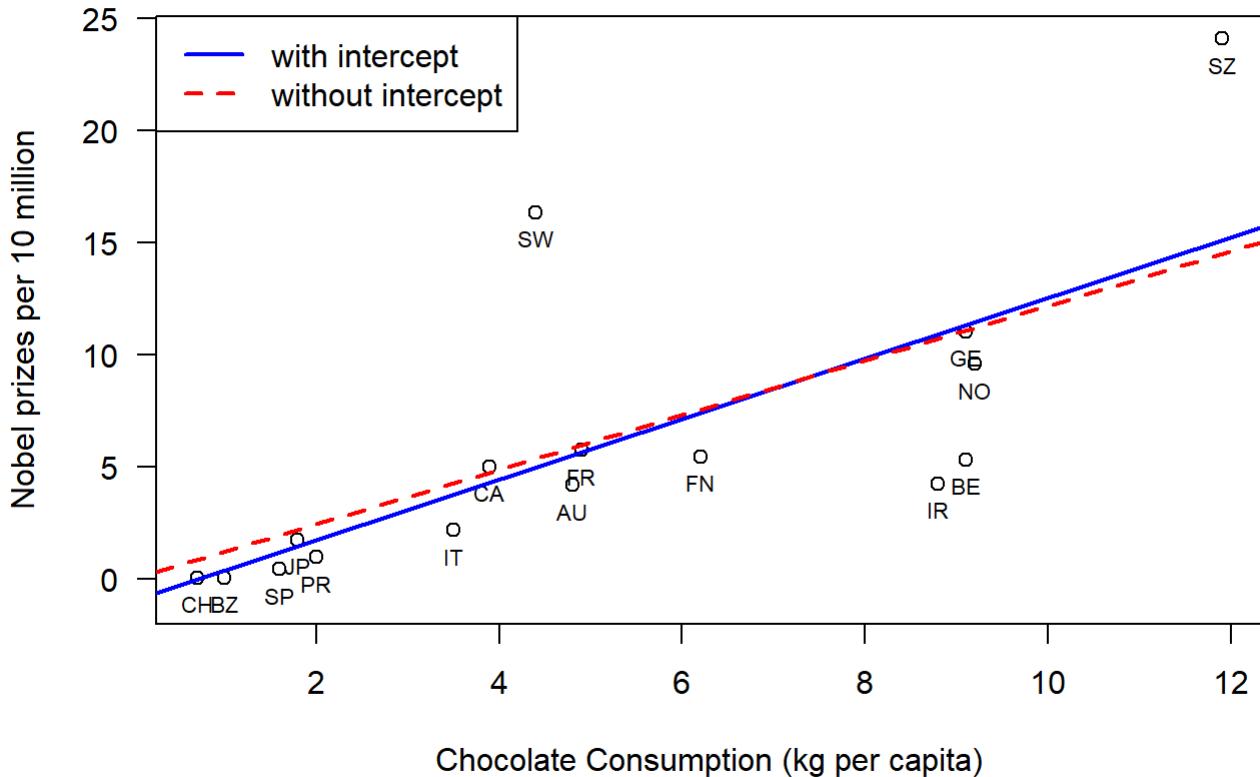
	Estimate	Std. Error	t value	Pr(> t)
choco	1.2205	0.1835	6.651	7.73e-06 ***

Signif. codes:	0 ***	0.001 **	0.01 *	0.05 .
	'	'	'	'

Residual standard error: 4.543 on 15 degrees of freedom
Multiple R-squared: 0.7468, Adjusted R-squared: 0.7299
F-statistic: 44.24 on 1 and 15 DF, p-value: 7.725e-06

```
# Plot data
plot(
  dat$choco, dat$nobel,
  xlab="Chocolate Consumption (kg per capita)",
  ylab="Nobel prizes per 10 million",
  las=1,#xlim=c(0,max(dat$choco)+1),
  ylim=c(-1,max(dat$nobel))
);
# Label data
text(
  x=dat$choco, y=dat$nobel,
  labels=dat$abbrev,
  pos=1,cex=0.7
)

# plot regression line
abline(lmDat,col='blue',lwd=2,lty=1)
abline(lmDat0,col='red',lwd=2,lty=2)
# Add a legend
legend(
  "topleft",legend = c("with intercept","without intercept"),
  col = c("blue","red"),lwd=2,lty=1:2
)
```



1.3.2 Regression coefficients

Given that the previous F test results in a significant p-value, the subsequent question is to ask which of the p regressors are significant? Hence, we have the following hypotheses for $j = 0, 1, \dots, p$.

$$H_{0,j} : \beta_j = 0 \quad H_{1,j} : \beta_j \neq 0.$$

Each individual $\hat{\beta}_j \sim \mathcal{N} \left(\beta_j, \sigma^2 (X^T X)_{j,j}^{-1} \right)$ where $(X^T X)_{j,j}^{-1}$ is the j th entry in the diagonal of $(X^T X)_{j,j}^{-1}$.

Remark 1.3. We will index the entries of the matrix from $0, 1, \dots, p$ to conform with the indexing of the β 's. Note that this is a $(p + 1) \times (p + 1)$ matrix.

Thus, under the null hypothesis that $\beta_j = 0$, we have that

$$\hat{\beta}_j / \sqrt{\sigma^2 (X^T X)_{j,j}^{-1}} \sim \mathcal{N}(0, 1).$$

However, we cannot perform a z test as σ^2 is unknown. To rectify this, the unbiased estimator for σ^2 is used in its place resulting in

$$\frac{\hat{\beta}_j}{\sqrt{(X^T X)_{j,j}^{-1} S S_{\text{res}} / (n - p - 1)}} \sim t(n - p - 1),$$

and a t test can be performed. If the value of the test statistic is large, then there may be sufficient evidence to reject the null that $\beta_j = 0$. The denominator is often referred to as the *standard error*. To simplify future formulae, this will be denoted as $\text{se}(\beta_j)$.

