

Statistical Computing and Data Visualization in R

Lecture 9

Linear Models

Linear models: an example

- The data frame `whiteside` contains weekly gas consumption and average external temperature at a house in south-east England during two ‘heating seasons’, one before and one after cavity-wall insulation was installed.
- The goal of the exercise was to assess the effect of the insulation on gas consumption.

Linear models: an example

- The dataframe is in the MASS package

```
> library(MASS)
```

```
> head(whiteside)
```

	Insul	Temp	Gas
1	Before	-0.8	7.2
2	Before	-0.7	6.9
3	Before	0.4	6.4
4	Before	2.5	6.0
5	Before	2.9	5.8
6	Before	3.2	5.8

- The response (y) is **Gas**, and you have two predictors (x), one that is continuous (**Temp**) and one that is categorical (**Insul**)

Linear models: exploratory data analysis

- Note that the variable Gas is continuous (numeric), but the variable Insul is categorical (factor):

```
> is.numeric(Gas)
```

```
[1] TRUE
```

```
> is.factor(Insul)
```

```
[1] TRUE
```

- Let's plot the data:

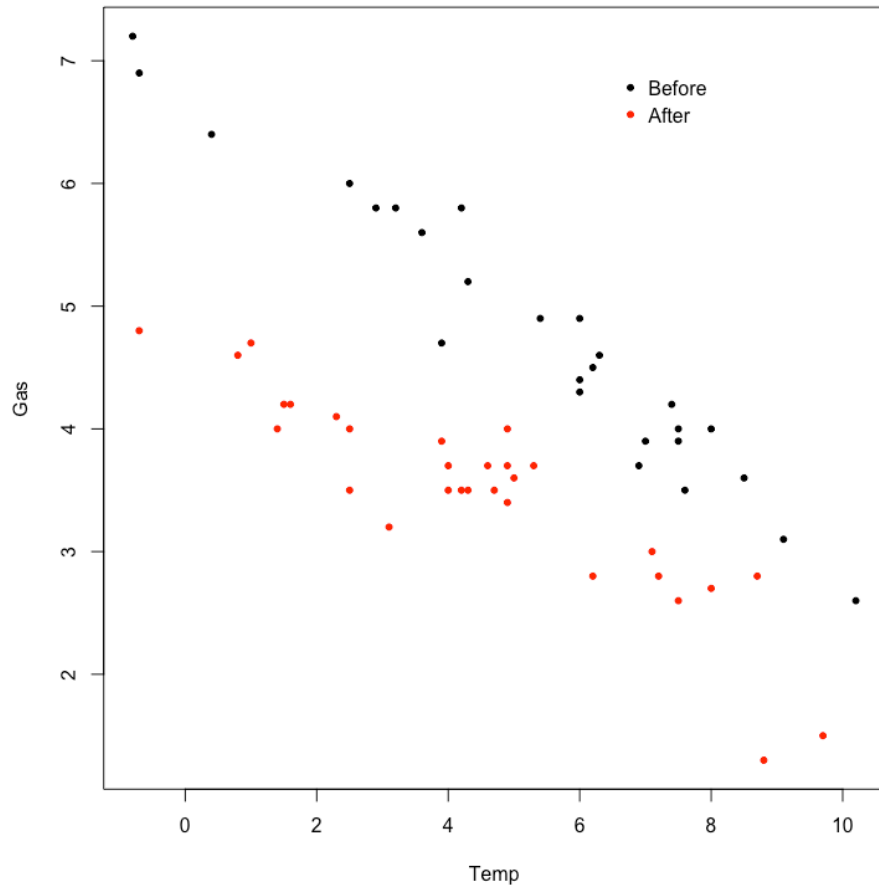
```
> attach(whiteside)
```

```
> par(mar=c(4,4,1,1)+0.1)
```

```
> plot(Temp, Gas, pch=20,  
col=as.numeric(Insul))
```

```
> legend(6.5, 7, c("Before", "After"),  
col=c("black", "red"), pch=20, bty="n")
```

Linear models: exploratory data analysis



- There is evidence of a linear relationship between Temperature and Gas consumption.
- However, the form of the relationship (both intercept and slope) depends on the type of insulation: after seems much better at low temperatures, but the differences tend to be minor at moderate ones.

Formulating a linear model

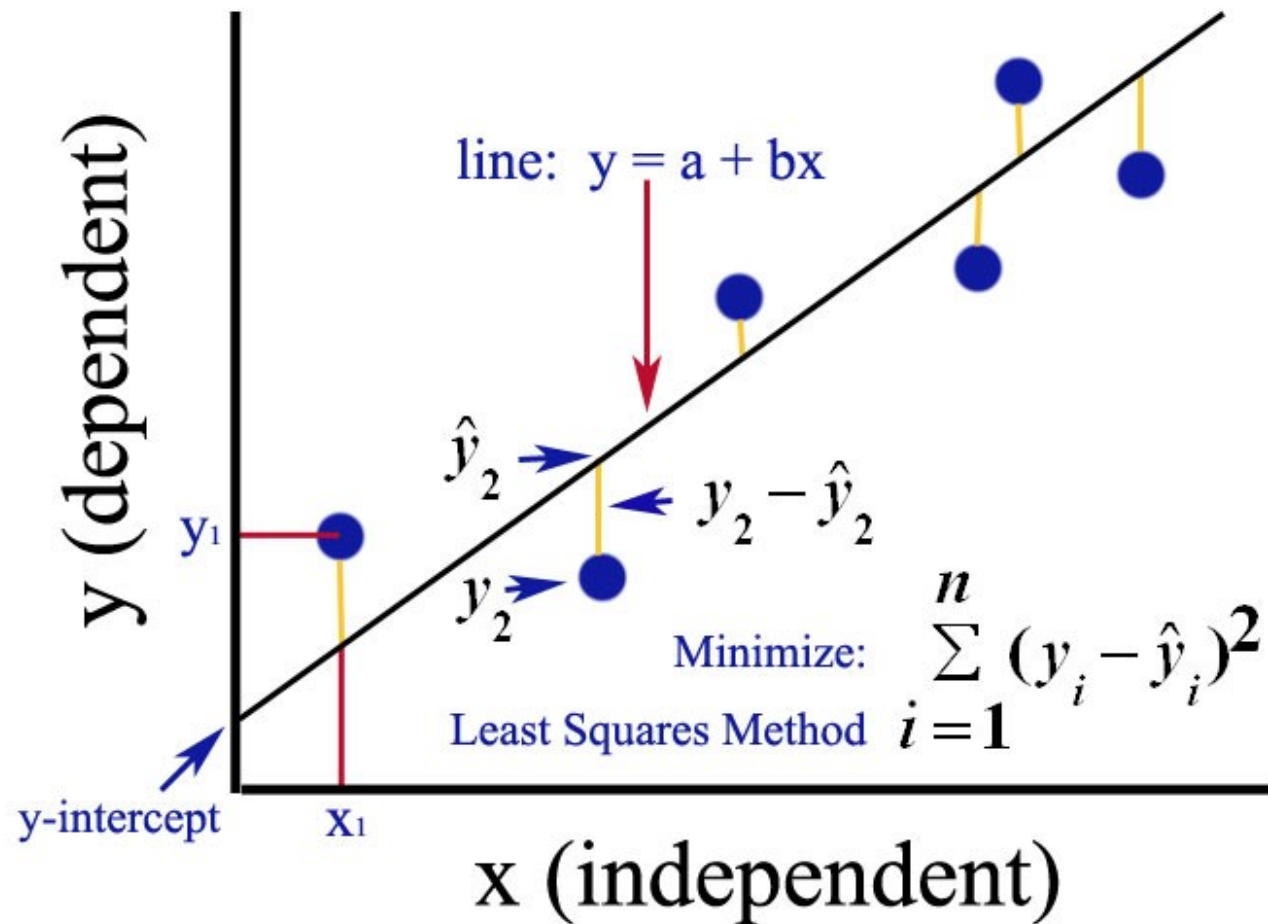
- The scatterplot suggests that fitting two separate simple linear regressions to each of the two groups of observations.

