

R notes for Lecture 8

For all of these examples we use the tree data.

```
> tree.table <- read.table("tree.txt", header=T, sep="\t")
> names(tree.table)
[1] "Diameter" "Height"   "Volume"
> attach(tree.table)
```

1. Fit a regression

```
> plot(Height, Volume)
> tree.ht <- lm(Volume ~ Height)
> abline(tree.ht)
```

2. Evaluate residuals and find Cook's distances

```
> resid.ht <- residuals(tree.ht) #saves residuals
> hist(resid.ht) # crude check for normality, outliers
> qqnorm(resid.ht) # better check for normality
> qqline(resid.ht)
> plot(Height, resid.ht) # check for deviations from
linearity, non-constant variance
> plot(tree.ht)
```

Hit <Return> to see next plot:

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Hit <Return> to see next plot:

Hit <Return> to see next plot:

#2 out of 4 of these plots will show you residual diagnostic plots. Plot #4 gives Cook's distances.

3. Check the coefficients of the model

```
> coefficients(tree.ht)
(Intercept)      Height
-87.123614      1.543350
```

4. T-tests that the slope is 0 or intercept is 0, find r-squared, find standard errors of estimated intercept and slope

```
> summary(tree.ht)
```

Call:

```
lm(formula = Volume ~ Height)
```

Residuals:

Min	1Q	Median	3Q	Max
-21.274	-9.894	-2.894	12.067	29.852

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	-87.1236	29.2731	-2.976	0.005835	**
Height	1.5433	0.3839	4.021	0.000378	***

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

Residual standard error: 13.4 on 29 degrees of freedom

Multiple R-Squared: 0.3579, Adjusted R-squared: 0.3358

F-statistic: 16.16 on 1 and 29 DF, p-value: 0.0003784

5. Estimate mean volume for all trees with height 72, 78, 79.

```
> new <- data.frame(Height=c(72,78,79)) # You must first  
create a data frame with the new values in it.
```

```
> predict.lm(tree.ht, new, interval="confidence")
```

	fit	lwr	upr
1	23.99757	18.15976	29.83538
2	33.25767	28.09207	38.42327
3	34.80102	29.34525	40.25678

```
>
```

First column are the predictions: So the mean volume of trees 72 feet tall is 23.998. The next two columns give the lower and upper bounds on a 95% confidence interval. So the confidence interval for trees 72 feet tall is (18.16, 29.8).

6. Predict the volume of a tree that is 72 feet tall. Also one that is 78, and one that is 79.

```
> predict.lm(tree.ht, new, interval="prediction")
```

	fit	lwr	upr
1	23.99757	-4.017335	52.01247

2	33.25767	5.375087	61.14025
3	34.80102	6.863226	62.73881

The only change in the command was that interval is set to "prediction" instead of "confidence". Note that the first column is the same, but the confidence limits have changed. And the first set of limits show some problems with our model (we are predicting trees with negative volume).