It is worth noting that this test looks for significant influence of the j th regressor on the response given all of the other regressors. Hence, it quantifies the marginal as opposed to the absolute effect of that variable on the model. These ideas will be investigated further when discussing variable selection later in this book. However, as a quick word of caution, when p hypothesis tests are performed, the analyst needs to consider [multiple testing corrections](#).

1.3.2.1 t test on Chocolate-Nobel data

The R commands `lm()` and `summary()` will return a table of regression coefficients, t test statistics and p-values associated with each coefficient. For the Chocolate-Nobel prize data, the table looks like

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-0.9910	2.1327	-0.465	0.64932
choco	1.3545	0.3446	3.931	0.00151

```
summary(lmData)
```

Call:

```
lm(formula = nobel ~ choco, data = dat)
```

Residuals:

Min	1Q	Median	3Q	Max
-6.6876	-1.6504	-0.5288	0.1484	11.3922

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
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choco	1.3545	0.3446	3.931	0.00151 **

Signif. codes:	0 ***	0.001 **	0.01 *	0.05 .
	'***'	'**'	'.'	'.'

Residual standard error: 4.666 on 14 degrees of freedom

Multiple R-squared: 0.5246, Adjusted R-squared: 0.4907

F-statistic: 15.45 on 1 and 14 DF, p-value: 0.001508

1.3.3 Partial F-test

In the previous two sections, we first tested as to whether or not there exists at least one β_j , $j = 1, \dots, p$, that is non-zero. Then, we tested whether or not a specific β_j is non-zero. The next logical question is whether or not some collection of β_j 's of size strictly between 1 and p has a non-zero element. That is, for a fixed q ,

$$H_0 : \underbrace{\beta_{p-q+1} = \dots = \beta_p = 0}_{\text{lost in prediction = 0.}} \quad H_1 : \exists i \geq p - q + 1 \text{ s.t. } \beta_i \neq 0. \quad (1.4)$$

at least 1 of the q's are significant.

Here, we are comparing two different models, which are the partial and full models, respectively,

$$Y = \beta_{0:p-q} X + \varepsilon, \quad \text{and} \quad Y = \beta_{0:p-q} X + \underbrace{\beta_{p-q+1:p} X}_{\text{Extra q added}} + \varepsilon,$$

and want to know whether the final q regressors add any significant explanation to our model given the other

$p - q$. For the above notation,

$$\beta_{i:j} = (0, \dots, 0, \beta_i, \beta_{i+1}, \dots, \beta_j, 0, \dots, 0)^T.$$

To run the hypothesis test in [Equation 1.4](#), we would have to compute the least squares estimator in the partial model, $\hat{\beta}_{1:p-q}$, and the standard least squares estimator in the full model, $\hat{\beta}$. Then, we will have to compute the additional explained sum of squares gained from adding the q extra regressors to our model, which is

$$SS_{\text{exp}}(\beta_{p-q+1:p} | \beta_{1:p-q}) = SS_{\text{exp}}(\beta) - SS_{\text{exp}}(\beta_{1:p-q}),$$

the explained sum of squares from the full model minus the explained sum of squares from the partial model.

Similarly to the full F-test from above, we have under the null hypothesis that

$$SS_{\text{exp}}(\beta_{p-q+1:p} | \beta_{1:p-q}) / \sigma^2 \sim \chi^2(q). \text{ Hence,}$$

$$\frac{SS_{\text{exp}}(\beta_{p-q+1:p} | \beta_{1:p-q}) / q}{SS_{\text{res}} / (n - p - 1)} \sim F(q, n - p - 1),$$

so if this test statistic is large, then we have evidence to suggest that at least one of the additional q regressors adds some explanatory power to our model.

1.4 Interval Estimators

1.4.1 Confidence Intervals

Confidence intervals play a complementary role with hypothesis testing. From the development of the above test for an individual β_j , we have that

$$\frac{\hat{\beta}_j - \beta_j}{\text{se}(\beta_j)} \sim t(n - p - 1),$$

Hence, a $1 - \alpha$ confidence interval for the parameter β_j is

$$\hat{\beta}_j - t_{\alpha/2, n-p-1} \text{se}(\beta_j) \leq \beta \leq \hat{\beta}_j + t_{\alpha/2, n-p-1} \text{se}(\beta_j)$$

where $t_{\alpha/2, n-p-1} \in \mathbb{R}^+$ is such that $P(T \leq t_{\alpha/2, n-p-1}) = \alpha/2$ when $T \sim t(n - p - 1)$.

While the above can be used to produce a confidence interval for each individual parameter, combining these intervals will not result in a $1 - \alpha$ confidence set for the entire parameter vector. To construct such a confidence region, a little more care is required. Also, we will construct a confidence set for the entire vector $(\beta_0, \beta_1, \dots, \beta_p)$, which results in $p + 1$ degrees of freedom in what follows. As $\hat{\beta} \sim \mathcal{N}(\beta, \sigma^2(X^T X)^{-1})$ we have that

$$\sigma^{-2}(\hat{\beta} - \beta)^T X^T X (\hat{\beta} - \beta) \sim \chi^2(p + 1).$$

From before, we have that $SS_{\text{res}} / \sigma^2 \sim \chi^2(n - p - 1)$. Hence

$$\frac{(\hat{\beta} - \beta)^T X^T X (\hat{\beta} - \beta) / (p + 1)}{SS_{\text{res}} / (n - p - 1)} \sim F(p + 1, n - p - 1).$$

Thus, a $1 - \alpha$ confidence ellipsoid can be constructed as

$$\frac{(\hat{\beta} - \beta)^T X^T X (\hat{\beta} - \beta)/(p+1)}{SS_{\text{res}}/(n-p-1)} \leq F_{\alpha, p+1, n-p-1}.$$

A 95% and a 99% confidence ellipsoid for the Chocolate-Nobel prize data is displayed in the code below. Notice that both ellipses contain $\hat{\beta}_0 = 0$ which had a t statistic p-value of 0.649. Meanwhile neither contain $\hat{\beta}_1 = 0$ whose p-value was the very significant 0.0015. The confidence ellipses were plotted with help from the R library `ellipse`.