$$y_j = \alpha_0 + \beta_0 x_j + \varepsilon_j \quad \varepsilon_j \sim N(0, \sigma_0^2) \quad \text{If insulation = "Before"}$$

$$y_i = \alpha_1 + \beta_1 x_j + \omega_i \quad \omega_j \sim N(0, \sigma_1^2) \quad \text{If insulation = "After"}$$

- Recall that MLEs $(\hat{\alpha}_k, \hat{\beta}_k)$ for (α_k, β_k) under this model correspond to the least square estimators, so that $\hat{y}_j = \hat{\alpha}_k + \hat{\beta}_k x_j$.

Ordinary least squares



A joint model

- In the context of this formulation, the questions of interest translate into: Is $\alpha_0 = \alpha_1$ or $\beta_0 = \beta_1$?
- However, to be able to test these hypotheses we need to setup a joint model for both groups.

A joint model

- Let $y_{i,j}$ represent the j -th observation in group i , with $i = 0$ for “Before” and $i = 1$ for “After”. Define $x_{i,j}$ similarly.
- In this case we have 26 observations in group 0 (Before) and 30 in group 1 (After):

```
> table(Insul)
```

```
Insul
```

```
Before  After
```

```
      26      30
```

- Hence, i runs from 0 to 1 and j runs from 1 to 26 when $i = 0$ and from 1 to 30 when $i = 1$.

A joint model

- Under this notation, the joint model can be written

$$y_{i,j} = \alpha_i + \beta_i x_{i,j} + \varepsilon_{i,j} \qquad \varepsilon_{i,j} \sim N(0, \sigma^2)$$

- This model also allows for two distinct intercepts and slopes. However, we have now a single variance for the errors
- The model can also be written as

$$y_{i,j} = \mu + \alpha_i^* + (\eta + \beta_i^*)x_{i,j} + \varepsilon_{i,j} \qquad \varepsilon_{i,j} \sim N(0, \sigma^2)$$

where $\alpha_i = \mu + \alpha_i^*$, $\beta_i = \eta + \beta_i^*$, $\alpha_0 = 0$ and $\beta_0 = 0$. (α_1^* and β_1^* interpreted as differences with respect to the baseline category values given by μ and η).

Fitting linear models in R

- The function `lm` allows you fit linear models in R. The structure of the function is:

```
lm(formula, data, weights, subset, na.action)
```

- Some of the main arguments are

<code>formula</code>	is the model formula (the only required argument).
<code>data</code>	in an optional data frame.
<code>weights</code>	is a vector of positive weights, if non-uniform weights are needed
<code>subset</code>	is an index vector specifying a subset of the data to be used (by default all items are used)
<code>na.action</code>	is a function specifying how missing values are to be handled

Fitting simple linear models in R

- For example, to fit the individual models (simple linear regression).

```
> mod.before <- lm(Gas~Temp, data=whiteside,  
subset=(Insul=="Before"))
```

```
> mod.before
```

```
Call:lm(formula = Gas ~ Temp, data = whiteside, subset = (Insul  
== "Before"))
```

```
Coefficients:
```

```
(Intercept)      Temp  
    6.8538      -0.3932
```

```
> mod.after <- lm(Gas~Temp, data=whiteside,  
subset=(Insul=="After"))
```

```
> mod.after
```

```
Call:lm(formula = Gas ~ Temp, data = whiteside, subset = (Insul  
== "After"))
```

```
Coefficients:
```

```
(Intercept)      Temp  
    4.7238      -0.2779
```

Fitting simple linear models in R

- Simply printing the object will only give you back the call you used and the coefficients. More detail can be obtained using the function `summary()`

```
> summary(mod.before)
```

```
Call:lm(formula = Gas ~ Temp, data = whiteside, subset = (Insul == "Before"))
```

```
Residuals:      Min        1Q      Median        3Q       Max
-0.62020 -0.19947  0.06068  0.16770  0.59778
```

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	6.85383	0.11842	57.88	<2e-16 ***	H0: $\alpha = 0$ vs Ha: $\alpha \neq 0$
Temp	-0.39324	0.01959	-20.08	<2e-16 ***	H0: $\beta = 0$ vs Ha: $\beta \neq 0$

```
---
```

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

```
Residual standard error: 0.2813 on 24 degrees of freedom
```

```
Multiple R-squared: 0.9438, Adjusted R-squared: 0.9415
```

```
F-statistic: 403.1 on 1 and 24 DF, p-value: < 2.2e-16
```

lm objects

- An `lm` object is a list that contains a number of pieces. Some of the key ones

<code>coefficients</code>	a named vector of coefficients
<code>residuals</code>	residuals (response minus fitted values)
<code>fitted.values</code>	fitted mean values
<code>df.residual</code>	degrees of freedom of the residuals

