

IBM Cloud

Data Science & Machine Learning 101 Technical Boot Camp

Lab Guide



Data Science



Machine Learning



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Lab Environment Overview

Installed Software and Tools

Software	Link
IBM Data Science Experience (DSX)	https://datascience.ibm.com/
IBM SPSS Statistics	http://www-03.ibm.com/software/products/no/spss-stats-base
Jupyter	http://jupyter.org/
GitHub	https://github.org/
Anaconda	https://www.anaconda.com/
RStudio	https://www.rstudio.com/

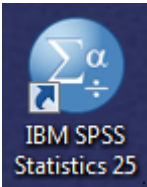
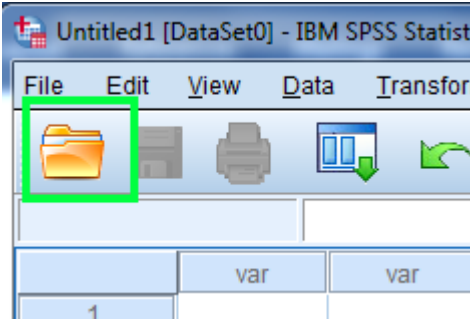
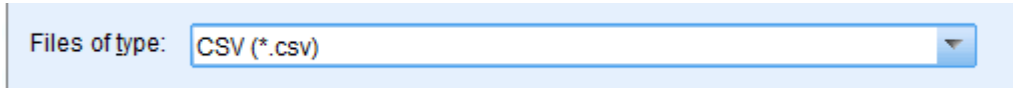
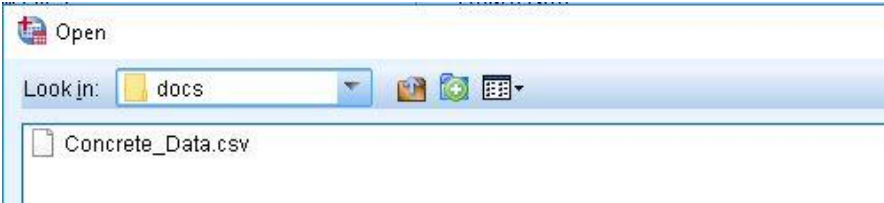
Module 2: Statistics

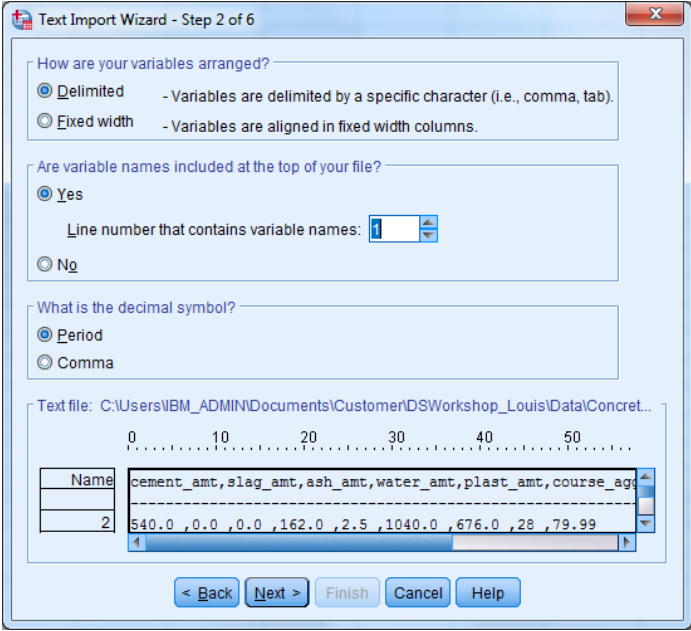
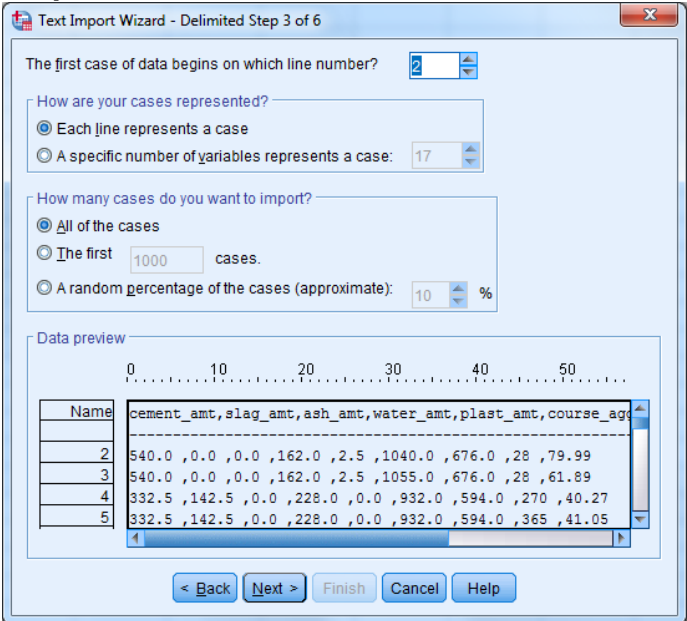
Purpose:	<p>This lab introduces the subject of statistics and the process of performing statistical analysis. After completing the lab, you should be able to:</p> <ul style="list-style-type: none">• Ingest an external data into IBM SPSS Statistics• Explore the characteristics of the dataset• Examine its descriptive statistics• Create a statistical model
Tasks:	<p>Tasks you will complete in this lab exercise include:</p> <ul style="list-style-type: none">• Load data• Exploratory Analysis<ul style="list-style-type: none">○ Analyze the data using visualizations○ Test the data for correlations○ Create a statistical model○ Measure model performance