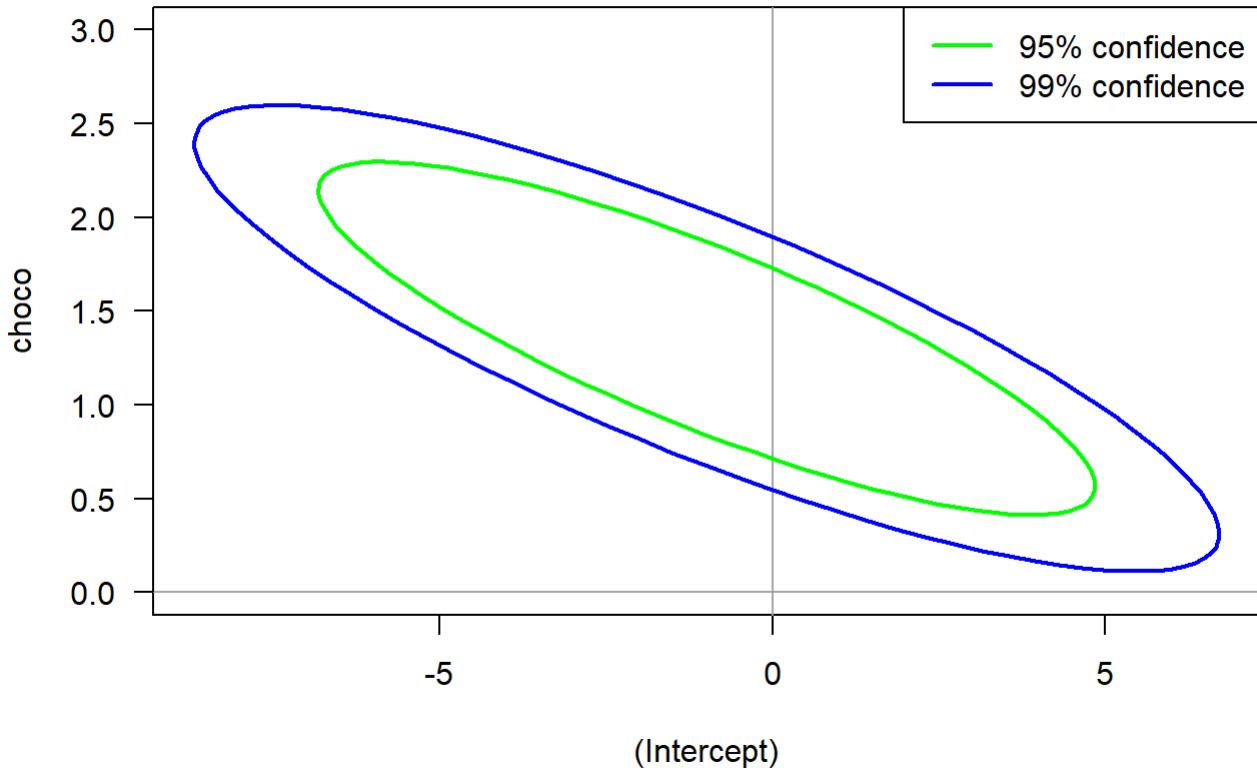
```
library(ellipse)
```

Attaching package: 'ellipse'

The following object is masked from 'package:graphics':

```
pairs
```

```
plot(
  ellipse(lmDat, level=.99), type='l',
  ylim=c(0,3), col='blue', lwd=2, las=1
);
abline(h=0, v=0, col='darkgray');
lines(ellipse(lmDat, level=.99), type='l', col='blue', lwd=2);
lines(ellipse(lmDat, level=.95), type='l', col='green', lwd=2);
legend(
  "topright", legend = c("95% confidence", "99% confidence"),
  col = c("green", "blue"), lwd = 2
)
```



1.4.2 Prediction Intervals for an expected observation ?

Given the least squares model, the analyst may be interested in estimating the expected value of Y have some specific input $x = (1, x_1, \dots, x_p)$. Our new random variable is $\hat{Y}_0 = \hat{\beta} \cdot X$ where X is fixed and $\hat{\beta}$ is random. Of course, the expected value is just

$$E(\hat{Y}_0 | X = x) = E\hat{\beta} \cdot x = \beta_0 + \sum_{i=1}^p \beta_i x_i.$$

To find a $1 - \alpha$ interval estimate for \hat{Y}_0 at $X = x$, recall once again that $\hat{\beta} \sim \mathcal{N}(\beta, \sigma^2(X^T X)^{-1})$. Thus,

$$\hat{Y}_0 | (X = x) \sim \mathcal{N}(\beta \cdot x, \sigma^2 x^T (X^T X)^{-1} x).$$