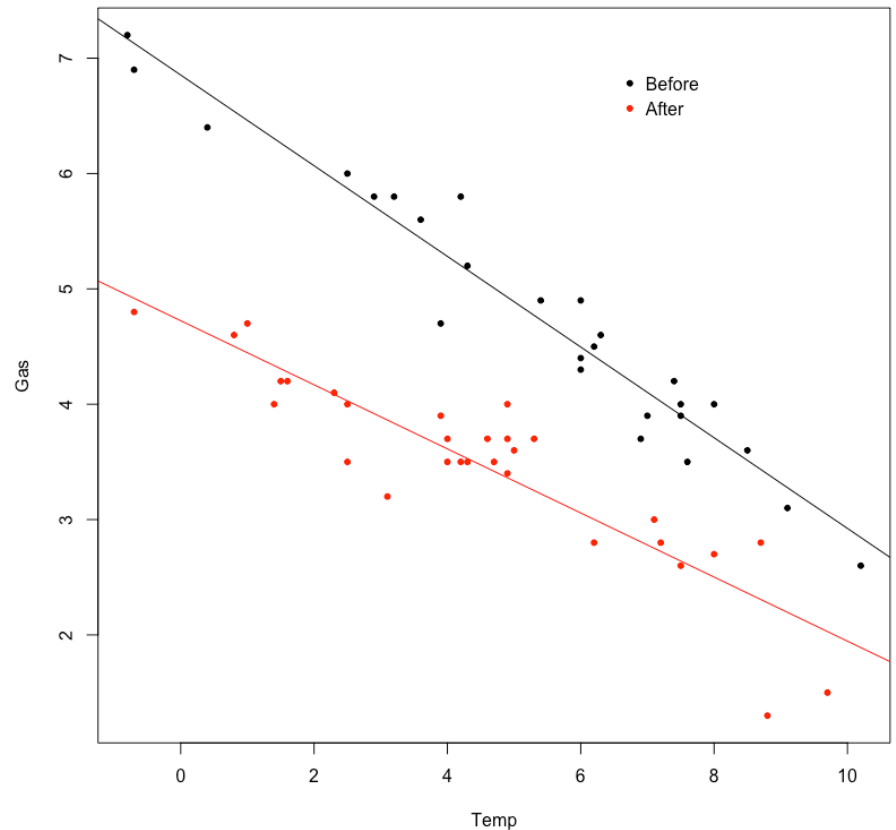
- For example, the estimate of the variance can be computed as

```
> sqrt(sum(mod.before$residuals^2) /  
df.residual)  
[1] 0.2813337
```

Plotting regression lines

- You can also use the elements of the object to add lines to the graph:

```
> par(mar=c(4,4,1,1)+0.1)
> plot(Temp, Gas, pch=20,
col=as.numeric(Insul))
> legend(6.5, 7, c("Before",
"After"),
col=c("black","red"), pch=20,
bty="n")
>
> abline(mod.before$coefficients,
col="black")
>
> abline(mod.after$coefficients,
col="red")
```



Fitting general linear models in R

- Let's do the joint model now

```
> mod.joint <- lm(Gas~Temp*Insul, data=whiteside)
> summary(mod.joint)
```

Call:lm(formula = Gas ~ Temp*Insul, data = whiteside)

Residuals:

	Min	1Q	Median	3Q	Max
	-0.97802	-0.18011	0.03757	0.20930	0.63803

Coefficients:

		Estimate	Std. Error	t value	Pr(> t)	
μ	(Intercept)	6.85383	0.13596	50.409	< 2e-16	***
α_1^*	Temp	-0.39324	0.02249	-17.487	< 2e-16	***
η	InsulAfter	-2.12998	0.18009	-11.827	2.32e-16	***
β_1^*	Temp:InsulAfter	0.11530	0.03211	3.591	0.000731	***

Signif. codes:

0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.323 on 52 degrees of freedom

Multiple R-squared: 0.9277, Adjusted R-squared: 0.9235

F-statistic: 222.3 on 3 and 52 DF, p-value: < 2.2e-16

Hypotheses testing for linear models

- Now we can answer the question we started with: what is the effect of insulation on gas consumption?
- This is really two questions:
 - Is the impact of insulation on gas consumption the same no matter what the external temperature is? (i.e., is the slope the same?, i.e., is $\beta_1^* \neq 0$?)
 - If the impact of insulation is the same at all temperatures, is that constant impact different non-negligible? (i.e., is the intercept the same?, i.e., is $\alpha_1^* \neq 0$?)
- Note that we ask these questions sequentially: if the answer to the first is yes the answer to the second is automatically yes too!

Hypotheses testing for linear models

- To answer the first question we can look at the p-value associated with β_1^*

```
> summary(mod.joint)
```

```
Call:lm(formula = Gas ~ Temp*Insul, data = whiteside)
```

```
Residuals:
```

Min	1Q	Median	3Q	Max
-0.97802	-0.18011	0.03757	0.20930	0.63803

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	6.85383	0.13596	50.409	< 2e-16	***
Temp	-0.39324	0.02249	-17.487	< 2e-16	***
InsulAfter	-2.12998	0.18009	-11.827	2.32e-16	***
Temp:InsulAfter	0.11530	0.03211	3.591	0.000731	***

```
---
```

```
Signif. codes:
```

```
0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

$H_0: \beta_1^* = 0$ vs. $H_a: \beta_1^* \neq 0$

Since the p-value is less than 0.05, we reject H_0 → The impact of insulation on gas consumption varies with the temperature. No further test is needed.

Hypotheses testing for linear models

- An alternative way to contrast these hypotheses is to run an F test between the model with different slopes and a model with a common slope:

```
> mod.joint2 <- lm(Gas~Temp+Insul, data=whiteside)
> aov(mod.joint2, mod.joint)
Analysis of Variance Table
Model 1: Gas ~ Temp + Insul
Model 2: Gas ~ Temp * Insul
```

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	53	6.7704				
2	52	5.4252	1	1.3451	12.893	0.0007307 ***

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

- Note in that this is the same p-value as before (in this case the tests are equivalent).
- More generally, F tests allow you to test multiple coefficients simultaneously (unlike t tests).

The general linear model

- The function `lm` allows you to fit any general linear model

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon} \quad \boldsymbol{\varepsilon} \sim N_n(0, \sigma^2 \text{diag}\{w_1, \dots, w_n\})$$

Where \mathbf{y} is a vector of observations, \mathbf{X} is the known design matrix, $\boldsymbol{\beta}$ is a vector of unknown coefficients, σ^2 is the variance and w_1, \dots, w_n are known weights.

- Examples of the general linear model include:
 - Simple and multiple regression.
 - Analysis of Variance (ANOVA) models.
 - Analysis of Covariance (ANCOVA) models.

Using formulas with lm

Symbol	Example	Meaning
+	+X	Include this variable
-	-X	Delete this variable
:	X : Z	Include the interaction between the variables
*	X * Y	Include these variables as well as their interactions
	X Z	Conditioning/nesting: include x given z
^	(X+Z+W) ^ 3	Include all these variables and all interactions up to 3-way
I	I (X * Z)	Isolate/as is: include a new variable obtained by performing the isolated operation
1	X-1	Intercept: do not include an intercept (default is to include)

Note that all of the following are equivalent:

$$Y \sim X + Z + W + X:Z + Z:W + Z:W$$

$$Y \sim X * Z * W - X:Z:W$$

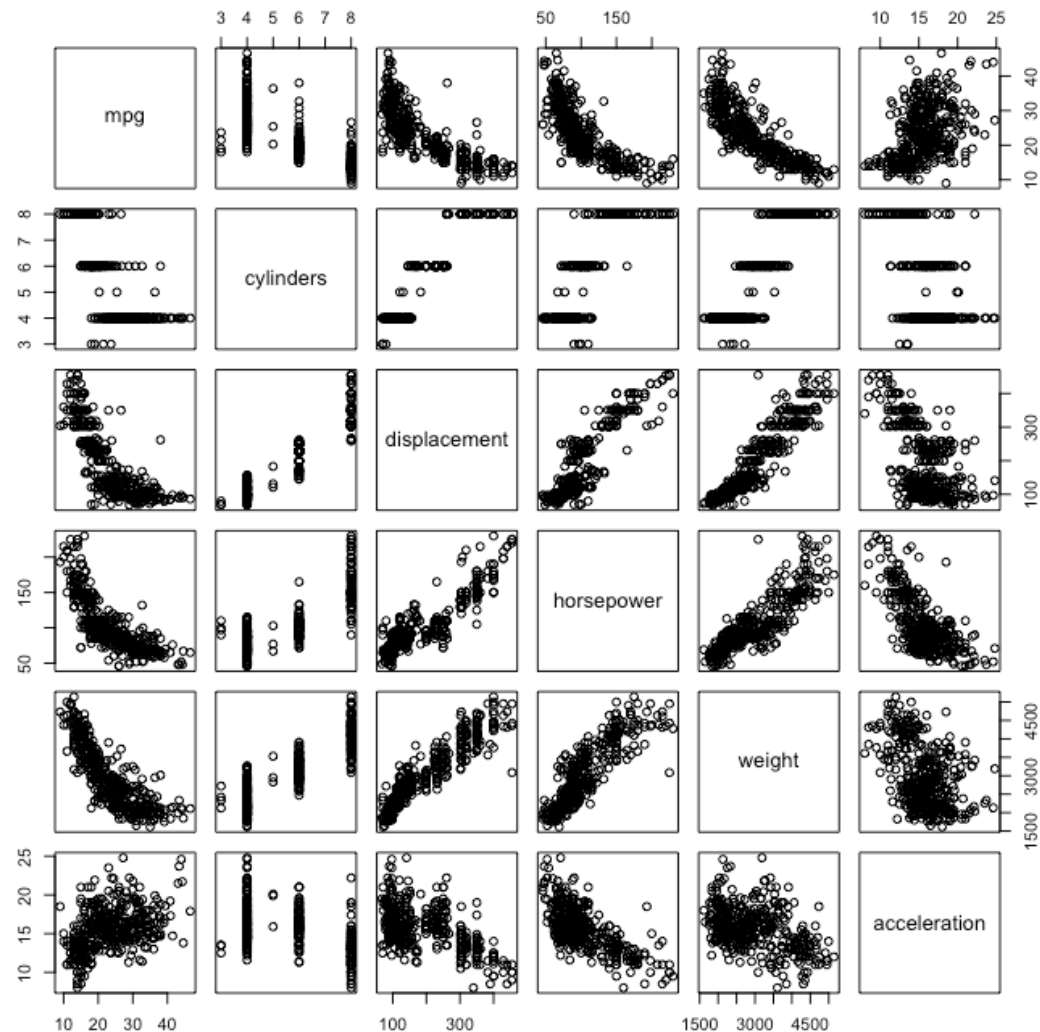
$$Y \sim (X + Z + W) ^ 2$$

A multiple regression example: Car fuel consumption

- Consider now a second example where we are trying to understand how different factors affect city-cycle fuel consumption in miles per gallon for various car models.
- The dataset was used as the testbed for graphical analysis packages at the 1983 American Statistical Association Exposition
- We have one continuous response variable as well as 5 continuous and 3 categorical predictors. We are going to ignore the 3 categorical ones (model year, origin and car name).

Descriptive analysis

```
> dat = read.table(file="auto-mpg.txt", header=T)
> pairs(~ mpg + cylinders + displacement+ horsepower + weight + acceleration, data=dat)
```



Car fuel consumption

- In this case we are going to fit a model without any interactions:

```
> dat = read.table(file="auto-mpg.txt", header=T)
> names(dat)
> mod = lm(mpg ~ cylinders + displacement + horsepower + weight + acceleration,
data=dat)
> summary(mod)
```

$$y_i = \beta_0 + \beta_1 x_{1,i} + \beta_2 x_{2,i} + \beta_3 x_{3,i} + \beta_4 x_{4,i} + \beta_5 x_{5,i} + \varepsilon_i$$

Call:

```
lm(formula = mpg ~ cylinders + displacement + horsepower + weight +
    acceleration, data = dat)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-11.5816	-2.8618	-0.3404	2.2438	16.3416

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
β_0 (Intercept)	4.626e+01	2.669e+00	17.331	<2e-16	***
β_1 cylinders	-3.979e-01	4.105e-01	-0.969	0.3330	
β_2 displacement	-8.313e-05	9.072e-03	-0.009	0.9927	
β_3 horsepower	-4.526e-02	1.666e-02	-2.716	0.0069	**
β_4 weight	-5.187e-03	8.167e-04	-6.351	6e-10	***
β_5 acceleration	-2.910e-02	1.258e-01	-0.231	0.8171	

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

Residual standard error: 4.247 on 386 degrees of freedom
(6 observations deleted due to missingness)

Multiple R-squared: 0.7077, Adjusted R-squared: 0.7039

F-statistic: 186.9 on 5 and 386 DF, p-value: < 2.2e-16

Backward selection

- We can start eliminating variables and refitting the model. We first drop displacement:

```
> mod.a = lm(mpg ~ cylinders + horsepower + weight + acceleration,  
data=dat)  
> summary(mod.a)
```

```
Call:lm(formula = mpg ~ cylinders + horsepower + weight + acceleration,  
data = dat)
```

Residuals:

Min	1Q	Median	3Q	Max
-11.5807	-2.8628	-0.3409	2.2427	16.3422

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	46.2739915	2.4481591	18.902	< 2e-16	***
cylinders	-0.4004602	0.3032615	-1.321	0.18744	
horsepower	-0.0452970	0.0160604	-2.820	0.00504	**
weight	-0.0051902	0.0007341	-7.070	7.26e-12	***
acceleration	-0.0289828	0.1248944	-0.232	0.81661	

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

Backward selection

- Next we drop acceleration:

```
> mod.a = lm(mpg ~ cylinders + horsepower + weight, data=dat)
> summary(mod.a)
```

```
Call:lm(formula = mpg ~ cylinders + horsepower + weight, data =
dat)
```