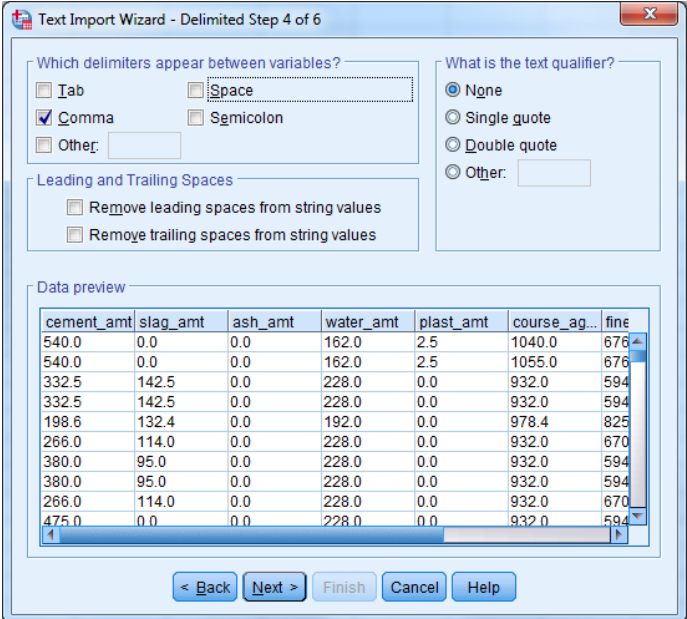
Module 2: Lab Workflow Overview

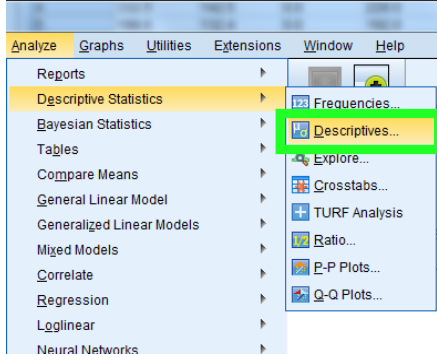
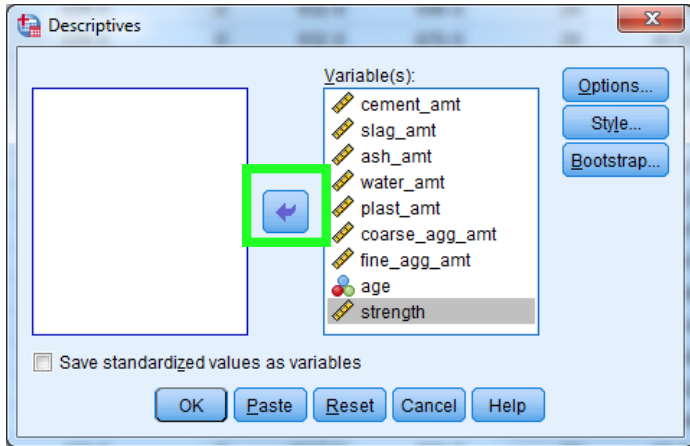
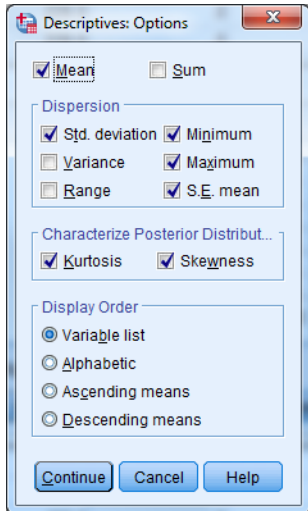
- 1 • Ingest external data
- 2 • Examine the data
- 3 • Investigate frequencies
- 4 • Explore correlations
- 5 • Create model and measure performance

Module 2: Lab Instructions

Step	Action
1	<p><u>Ingest external data</u></p> <ol style="list-style-type: none"> Double-click the IBM SPSS Statistics 25 icon.  <ol style="list-style-type: none"> Close the dialog box that appears. Click on the Open Data icon in the toolbar.  <ol style="list-style-type: none"> Change the file type, in the middle of the dialog box, to CSV by clicking in the dropdown box.  <ol style="list-style-type: none"> At the top of the dialog box, navigate to the Concrete_Data.CSV file. C:\Data Science Bootcamp\docs  <ol style="list-style-type: none"> Click on file name and then Open.

Step	Action
	<p data-bbox="370 310 1477 384">g. There are many options when reading a flat file. You will be lead through 6 steps to ensure the data is properly read. Just click Next on Step 1.</p> <p data-bbox="418 422 1073 457">In step 2, 3 and 4, match the selections below.</p> <p data-bbox="418 499 516 535">Step 2</p>  <p data-bbox="418 1203 516 1239">Step 3</p> 

Step	Action