Hence,

$$\frac{\hat{\beta} \cdot x - E(\hat{Y}_0 | X = x)}{\sqrt{\sigma^2 x^T (X^T X)^{-1} x}} \sim \mathcal{N}(0, 1),$$

and

$$\frac{\hat{\beta} \cdot x - E(\hat{Y}_0 | X = x)}{\sqrt{(SS_{\text{res}}/(n-p-1)) x^T (X^T X)^{-1} x}} \sim t(n-p-1),$$

which results in the following $1 - \alpha$ confidence interval:

$$\begin{aligned}\hat{\beta} \cdot x - t_{\alpha/2, n-p-1} \sqrt{\frac{SS_{\text{res}}}{n-p-1} x^T (X^T X)^{-1} x} &\leq \\ &\leq E(\hat{Y}_0 | X = x) = \beta \cdot x \leq \\ &\leq \hat{\beta} \cdot x + t_{\alpha/2, n-p-1} \sqrt{\frac{SS_{\text{res}}}{n-p-1} x^T (X^T X)^{-1} x}.\end{aligned}$$

1.4.3 Prediction Intervals for a new observation

In the previous subsection, we asked for a confidence interval for the expected value of the response given a new vector of regressors, which was a confidence interval for $E(\hat{Y}_0 | X = x) = \beta \cdot x$ based on $\hat{\beta} \cdot x$. Now, we want to determine a confidence interval for the future response given a vector of regressors. That is, we want an interval for $Y_0 = \beta \cdot x + \varepsilon_0 \sim \mathcal{N}(\beta \cdot x, \sigma^2)$, but, as usual, β unknown.

To circumvent this, note that

$$Y_0 - \hat{Y}_0 = (\beta \cdot x + \varepsilon_0) - \hat{\beta} \cdot x \sim \mathcal{N}(0, \sigma^2(1 + x^T (X^T X)^{-1} x)),$$

because the variances of ε_0 and $\hat{\beta} \cdot x$ sum as these are independent random variables. Hence, applying the usual rearrangement of terms and replacement of σ^2 with $SS_{\text{res}}/(n-p-1)$ results in

$$\begin{aligned}\hat{\beta} \cdot x - t_{\alpha/2, n-p-1} \sqrt{\frac{(1 + x^T (X^T X)^{-1} x) SS_{\text{res}}}{n-p-1}} &\leq Y_0 \leq \\ &\leq \hat{\beta} \cdot x + t_{\alpha/2, n-p-1} \sqrt{\frac{(1 + x^T (X^T X)^{-1} x) SS_{\text{res}}}{n-p-1}}.\end{aligned}$$

To demonstrate these prediction intervals, we once again consider the Chocolate-Nobel prize data for both the expected mean and for a new observation.

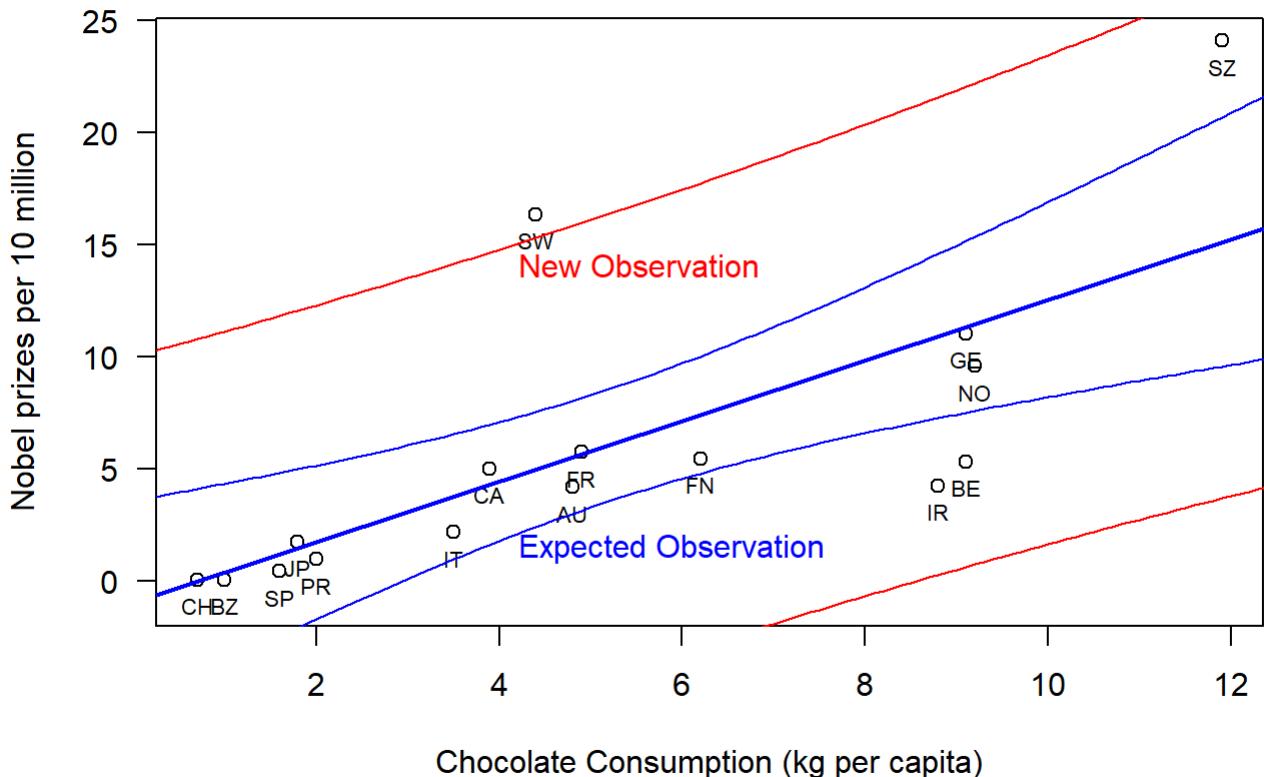
```
# Plot data
plot(
  dat$choco, dat$nobel,
  xlab="Chocolate Consumption (kg per capita)",
  ylab="Nobel prizes per 10 million",
  las=1, #xlim=c(0,max(dat$choco)+1),
  ylim=c(-1,max(dat$nobel))
);
# Label data
text(
  x=dat$choco, y=dat$nobel,
  labels=dat$abbrev,
  pos=1, cex=0.7
)
# plot regression line
abline(lmDat, col='blue', lwd=2)

# plot 95% confidence and prediction intervals
tt = seq(0, 13, 0.1)
prDat = predict(
```

```

lmDat,newdata=data.frame(choco=tt),
interval='confidence',level=0.95
)
prDat2 = predict(
  lmDat,newdata=data.frame(choco=tt),
  interval='prediction',level=0.95
)
lines(tt,prDat[,2],col='blue')
lines(tt,prDat[,3],col='blue')
lines(tt,prDat2[,2],col='red')
lines(tt,prDat2[,3],col='red')
text(
  c(4,4),c(1.5,14),
  labels=c("Expected Observation","New Observation"),
  col=c("blue","red"),pos=4
)

