

R for Regression **Dolores Romero Morales**

The aim of this workshop is to perform Regression with R using a brief guide to R for Regression and three datasets (tupelo.txt, blood.txt, housing.txt). All files can be found in the course directory. The tupelo and the blood datasets were used during the lecture. More information on the housing dataset can be found in <https://archive.ics.uci.edu/ml/machine-learning-databases/housing/>.

Once you are ready with the guide, use the datasets to perform the following steps using R:

Task 1

Tupelo dataset: Linear Regression

Step 1. Read the tupelo.txt file into a data frame.

Answer:

```
mytupelo <- read.table(file.choose(),header=TRUE)
```

Note:

With `file.choose()` a new window will pop up and you will need to find the file.

Step 2. Regress the Cost variable against Capacity and Year.

Answer:

```
model1 <- lm(mytupelo$Cost ~ .,mytupelo)
```

Note:

You can use `attach()` to shorten the names of the variables of a data frame. For instance, by calling `attach(mytupelo)`, you can use `Capacity` instead of `mytupelo$Capacity`. Use `detach(mytupelo)` when you want to end the attachment. Be careful with `attach()` in case you are working with several data frames.

Step 3. Give a summary of the model. Check the fit of the model and the significance of the explanatory variables.

Answer:

```
summary(model1)
```

Note1: This is the output we get with `summary()`

Call:

```
lm(formula = mytupelo$Cost ~ ., data = mytupelo)
```

Residuals:

Min	1Q	Median	3Q	Max
-2.4966	-1.4692	-0.2149	1.2054	3.2877

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	20.5488	3.1186	6.589	0.000307 ***
Year	0.6187	0.3521	1.757	0.122312
Capacity	-3.2121	1.1368	-2.826	0.025567 *

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

Residual standard error: 2.079 on 7 degrees of freedom

Multiple R-squared: 0.7903, Adjusted R-squared: 0.7304

F-statistic: 13.19 on 2 and 7 DF, p-value: 0.004223

Note2:

Fit of a linear regression model is given by the Adjusted R Squared, while the significance of explanatory variables by their p-value. We can see that Year has a p-value of 0.122312. If we choose a significance of 0.05, this variable is not significant to the model.

Note3:

At this point, it would be good to detach the data frame, i.e.,
detach(mytupelo)

Task 2

Blood dataset: Logistic Regression

Step 1. Read the blood.txt file into a data frame.

Answer:

```
myblood <- read.table(file.choose(), header=TRUE)
```

Step 2. Regress the Blood variable against Smoke and Alcohol.

Answer:

```
model2 <- glm(Blood ~ ., family='binomial', myblood)
```

Step 3. Give a summary of the model, and check the fit of the model and the significance of the explanatory variables.

Answer:

```
summary(model2)  
logLik(model2)
```

Note1:

The output of the summary() function is

Call:

```
glm(formula = Blood ~ ., family = "binomial", data = myblood)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-2.5205	-0.6426	-0.5371	0.4434	1.9358

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-1.39440	0.37295	-3.739	0.000185 ***
Smoke	2.26416	0.51287	4.415	1.01e-05 ***
Alcohol	-0.07816	0.08500	-0.920	0.357792

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

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 137.186 on 99 degrees of freedom
Residual deviance: 77.999 on 97 degrees of freedom
AIC: 83.999

Number of Fisher Scoring iterations: 7

Note2:

The significance of explanatory variables is again measured by their p-value. We can see that Alcohol has a p-value of 0.357792. If we choose a significance of 0.05, this variable is not significant to the model.

Note3:

In logistic regression, when used for probability estimation, has not got a natural way to measure fit. Both, Cox-Snell and Nagelkerke R Square, are a try to simulate what R Square does in Linear Regression.

An alternative way to measure fit is the log likelihood. This is always a negative number. When comparing two models in terms of fit, we would like to choose the one with the smallest absolute value of the log likelihood.

Task 3

Housing dataset: Multiple Regression

Step 1. Read the housing.txt file into a data frame.

Answer:

```
myhousing <- read.table(file.choose(), header=TRUE)
```

Note:

Although not required, it is advisable to call the attach function and the dimension one.

Step 2. Regress the last variable against the rest.

Answer:

```
modelwith13 <- lm(myhousing$MEDV ~ ., myhousing)
```

Step 3. Give a summary of the model, and check the fit of the model and the significance of all explanatory variables.

Answer:

```
summary(modelwith13)
```

Note1: See the output below.

Call:

```
lm(formula = MEDV ~ ., data = myhousing)
```

Residuals:

Min	1Q	Median	3Q	Max
-15.595	-2.730	-0.518	1.777	26.199

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	3.646e+01	5.103e+00	7.144	3.28e-12 ***
CRIM	-1.080e-01	3.286e-02	-3.287	0.001087 **
ZN	4.642e-02	1.373e-02	3.382	0.000778 ***
INDUS	2.056e-02	6.150e-02	0.334	0.738288
CHAS	2.687e+00	8.616e-01	3.118	0.001925 **
NOX	-1.777e+01	3.820e+00	-4.651	4.25e-06 ***
RM	3.810e+00	4.179e-01	9.116	< 2e-16 ***
AGE	6.922e-04	1.321e-02	0.052	0.958229
DIS	-1.476e+00	1.995e-01	-7.398	6.01e-13 ***
RAD	3.060e-01	6.635e-02	4.613	5.07e-06 ***
TAX	-1.233e-02	3.760e-03	-3.280	0.001112 **
PTRATIO	-9.527e-01	1.308e-01	-7.283	1.31e-12 ***
B	9.312e-03	2.686e-03	3.467	0.000573 ***
LSTAT	-5.248e-01	5.072e-02	-10.347	< 2e-16 ***

Signif. codes:	0 ‘***’	0.001 ‘**’	0.01 ‘*’	0.05 ‘.’
	0.1 ‘ ’	1		

Residual standard error: 4.745 on 492 degrees of freedom
Multiple R-squared: 0.7406, Adjusted R-squared: 0.7338
F-statistic: 108.1 on 13 and 492 DF, p-value: < 2.2e-16

Note2: The question is what set of variables we should build the model on. The step() function performs model selection in a stepwise fashion based on the Akaike Information Criterion (AIC). This criterion is a tradeoff between the number of variables used and the fit of the model obtained with those variables. We can see below that we would end up with 11 out of the 13 explanatory variables.

step(modelwith13)

Start: AIC=1589.64
myhousing\$MEDV ~ CRIM + ZN + INDUS + CHAS + NOX + RM + AGE +
DIS + RAD + TAX + PTRATIO + B + LSTAT

	Df	Sum of Sq	RSS	AIC
- AGE	1	0.06	11079	1587.7
- INDUS	1	2.52	11081	1587.8
<none>			11079	1589.6
- CHAS	1	218.97	11298	1597.5
- TAX	1	242.26	11321	1598.6
- CRIM	1	243.22	11322	1598.6
- ZN	1	257.49	11336	1599.3
- B	1	270.63	11349	1599.8
- RAD	1	479.15	11558	1609.1
- NOX	1	487.16	11566	1609.4
- PTRATIO	1	1194.23	12273	1639.4
- DIS	1	1232.41	12311	1641.0
- RM	1	1871.32	12950	1666.6
- LSTAT	1	2410.84	13490	1687.3

Step: AIC=1587.65

myhousing\$MEDV ~ CRIM + ZN + INDUS + CHAS + NOX + RM + DIS +
RAD + TAX + PTRATIO + B + LSTAT

	Df	Sum of Sq	RSS	AIC
- INDUS	1	2.52	11081	1585.8
<none>			11079	1587.7
- CHAS	1	219.91	11299	1595.6
- TAX	1	242.24	11321	1596.6
- CRIM	1	243.20	11322	1596.6
- ZN	1	260.32	11339	1597.4
- B	1	272.26	11351	1597.9
- RAD	1	481.09	11560	1607.2
- NOX	1	520.87	11600	1608.9
- PTRATIO	1	1200.23	12279	1637.7
- DIS	1	1352.26	12431	1643.9
- RM	1	1959.55	13038	1668.0
- LSTAT	1	2718.88	13798	1696.7

Step: AIC=1585.76

myhousing\$MEDV ~ CRIM + ZN + CHAS + NOX + RM + DIS + RAD + TAX +
PTRATIO + B + LSTAT

	Df	Sum of Sq	RSS	AIC
<none>		11081	1585.8	
- CHAS	1	227.21	11309	1594.0
- CRIM	1	245.37	11327	1594.8
- ZN	1	257.82	11339	1595.4
- B	1	270.82	11352	1596.0
- TAX	1	273.62	11355	1596.1
- RAD	1	500.92	11582	1606.1
- NOX	1	541.91	11623	1607.9
- PTRATIO	1	1206.45	12288	1636.0
- DIS	1	1448.94	12530	1645.9
- RM	1	1963.66	13045	1666.3
- LSTAT	1	2723.48	13805	1695.0

Call:

lm(formula = myhousing\$MEDV ~ CRIM + ZN + CHAS + NOX + RM + DIS +
RAD + TAX + PTRATIO + B + LSTAT, data = myhousing)

Coefficients:

(Intercept)	CRIM	ZN	CHAS	NOX	RM	DIS	RAD
TAX	PTRATIO						
36.341145	-0.108413	0.045845	2.718716	-17.376023	3.801579	-1.492711	
0.299608	-0.011778	-0.946525					
B	LSTAT						
0.009291	-0.522553						