	<p>Step 4</p>  <p>In step 5, click Next. In step 6, click Finish.</p> <ul style="list-style-type: none"> h. A new window, called the Output window, appears and then moves to the background. This records all activities as the product is used and is where the results of the procedures performed on the data will be shown. i. It is helpful to know that SPSS can read many data formats. You can get a sense for this by exploring the various options for opening data files in the File menu.
2	<p><u>Examine the data</u></p> <ul style="list-style-type: none"> a. Click the Analyze menu in the menu bar. Hover over Descriptive Statistics and click Descriptives in the sub-menu.

Step	Action
	<div data-bbox="410 268 844 621">  </div> <p data-bbox="370 657 1477 730">b. Select all the variables; then, click the arrow box to move all the variables to the Variable(s) box on the right.</p> <div data-bbox="410 762 1096 1205">  </div> <p data-bbox="370 1245 1205 1281">c. Click Options. Make the selections as indicated below.</p> <div data-bbox="410 1312 716 1820">  </div>

Step

Action

d. Click **Continue**. Then, click **OK** in the Descriptives dialog.

e. Examine the statistics about each variable.

```

DATASET NAME DataSet1 WINDOW=FRONT.
DESCRIPTIVES VARIABLES=cement_amt slag_amt ash_amt water_amt plast_amt coarse_agg_amt fine_agg_amt
age strength
/STATISTICS=MEAN STDDEV MIN MAX SEMEAN KURTOSIS SKEWNESS.