```



1.5 Indicator Variables and ANOVA

1.5.1 Indicator variables

Thus far, we have considered models of the form

$$y = \beta_0 + \beta_1 x_1 + \dots + \beta_p x_p + \varepsilon$$

where the regressors $x_1, \dots, x_p \in \mathbb{R}$ can take on any real value. However, very often in practice, we have

regressors that take on categorical values. For example, male vs female, employed vs unemployed, treatment vs placebo, Edmonton vs Calgary, etc. When there is a binary choice as in these examples, we can choose one category to correspond to zero and the other category to correspond to one.

As an example, consider

$$y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \varepsilon \quad (1.5)$$

where $x_1 \in \mathbb{R}$ and $x_2 \in \{0, 1\}$. Then, we effectively have two models:

$$\begin{aligned} y &= \beta_0 + \beta_1 x_1 + 0 + \varepsilon \\ y &= \beta_0 + \beta_1 x_1 + \beta_2 + \varepsilon = (\beta_0 + \beta_2) + \beta_1 x_1 + \varepsilon. \end{aligned}$$

What we have is two models with the same slope β_1 but with two different intercepts β_0 and $\beta_0 + \beta_2$, which are two parallel lines.

Remark 1.4. A first thought is to merely split the data and train two separate models. However, we want to use the entire dataset at once specifically to estimate the common slope β_1 with as much accuracy as possible.

While the range of the regressors has changed, we will fit the least squares estimate to the model precisely as before. Now, considering the model in [Equation 1.5](#), assume that we have m samples with $x_2 = 0$ and n samples with $x_2 = 1$. Our design matrix takes on a new form:

$$Y = \begin{pmatrix} Y_1 \\ \vdots \\ Y_m \\ Y_{m+1} \\ \vdots \\ Y_{m+n} \end{pmatrix}, \quad X = \begin{pmatrix} 1 & x_1 & 0 \\ \vdots & \vdots & \vdots \\ 1 & x_m & 0 \\ 1 & x_{m+1} & 1 \\ \vdots & \vdots & \vdots \\ 1 & x_{m+n} & 1 \end{pmatrix}, \quad \beta = \begin{pmatrix} \beta_0 \\ \beta_1 \\ \beta_2 \end{pmatrix}, \quad \varepsilon = \begin{pmatrix} \varepsilon_1 \\ \vdots \\ \varepsilon_{m+n} \end{pmatrix}.$$

However, the least squares estimate is computed as before as $\hat{\beta} = (X^T X)^{-1} X^T Y$. Furthermore, we can perform hypothesis tests on the fitted model such as

$$H_0 : \beta_2 = 0 \quad H_1 : \beta_2 \neq 0,$$

which is equivalently asking whether or not the regression lines have the same intercept.

Models can be expanded to include multiple indicator variables as long as the matrix $X^T X$ is still invertible. For example, let's suppose we want to look at wages in Alberta with respect to age but partitioned for male vs female and for Edmonton vs Calgary. Then, the model would look like

$$(\text{wage}) = \beta_0 + \beta_1(\text{age}) + \beta_2(\text{Is male?}) + \beta_3(\text{Is from Edmonton?}).$$

In the silly case that our data only consisted of men from Calgary and women from Edmonton, then the final regressor is redundant and $X^T X$ will not be invertible. While this extreme case should not occur, it is possible to have an imbalance in the categories, which we will discuss later.

1.5.2 ANOVA

ANOVA, or the Analysis of Variance, is a slightly overloaded term in statistics. We already considered ANOVA tables when comparing nested models in the hypothesis tests. However, ANOVA can also be used in the setting of the so-called [One-Way Analysis of Variance](#). In this case, we want to compare k samples for equality of the

means. For example, we take height measurements from randomly selected citizens from different countries and ask whether or not there is significant evidence to reject the claim that all nations have roughly the same height distribution.

The reason for discussing ANOVA in these notes is that it can be written in a linear regression context as follows. Imagine that we have k different groups of observations with sample sizes $n_j, j = 1, \dots, k$ for each group. Let $y_{i,j}$ be the i th observation from the j th group where $i \in \{1, \dots, n_j\}$ and $j \in \{1, \dots, k\}$. The model is

$$y_{i,j} = \mu_j + \varepsilon_{i,j},$$

which is each observation is just some group mean, μ_j , with the addition of random noise.