Residuals:

Min	1Q	Median	3Q	Max
-11.5260	-2.7955	-0.3559	2.2567	16.3209

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	45.7368172	0.7959566	57.461	< 2e-16 ***
cylinders	-0.3889745	0.2988302	-1.302	0.193806
horsepower	-0.0427277	0.0116196	-3.677	0.000269 ***
weight	-0.0052723	0.0006424	-8.208	3.37e-15 ***

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

Backward selection

- Finally, we drop cylinders:

```
> mod.a = lm(mpg ~ cylinders + horsepower + weight, data=dat)
> summary(mod.a)
```

```
Call:lm(formula = mpg ~ horsepower + weight, data = dat)
```

Residuals:

Min	1Q	Median	3Q	Max
-11.0762	-2.7340	-0.3312	2.1752	16.2601

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	45.6402108	0.7931958	57.540	< 2e-16 ***
horsepower	-0.0473029	0.0110851	-4.267	2.49e-05 ***
weight	-0.0057942	0.0005023	-11.535	< 2e-16 ***

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

The variables that seem to explain mpg are horsepower and weight.

Continuous vs Categorical

- Note that we are treating cylinders a continuous variable, so `lm` fits a linear regression:

```
> mod1 = lm(mpg ~ cylinders, data=dat)
> summary(mod1)
```

Call:

```
lm(formula = mpg ~ cylinders, data = dat)
```

Residuals:

Min	1Q	Median	3Q	Max
-14.2607	-3.3841	-0.6478	2.5538	17.9022

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	42.9493	0.8330	51.56	<2e-16 ***
cylinders	-3.5629	0.1458	-24.43	<2e-16 ***

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

Residual standard error: 4.942 on 396 degrees of freedom

Multiple R-squared: 0.6012, Adjusted R-squared: 0.6002

F-statistic: 597.1 on 1 and 396 DF, p-value: < 2.2e-16

Continuous vs Categorical

- If we turn the number of cylinder into a categorical variable, `lm` fits an ANOVA model instead:

```
> mod2 = lm(mpg ~ factor(cylinders), data=dat)
> summary(mod2)
```

Call:

```
lm(formula = mpg ~ factor(cylinders), data = dat)
```

Residuals:

Min	1Q	Median	3Q	Max
-11.2868	-2.9631	-0.9631	2.3890	18.0143

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	20.5500	2.3657	8.687	< 2e-16	***
factor(cylinders)4	8.7368	2.3888	3.657	0.000289	***
factor(cylinders)5	6.8167	3.6137	1.886	0.059985	.
factor(cylinders)6	-0.5643	2.4214	-0.233	0.815849	
factor(cylinders)8	-5.5869	2.4112	-2.317	0.021014	*

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

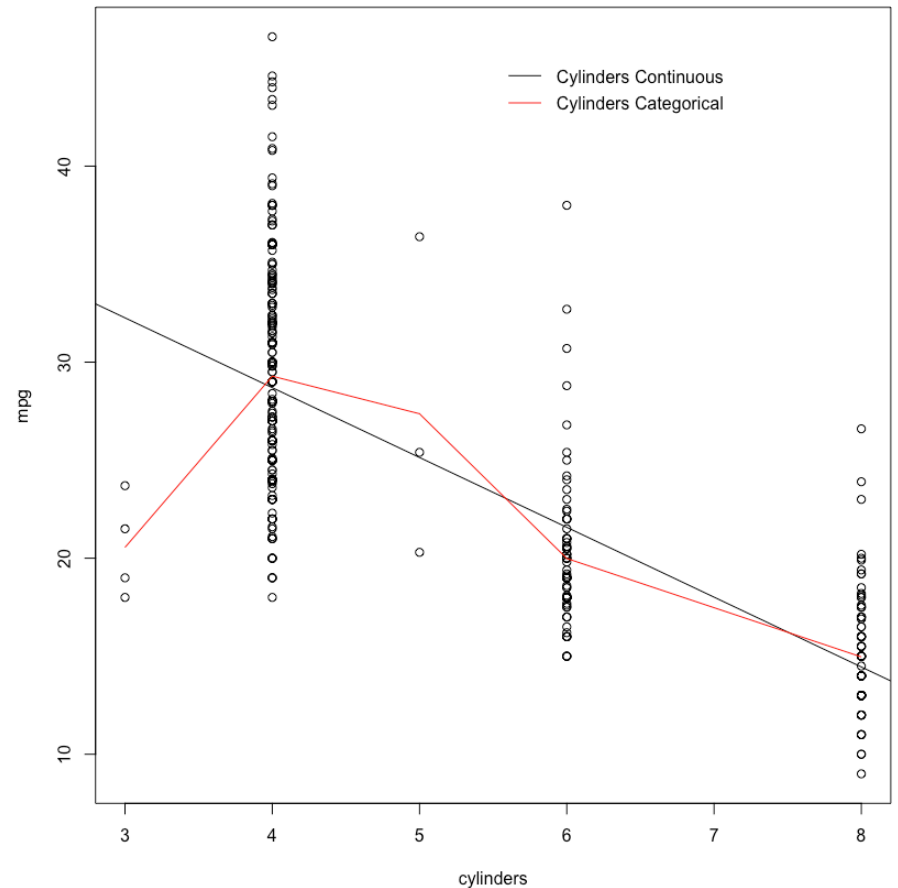
Residual standard error: 4.731 on 393 degrees of freedom

Multiple R-squared: 0.6372, Adjusted R-squared: 0.6335

F-statistic: 172.6 on 4 and 393 DF, p-value: < 2.2e-16

Continuous vs Categorical

- What is more appropriate in the previous example, to treat the number of cylinders as a continuous or a categorical variable?