```

➔ Descriptives

[DataSet1]

Descriptive Statistics

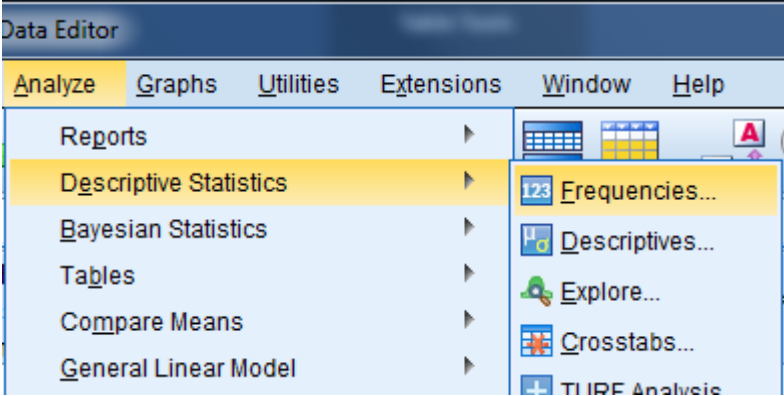
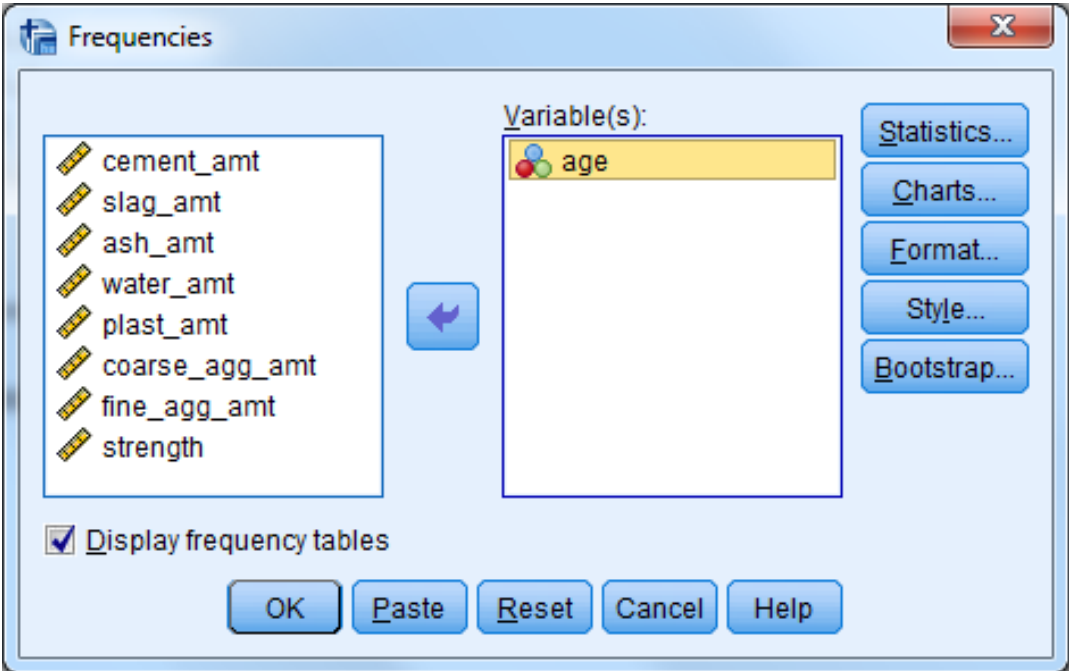
	N	Minimum	Maximum	Mean		Std. Deviation	Skewness		Kurtosis	
	Statistic	Statistic	Statistic	Statistic	Std. Error	Statistic	Statistic	Std. Error	Statistic	Std. Error
cement_amt	1030	102.0	540.0	281.168	3.2563	104.5064	.509	.076	-.521	.152
slag_amt	1030	.0	359.4	73.896	2.6884	86.2793	.801	.076	-.508	.152
ash_amt	1030	.0	200.1	54.188	1.9941	63.9970	.537	.076	-1.329	.152
water_amt	1030	121.8	247.0	181.567	.6654	21.3542	.075	.076	.122	.152
plast_amt	1030	.0	32.2	6.205	.1861	5.9738	.907	.076	1.411	.152
coarse_agg_amt	1030	801.0	1145.0	972.919	2.4227	77.7540	-.040	.076	-.599	.152
fine_agg_amt	1030	594.0	992.6	773.580	2.4982	80.1760	-.253	.076	-.102	.152
age	1030	1	365	45.66	1.968	63.170	3.269	.076	12.169	.152
strength	1030	2.33	82.60	35.8180	.52053	16.70574	.417	.076	-.314	.152
Valid N (listwise)	1030									