From here, one can show that the fitted values as just $\hat{y}_{i,j} = n_j^{-1} \sum_{l=1}^{n_j} y_{l,j}$, which is the j th sample mean. Then, an F-test can be performed similar to what we did above with F tests to test

$$H_0 : \mu_1 = \dots = \mu_k \quad H_1 : \exists j_1 \neq j_2 \text{ s.t. } \mu_{j_1} \neq \mu_{j_2}.$$

To reformulate the model to align with our F-test from before, we rewrite it as a linear regression with indicator variables for the regressors

$$y = \beta_0 + \beta_1 x_1 + \dots + \beta_{k-1} x_{k-1} + \varepsilon_{i,j}$$

with $\beta_0 = \mu_k$ and $\beta_j = \mu_j - \mu_k$ for $j = 1, \dots, k-1$. Then, we can test for whether or not there exists at least one $\beta_j \neq 0$ for $j = 1, \dots, k-1$. Here, the degrees of freedom for the explained sum of squares is $k-1$ and the degrees of freedom for the residual sum of squares is $N - (k-1) - 1 = N - k$ with $N = \sum_{j=1}^k n_j$.

If all of the n_j are equal, this reduces to $k(n-1)$.

In this case, the vector Y and the design matrix X will take on the form

$$Y = \begin{pmatrix} y_{1,1} \\ y_{2,1} \\ \vdots \\ y_{n_1,1} \\ y_{1,2} \\ \vdots \\ y_{n_k,k} \end{pmatrix}, \quad X = \begin{pmatrix} 1 & 1 & 0 & \dots & 0 \\ 1 & 1 & 0 & \dots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & 1 & 0 & \dots & 0 \\ 1 & 0 & 1 & \dots & 0 \\ 1 & 0 & 1 & \dots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & 0 & 0 & \dots & 1 \\ 1 & 0 & 0 & \dots & 1 \\ 1 & 0 & 0 & \dots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & 0 & 0 & \dots & 0 \\ 1 & 0 & 0 & \dots & 0 \end{pmatrix}.$$

1.6 Data Example: Exam Grades

To illustrate the topics discussed in this chapter, we will consider a dataset of course grades for a linear regression class. This dataset consists of $n = 56$ undergraduate students who are either in their 3rd year (22

students) or 4th year (33 students) of university studies. The year of study acts as a binary variable as discussed in the previous section. This dataset also has columns corresponding to the students overall mark on written homework assignments, online coding assignments, the midterm exam, and the final exam. In what follows, we will see what variables if any can be used to predict the final exam grade and, furthermore, if there is a difference between the final exam grades for the 3rd year and 4th year students.

```
# Read in Data
exams = read.csv("data/statGrades.csv")

# Tell R that "year" is a categorical variable
exams$year <- as.factor(exams$year)

# Look at the data
head(exams)
```

	year	written	online	midterm	final
1	4th Year	80.668	83.84127	70.00000	56.4948
2	3rd Year	96.668	100.00000	50.00000	54.0000
3	4th Year	90.000	93.73737	50.00000	43.5024
4	3rd Year	98.000	98.57143	56.66667	62.9964
5	3rd Year	46.000	48.72727	50.00000	43.5024
6	4th Year	75.334	96.34921	93.33333	81.0000

When comparing two independent groups, we can use the classic two sample t test using `t.test()` in R. The default is to assume that the variances between the two groups are not equal, which is the so-called Welch's t-test. Otherwise, we can tell the function to treat the variances as equal by setting `var.equal=T`.

```
t.test(
  final ~ year,
  data = exams
)
```

Welch Two Sample t-test

```
data: final by year
t = -0.42107, df = 43.775, p-value = 0.6758
alternative hypothesis: true difference in means between group 3rd Year and group 4th
Year is not equal to 0
95 percent confidence interval:
-11.709616  7.662733
sample estimates:
mean in group 3rd Year mean in group 4th Year
      65.67292           67.69636
```

```
t.test(
  final ~ year,
  data = exams,
  var.equal=T
)
```

Two Sample t-test

```
data: final by year
t = -0.43015, df = 54, p-value = 0.6688
alternative hypothesis: true difference in means between group 3rd Year and group 4th
Year is not equal to 0
95 percent confidence interval:
-11.454430  7.407547
sample estimates:
mean in group 3rd Year mean in group 4th Year
65.67292          67.69636
```

We can also view a two sample t test as a linear regression of the response variable (final exam grade) on the dummy variable (student year). The regression equation becomes

$$(\text{final exam grade}) = \beta_0 + \beta_1(\text{student year}) + \varepsilon$$

What we can see from the output below is that this model is equivalent to the two sample t test assuming equal (homogeneous) variances among the two samples. In the end, there is no noticeable difference between the performance of 3rd and 4th year students.

```
# A two-sample test
md.year = lm(
  final ~ year,
  data = exams
)
summary(md.year)
```

Call:

```
lm(formula = final ~ year, data = exams)
```

Residuals:

Min	1Q	Median	3Q	Max
-31.700	-14.444	1.318	12.550	36.322

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	65.673	3.611	18.19	<2e-16 ***
year4th Year	2.023	4.704	0.43	0.669

Signif. codes:	0 ***	0.001 **	0.01 *	0.05 .
	'	'	'	'
	'	'	'	'

Residual standard error: 17.32 on 54 degrees of freedom

Multiple R-squared: 0.003415, Adjusted R-squared: -0.01504

F-statistic: 0.185 on 1 and 54 DF, p-value: 0.6688

However, the `lm()` function allows us to fit more complex models than just a two sample comparison. Here, we use the students performance on the written homework, the online coding homework, and the midterm exam as predictors of performance on the final exam. The result below is that the midterm mark is a strongly significant predictor of the final exam mark with an estimated coefficient of 0.953 and that the written assignment mark is a weak but possibly still relevant predictor of the final exam mark.

```
# Fit Linear Model
md.exams = lm(
  final ~ year+written+online+midterm,
  data = exams
)
summary(md.exams)
```