Continuous vs Categorical

- When you have different classes of predictors, `lm` fits an ANCOVA model (as in our first example):

```
> mod3 = lm(mpg ~ weight*factor(cylinders), data=dat)
> summary(mod3)
```

Call:

```
lm(formula = mpg ~ weight * factor(cylinders), data = dat)
```

Residuals:

Min	1Q	Median	3Q	Max
-10.2700	-2.4097	-0.4621	1.8307	17.0414

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	5.166320	22.673910	0.228	0.820
weight	0.006414	0.009416	0.681	0.496
factor(cylinders)4	44.742048	22.753837	1.966	0.050 *
factor(cylinders)5	25.440982	32.858673	0.774	0.439
factor(cylinders)6	31.801776	23.075818	1.378	0.169
factor(cylinders)8	24.277498	22.971639	1.057	0.291
weight:factor(cylinders)4	-0.015348	0.009451	-1.624	0.105
weight:factor(cylinders)5	-0.007458	0.012117	-0.616	0.539
weight:factor(cylinders)6	-0.011724	0.009510	-1.233	0.218
weight:factor(cylinders)8	-0.009933	0.009458	-1.050	0.294

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

Residual standard error: 4.037 on 388 degrees of freedom

Multiple R-squared: 0.7392, Adjusted R-squared: 0.7332

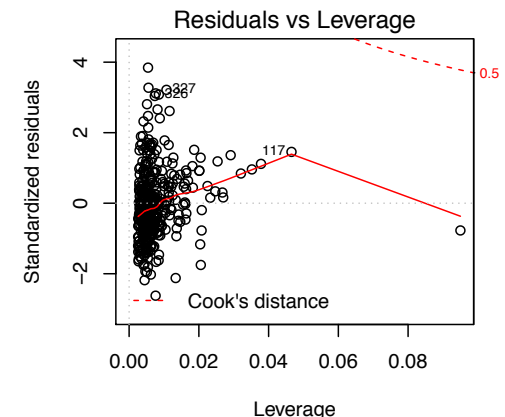
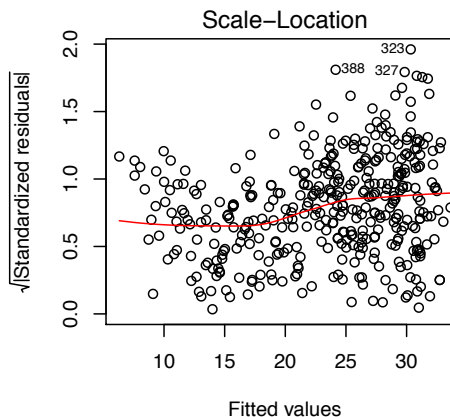
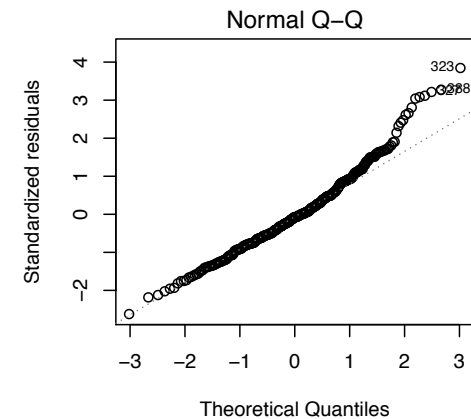
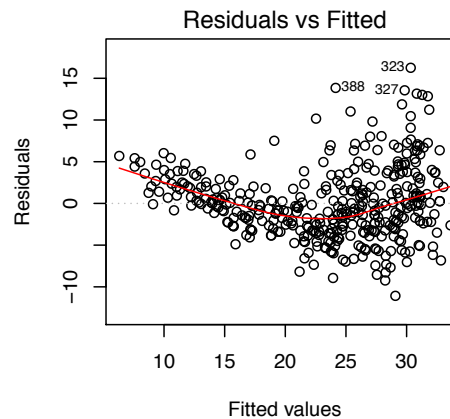
F-statistic: 122.2 on 9 and 388 DF, p-value: < 2.2e-16

Goodness of fit in linear models

- You can plot a model object to facilitate the investigation of model fit:

```
> quartz()  
> par(mfrow=c(2,2))  
> plot(mod)
```

- What should we do about the residual trend?
- How to deal with the heteroskedasticity?
- What should we do about the right tail of the residuals?



Dealing with lack of fit

- It is clear from the descriptive and residual plots that we might want to add quadratic terms to the regression:

```
> mod6 = lm(mpg ~ cylinders + horsepower + I(horsepower^2) + weight +  
I(weight^2) + displacement + I(displacement^2) + acceleration, data=dat)  
> summary(mod6)
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	6.256e+01	4.455e+00	14.042	< 2e-16	***
cylinders	7.847e-01	4.236e-01	1.852	0.064750	.
horsepower	-2.969e-01	5.328e-02	-5.571	4.78e-08	***
I(horsepower^2)	7.249e-04	1.819e-04	3.984	8.11e-05	***
weight	-2.050e-03	3.381e-03	-0.606	0.544653	
I(weight^2)	9.793e-08	4.584e-07	0.214	0.830935	
displacement	-8.150e-02	2.428e-02	-3.356	0.000869	***
I(displacement^2)	1.157e-04	4.353e-05	2.658	0.008193	**
acceleration	-3.866e-01	1.326e-01	-2.916	0.003758	**

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

Residual standard error: 4.731 on 393 degrees of freedom
Multiple R-squared: 0.6372, Adjusted R-squared: 0.6335
F-statistic: 172.6 on 4 and 393 DF, p-value: < 2.2e-16

Dealing with lack of fit

- After doing some stepwise regression

```
> mod7 = lm(mpg ~ cylinders + horsepower + I(horsepower^2) +  
displacement + I(displacement^2) + acceleration, data=dat)  
> summary(mod7)
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	63.9777080	3.1315133	20.430	< 2e-16	***
cylinders	0.8473024	0.3978732	2.130	0.033840	*
horsepower	-0.3396011	0.0443881	-7.651	1.62e-13	***
I(horsepower^2)	0.0008323	0.0001660	5.015	8.11e-07	***
displacement	-0.0933867	0.0178144	-5.242	2.62e-07	***
I(displacement^2)	0.0001239	0.0000344	3.601	0.000359	***
acceleration	-0.5072507	0.1074119	-4.722	3.27e-06	***