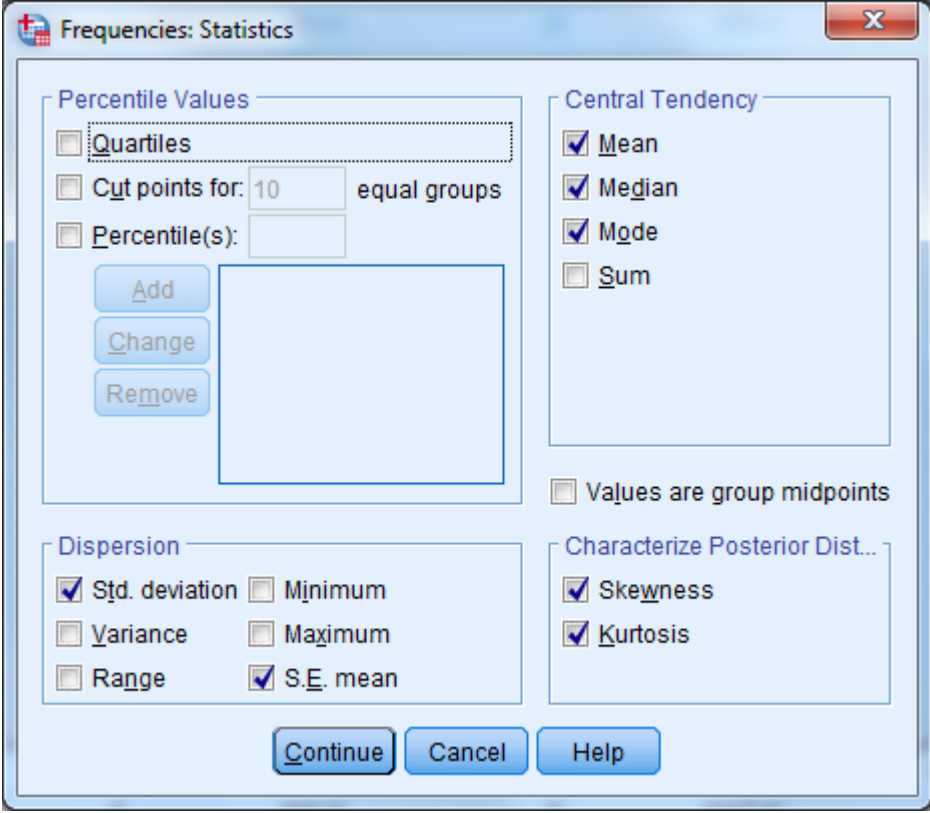
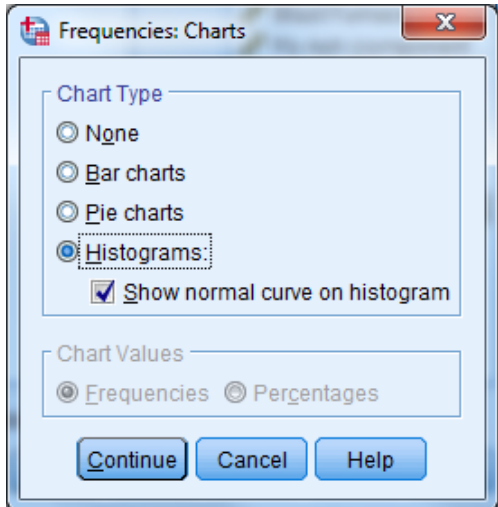
As shown above this table, there is SPSS generated code.

Do you see terms that you recognize?_____

What does the code mean? _____

Is there a relationship between the mean, standard deviation and skewness? If yes, explain.