Call:

```
lm(formula = final ~ year + written + online + midterm, data = exams)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-20.2987	-7.0874	-0.8767	6.4742	31.3874

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)							
(Intercept)	-14.3982	12.6807	-1.135	0.2615							
year4th Year	1.5523	3.0561	0.508	0.6137							
written	0.3707	0.1962	1.889	0.0646 .							
online	-0.2024	0.2143	-0.944	0.3494							
midterm	0.9527	0.1113	8.558	1.97e-11 ***							

Signif. codes:	0	'***'	0.001	'**'	0.01	'*'	0.05	'. '	0.1	' '	1

Residual standard error: 11.24 on 51 degrees of freedom

Multiple R-squared: 0.6032, Adjusted R-squared: 0.5721

F-statistic: 19.39 on 4 and 51 DF, p-value: 9.478e-10

We can also consider the simpler regression model that only takes the midterm mark into account as a predictor variable. The fitted model has a slope parameter of 0.938 and an intercept not significantly different from zero. This indicates that the student's final exam mark was on average $0.938 \times (\text{midterm mark})$, or simply that students had slightly lower grades on average on the final exam compared to the midterm exam. The cyan coloured region indicates those who had a higher midterm mark than final exam mark.

```
md.exams0 = lm(
  final ~ midterm,
  data = exams
)
summary(md.exams0)
```

Call:

```
lm(formula = final ~ midterm, data = exams)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-16.758	-8.683	-2.942	7.620	31.621

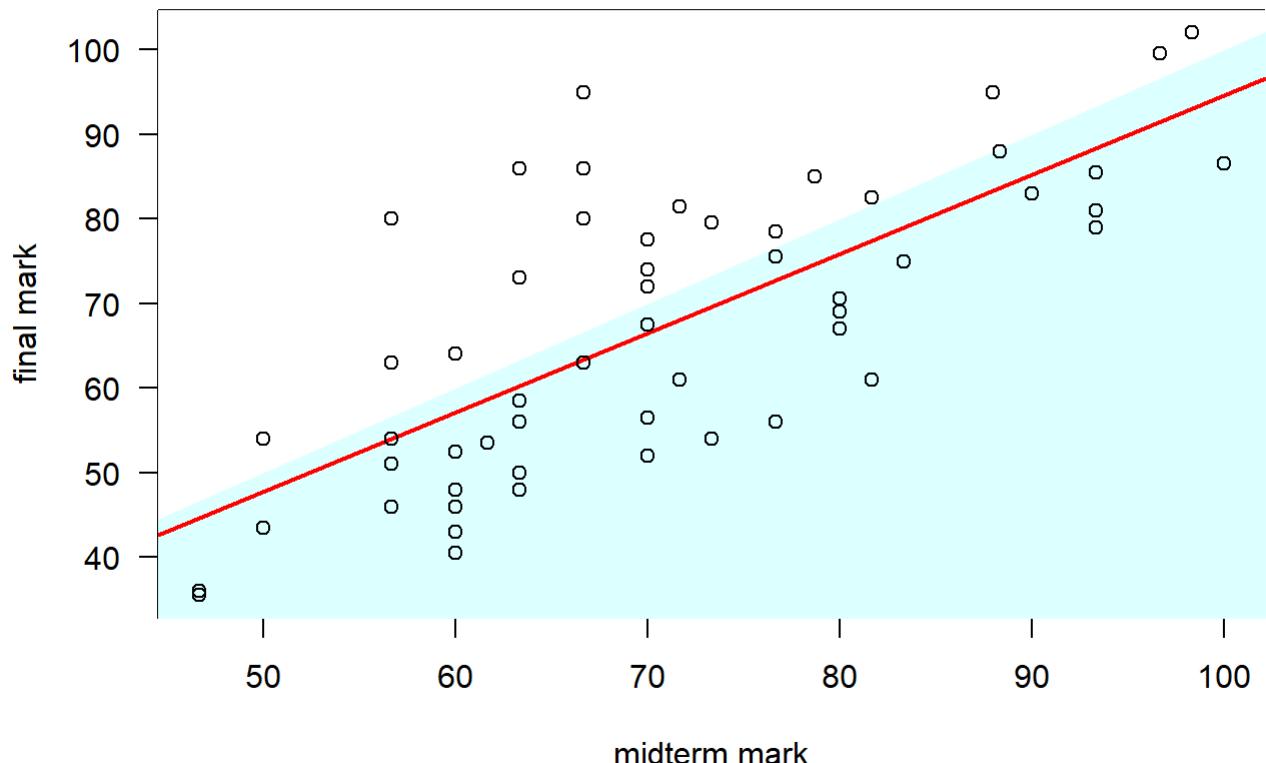
Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	0.8376	8.1986	0.102	0.919

```
midterm      0.9381     0.1144   8.201 4.68e-11 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Residual standard error: 11.58 on 54 degrees of freedom
 Multiple R-squared: 0.5547, Adjusted R-squared: 0.5464
 F-statistic: 67.25 on 1 and 54 DF, p-value: 4.684e-11

```
plot(
  exams$midterm, exams$final, las=1,
  xlab="midterm mark", ylab="final mark"
)
polygon(
  c(0,200,200),c(0,200,0),col = 'lightcyan',
  border = NA
)
polygon(
  c(0,0,200),c(0,200,200),col = 'white',
  border = NA
)
abline(md.exams0,col='red',lwd=2)
points(exams$midterm,exams$final)
```



We can now imagine a hypothetical student in their 4th year who got a 95% on the written assignments, 85% on the online assignments, and an 83% on the midterm exam. Using the `predict()` function in R, we get an expected final exam grade of 84.24%. Going further, a 95% prediction interval for a new value gives a very wide

range from 60.3% to 108.17%. This is where the model believes the final exam grade will lie for our hypothetical student.