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

Residual standard error: 4.731 on 393 degrees of freedom

Multiple R-squared: 0.6372, Adjusted R-squared: 0.6335

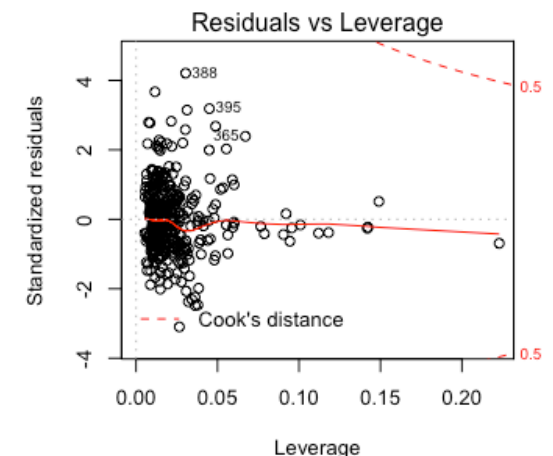
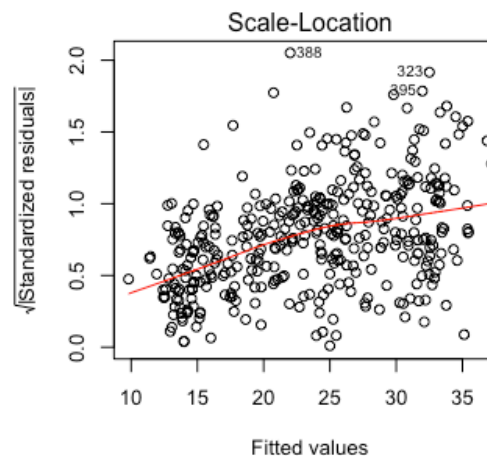
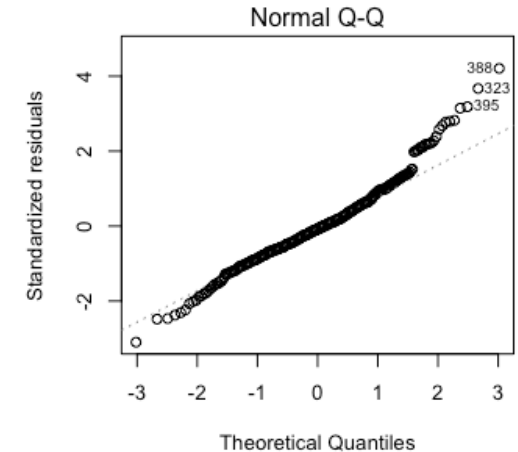
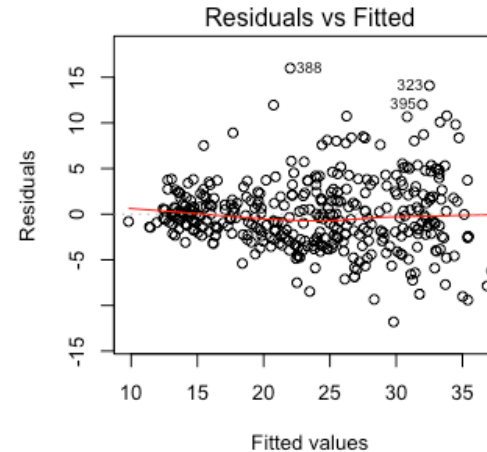
F-statistic: 172.6 on 4 and 393 DF, p-value: < 2.2e-16

Now weight is not significant, and cylinders, displacement and acceleration are.

Residual analysis

Residuals are not perfect (normal Q-Q plot still shows a heavy right tail and you see some heteroscedasticity), but at least the mean looks better.

You might need to log transform the response to address the remaining issue.



Predictions

- Predictions at new values are easy to obtain:

```
> mod5 = lm(mpg ~ horsepower + weight, data=dat)
> xn = data.frame(horsepower = c(60,90,120), weight = c(2500,
3250, 4000))
> gpm.pred = predict(mod5, xn, se.fit=TRUE)
> gpm.pred

$fit
      1      2      3
28.31665 22.55194 16.78724

$se.fit
      1      2      3
0.3767431 0.3581450 0.4317328

$df
[1] 389

$residual.scale
[1] 4.240169
```

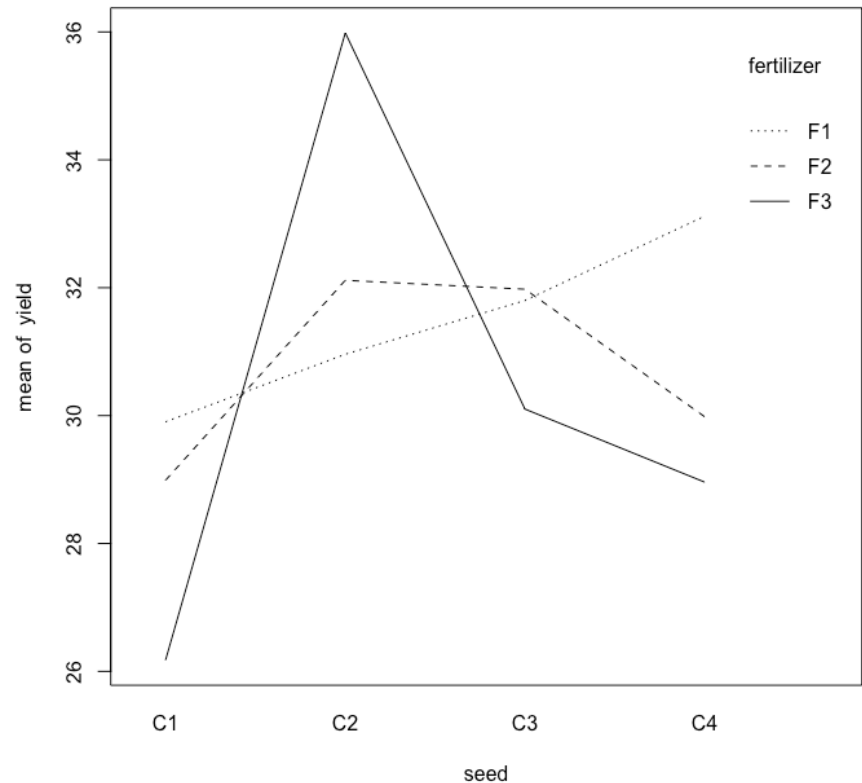
ANOVA models

- We want to investigate now how the type of fertilizer and seed used to grow maize affects the yield.
- In this case the response is continuous, but we have two categorical predictors, one with 3 levels (fertilizer) and one with 4 levels (seed).
- Four replicates were taken for each combination of factors.
- Questions of interest:
 - Does the effect of fertilizer depend on the type of seed?
 - If that is not the case, is there an effect from either fertilizer or seed on their own.

Descriptive analysis

- The first thing to do is to construct interaction plots:

```
> yields = read.table(  
  "yields.csv", header=T)  
> attach(yields)  
> interaction.plot(seed,  
  fertilizer, yield)
```
- The graph suggests the presence of an interaction between fertilizer and seed (i.e., the type of seed affects how well the fertilizer works).



Testing for the presence of an interaction

- We can use an F test to determine whether there is actually a significant interaction between the variables:

```
> mod = lm(yield ~ fertilizer*seed, data=yields)
> anova(mod)
```

Analysis of Variance Table

Response: yield

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
fertilizer	2	10.528	5.264	90.986	8.361e-15	***
seed	3	133.843	44.614	771.155	< 2.2e-16	***
fertilizer:seed	6	121.206	20.201	349.171	< 2.2e-16	***
Residuals	36	2.083	0.058			

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

- **The interaction (`fertilizer:seed`) appears to be highly significant in this dataset. Hence, we do no further testing.**
- If the interaction had not been significant, then we refit the model without it and would test for the significance of the main effects (`fertilizer` and `seed`).