Step	Action
3	<p><u>Investigate frequencies</u></p> <p>a. Click the Analyze menu in the menu bar. Hover over Descriptive Statistics and click Frequencies in the sub-menu.</p>  <p>b. Move the Age variable to the Variable(s) box. Ensure the 'Display frequency tables' box is checked.</p> 

Step	Action
	<p>c. Click Statistics and match the selections below.</p>  <p>d. Click Continue, then Charts. Match the selections below.</p> 

Step

Action

e. Click **Continue**; then, **OK**. You will see the descriptive statistics; the same information you saw in the previous exercise.

➔

Frequencies

Statistics

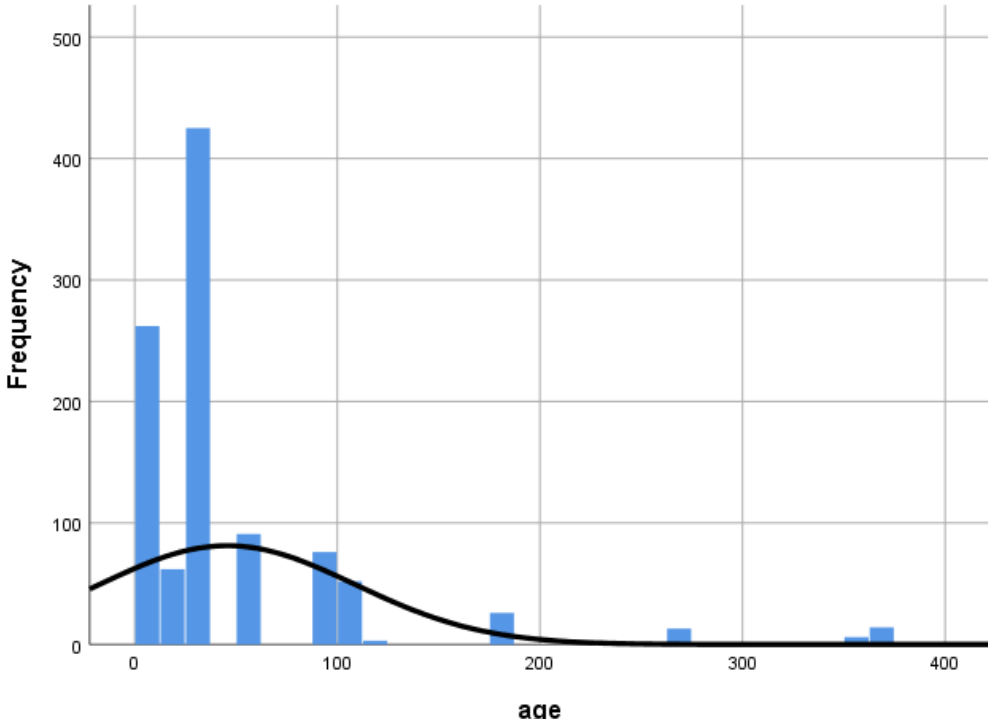
age

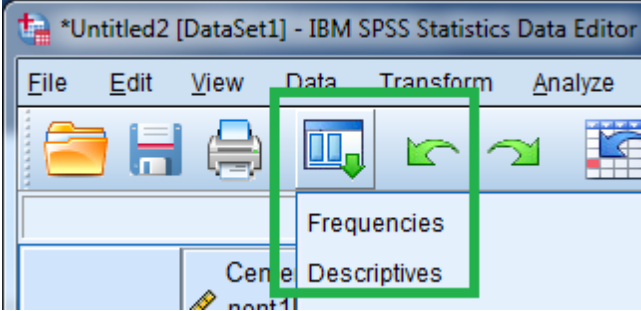
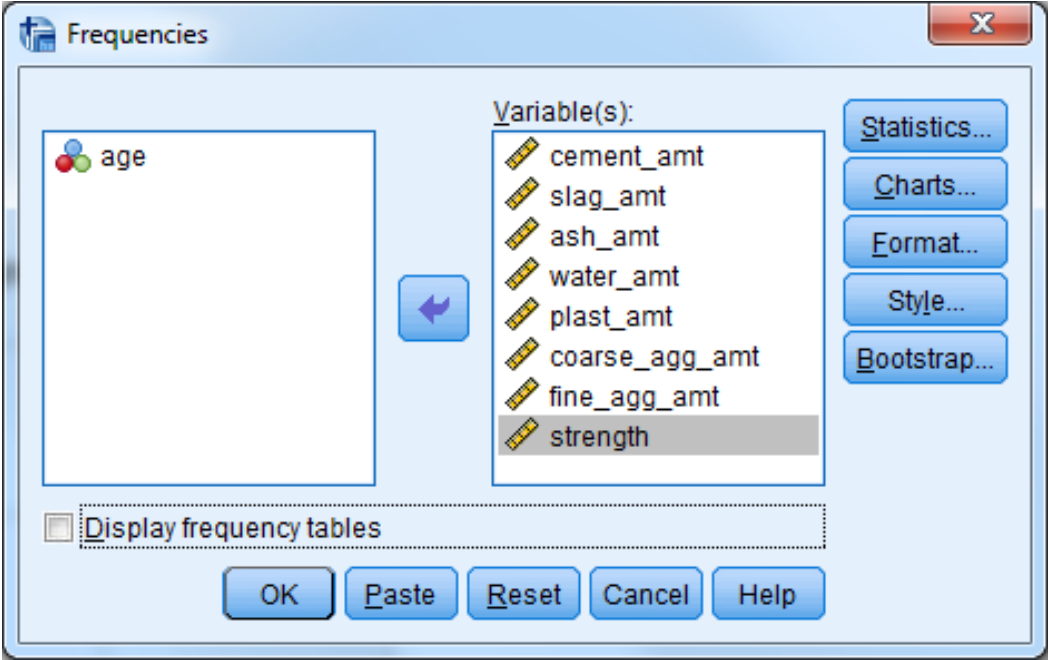
N	Valid	1030
	Missing	0
Mean		45.66
Std. Error of Mean		1.968
Median		28.00
Mode		28
Std. Deviation		63.170
Skewness		3.269
Std. Error of Skewness		.076
Kurtosis		12.169
Std. Error of Kurtosis		.152

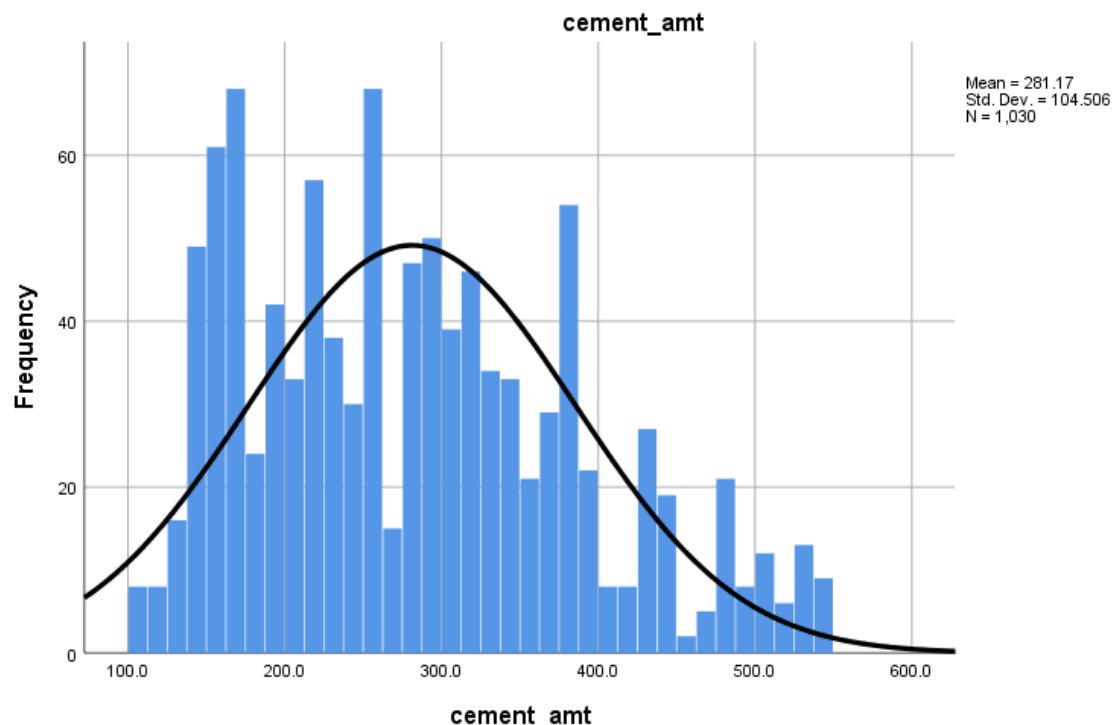
f. Followed by the frequency table.