```
predict(
  md.exams,
  newdata=data.frame(
    year="4th Year",written=95,online=85,midterm=83
  ), interval="prediction", level=0.95
)
```

	fit	lwr	upr
1	84.24118	60.30866	108.1737

We can use the `anova()` function to compare nested models, which will perform a partial F-test. In this example, we can compare the model that only includes the `year` variable to a model that contains model `year` and `midterm` to a model that contains all four predictor variables.

In the ANOVA table below, we have that comparing model 1 to model 2 results is a very significant p-value indicating that including `midterm` as a predictor results in a large decrease in the residual sum of squares (column 2 in the table below). Secondly, we compare model 2 to model 3 and get a weak but possibly significant p-value of 0.0597. This gives some weak evidence that including `written` and `online` assignment marks may further improve the fit of the model.

Note that the degrees of freedom for the partial F-tests performed can be found in columns 1 and 3 in the ANOVA table. These are the degrees of freedom for the residual sum of squares (column 1) and the explained sum of squares (column 3).

```
md.exams1 = lm(
  final ~ year+midterm,
  data = exams
)
anova(md.year,md.exams1,md.exams)
```

Analysis of Variance Table

Model 1: final ~ year					
Model 2: final ~ year + midterm					
Model 3: final ~ year + written + online + midterm					
Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	16195.2				
2	7200.8	1	8994.4	71.1454	3.058e-11 ***
3	6447.5	2	753.3	2.9792	0.05974 .

Signif. codes:	0	'***'	0.001	'**'	0.01
	*	'*'	0.05	.'	0.1
	'	'	1		

If we incorrectly try to compare non-nested models, we get erroneous results in our ANOVA table.

```
md.exams2 = lm(
  final ~ year+written+online,
  data = exams
```

```
)
anova(md.exams1,md.exams2)
```

Analysis of Variance Table

```
Model 1: final ~ year + midterm
Model 2: final ~ year + written + online
  Res.Df    RSS Df Sum of Sq F Pr(>F)
1      53  7200.8
2      52 15706.6  1   -8505.8
```

The general equation for a linear regression is $Y = X\beta + \varepsilon$. To recover the design matrix X , we can use the function `model.matrix()` in R.

Notice that the column corresponding to `year` is now encoded as a binary variable with a 1 indicating 4th year and a 0 indicating 3rd year.

```
model.matrix(md.exams)
```

	(Intercept)	year4th	Year	written	online	midterm
1	1	1	1	80.66800	83.84127	70.00000
2	1	1	0	96.66800	100.00000	50.00000
3	1	1	1	90.00000	93.73737	50.00000
4	1	1	0	98.00000	98.57143	56.66667
5	1	1	0	46.00000	48.72727	50.00000
6	1	1	1	75.33400	96.34921	93.33333
7	1	1	0	82.66600	89.77778	100.00000
8	1	1	0	80.33400	96.34921	60.00000
9	1	1	0	96.66600	96.34921	56.66667
10	1	1	0	82.66600	87.46609	70.00000
11	1	1	1	72.66667	100.00000	56.66667
12	1	1	0	95.33600	98.57143	63.33333
13	1	1	1	98.00000	100.00000	93.33333
14	1	1	0	92.00000	100.00000	88.33333
15	1	1	1	63.33200	74.66667	63.33333
16	1	1	1	98.00200	98.57143	76.66667
17	1	1	1	86.00000	85.73737	60.00000
18	1	1	1	100.00000	97.77778	73.33333
19	1	1	1	96.66600	98.57143	60.00000
20	1	1	1	100.00000	93.10245	81.66667
21	1	1	1	97.33400	98.57143	46.66667
22	1	1	0	98.66600	100.00000	98.33333
23	1	1	1	96.00000	100.00000	80.00000
24	1	1	1	95.33400	97.77778	60.00000
25	1	1	0	85.33400	94.53102	66.66667
26	1	1	1	77.33267	93.10245	81.66667
27	1	1	1	94.00000	97.14286	87.96667
28	1	1	1	96.00000	100.00000	70.00000
29	1	1	1	86.66600	93.14286	73.33333
30	1	1	1	82.66400	88.92064	70.00000
31	1	1	0	85.33400	91.55556	83.33333
32	1	1	0	86.66533	94.34921	96.66667
33	1	1	1	99.33400	100.00000	63.33333
34	1	1	0	91.33400	95.95960	70.00000
35	1	1	0	75.33400	93.73737	76.66667

```
36      1      0  76.66600  87.06205  80.00000
37      1      1  60.62733  92.69841  71.66667
38      1      1  85.33400  92.34921  66.66667
39      1      1  74.66800  79.79221  60.00000
40      1      1  92.00000  98.00000  78.70000
41      1      1  70.66400  88.34921  56.66667
42      1      1  93.33400  100.00000 71.66667
43      1      0  98.66800  85.44012  63.33333
44      1      1  97.33400  100.00000 56.66667
45      1      0  71.99800  93.77778  63.33333
46      1      1  96.66600  95.77778  61.66667
47      1      1  98.66600  100.00000 90.00000
48      1      1  0.00000   0.00000  93.33333
49      1      1  99.33400  100.00000 66.66667
50      1      0  97.33400  98.57143  60.00000
51      1      1  95.33400  100.00000 76.66667
52      1      0  94.66733  100.00000 70.00000
53      1      0  93.33200  98.57143  63.33333
54      1      0  87.33200  95.77778  80.00000
55      1      1  93.33400  96.00000  66.66667
56      1      0  70.00000  81.77778  46.66667
attr("assign")
[1] 0 1 2 3 4
attr("contrasts")
attr("contrasts")$year
[1] "contr.treatment"
```

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1. Messerli, F. H. (2012). Chocolate consumption, cognitive function, and Nobel laureates. *New England Journal of Medicine*, 367(16), 1562-1564. ↪