Pairwise multiple comparisons

- Since there is an interaction, we might want to know what fertilizer is most effective for each type of seed.
- We can get the mean yields for each combination of factor using `aggregate()`.

```
> aggregate(yield, by=list(seed,fertilizer), mean)
```

	Group.1	Group.2	x
1	C1	F1	29.90229
2	C2	F1	30.95948
3	C3	F1	31.79963
4	C4	F1	33.11874
5	C1	F2	28.98773
6	C2	F2	32.11534
7	C3	F2	31.97647
8	C4	F2	29.98227
9	C1	F3	26.17516
10	C2	F3	35.98370
11	C3	F3	30.10192
12	C4	F3	28.95868

Pairwise multiple comparisons

- Consider for example seed C1.
- In that case the highest yield seems to be associated with fertilizer F3. But are the differences statistically significant?
- To answer that question we need to run pairwise tests

```
> pairwise.t.test(yield[seed=="C1"], fertilizer[seed=="C1"],  
p.adjust.method="bonferroni")
```

Pairwise comparisons using t tests with pooled SD

data: yield[seed == "C1"] and fertilizer[seed == "C1"]

	F1	F2
F2	0.0041	-
F3	5.3e-08	6.2e-07

} All p-values are < 0.05, so all
differences in mean are
statistically significant!

Simplest possible correction
for multiple comparisons

P value adjustment method: bonferroni

Contrasts

- When categorical variables (factors) are included in a linear model, computation requires that they be encoded using dummy variables.
- There is an infinite number of potential encodings, each corresponding to one different set of constraints that ensure that the parameters of the model are estimable.

Contrasts

- Consider a simple example in which we are trying to explain writing skill as a function of race. The ANOVA model takes the form

$$y_{i,j} = \mu + \alpha_i + \varepsilon_{i,j} \quad j = 1, \dots, J_i \quad i = 1, \dots, I$$

- For example:

```
> hsb2 = read.table("hsb2.csv", sep=",", header=T)
> hsb2$race.f = factor(hsb2$race,
labels=c("Hispanic", "Asian", "African-Am",
"Caucasian"))
> tapply(hsb2$write, hsb2$race.f, mean)
Hispanic      Asian African-Am  Caucasian
46.45833    58.00000    48.20000    54.05517
```

- Here we have $I = 4$ groups and 5 parameters (μ plus 4 α s), so one constraint needs to be introduced!

Contrasts

- One popular options is for the first category to be the baseline level and the regression coefficients to represent differences with respect to the baseline level, i.e., $\alpha_1 = 0$.

```
> contrasts(hsb2$race.f) = contr.treatment(nlevels(hsb2$race.f))
> mod = lm(write ~ race.f, hsb2)
> summary(mod)
```

Call:

```
lm(formula = write ~ race.f, data = hsb2)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-23.0552	-5.4583	0.9724	7.0000	18.8000

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	46.458	1.842	25.218	< 2e-16	***
race.f2	11.542	3.286	3.512	0.000552	***
race.f3	1.742	2.732	0.637	0.524613	
race.f4	7.597	1.989	3.820	0.000179	***

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

Residual standard error: 9.025 on 196 degrees of freedom

Multiple R-squared: 0.1071, Adjusted R-squared: 0.0934

F-statistic: 7.833 on 3 and 196 DF, p-value: 5.785e-05


Contrasts

- Note that the estimate of the intercept is the mean for Hispanics (see previous slide), the one for `race.f2` is the difference between the mean of Hispanics and the mean for Asians, $58.000 - 46.458 = 11.542$, and so on ...
- It is illustrative to look at the matrix of contrasts:

```
> contr.treatment(nlevels(hsb2$race.f))
```

	2	3	4
1	0	0	0
2	1	0	0
3	0	1	0
4	0	0	1

Mean of the second factor is the
intercept plus the coefficient of
the first dummy variable



- This type of contrast is the default in R.

Contrasts

- You can change the baseline level for the contrast (now African-Americans are the baseline group):

```
> contrasts(hsb2$race.f) = contr.treatment(4)[c(3,2,1,4),]  
> mod = lm(write ~ race.f, hsb2)  
> summary(mod)
```

Call:

```
lm(formula = write ~ race.f, data = hsb2)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-23.0552	-5.4583	0.9724	7.0000	18.8000

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	48.200	2.018	23.884	< 2e-16	***
race.f2	9.800	3.388	2.893	0.00425	**
race.f3	-1.742	2.732	-0.637	0.52461	
race.f4	5.855	2.153	2.720	0.00712	**

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

Residual standard error: 9.025 on 196 degrees of freedom

Multiple R-squared: 0.1071, Adjusted R-squared: 0.0934

F-statistic: 7.833 on 3 and 196 DF, p-value: 5.785e-05

Contrasts

- Alternatively, we can interpret the intercept as the grand mean (mean of group means):

```
> contrasts(hsb2$race.f) = contr.sum(nlevels(hsb2$race.f))
> mod = lm(write ~ race.f, hsb2)
> summary(mod)
```

Call:

```
lm(formula = write ~ race.f, data = hsb2)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-23.0552	-5.4583	0.9724	7.0000	18.8000

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	51.6784	0.9821	52.619	< 2e-16	***
race.f1	-5.2200	1.6314	-3.200	0.00160	**
race.f2	6.3216	2.1603	2.926	0.00384	**
race.f3	-3.4784	1.7323	-2.008	0.04602	*

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

Residual standard error: 9.025 on 196 degrees of freedom

Multiple R-squared: 0.1071, Adjusted R-squared: 0.0934

F-statistic: 7.833 on 3 and 196 DF, p-value: 5.785e-05

Contrasts

- Note that:
 - The intercept is indeed the grand mean $(46.45833 + 58.00000 + 48.20000 + 54.05517)/4 = 51.67837$
 - The coefficient for race.f1 is the difference between the mean of Hispanics and the grand mean and $46.45833 - 51.67837 = -5.2200$ (“Hispanic effect”).
 - The coefficient for race.f2 is the difference between the mean of Asians and the grand mean and $58 - 51.67837 = 6.32163$ (“Asianeffect”).
 - The coefficient for race.f3 is the difference between the mean of African-Americans and the grand mean and $48.2 - 51.67837 = -3.47837$ (“African-American effect”).
 - The “Caucassian” effect is simply $3 \times 51.67837 - 46.45833 - 58.00000 - 48.2 = 2.37678$.

Contrasts

- The matrix of contrasts in this case is

```
> contr.sum(nlevels(hsb2$race.f))
```

	[,1]	[,2]	[,3]
1	1	0	0
2	0	1	0
3	0	0	1
4	-1	-1	-1

- Other options include `contr.poly` (for equally-space ordinal variables, useful to assess trends) and `contr.helmert` (comparing each level to the mean of all subsequent ones).
- You can create your own encoding by providing an appropriate matrix of contrasts!