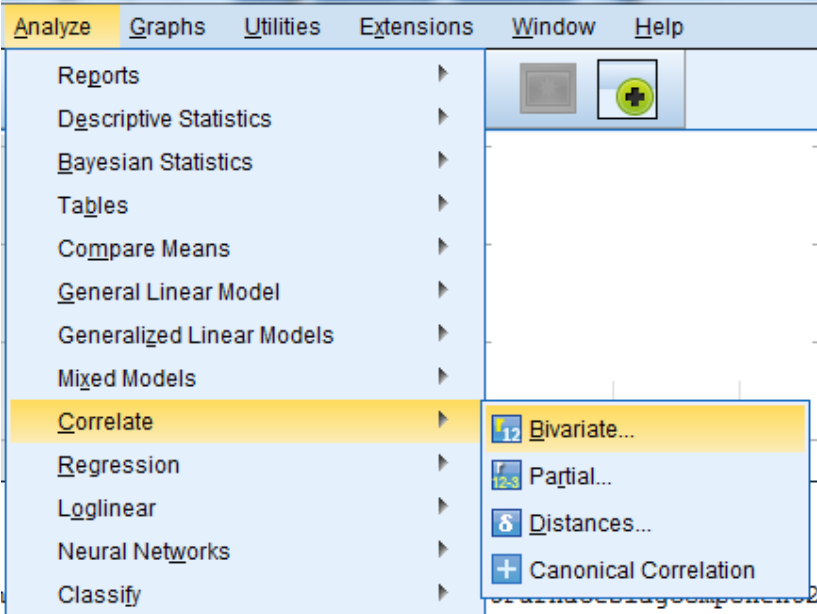
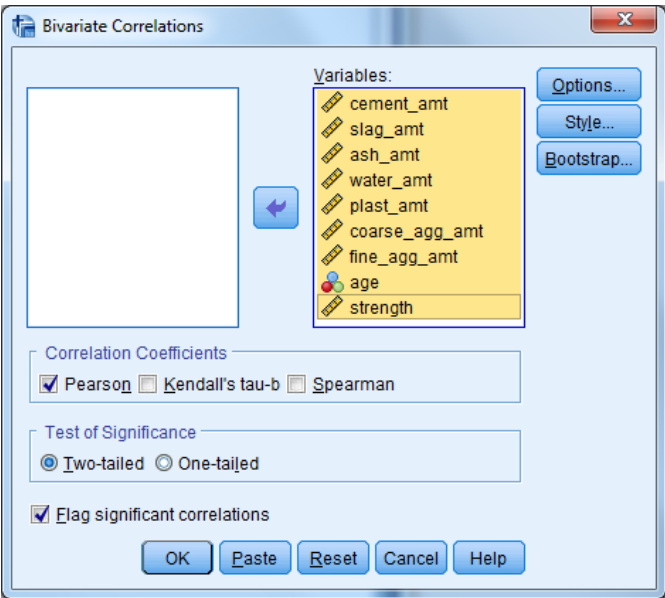
age

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	1	2	.2	.2	.2
	3	134	13.0	13.0	13.2
	7	126	12.2	12.2	25.4
	14	62	6.0	6.0	31.5
	28	425	41.3	41.3	72.7
	56	91	8.8	8.8	81.6
	90	54	5.2	5.2	86.8
	91	22	2.1	2.1	88.9
	100	52	5.0	5.0	94.0
	120	3	.3	.3	94.3
	180	26	2.5	2.5	96.8
	270	13	1.3	1.3	98.1
	360	6	.6	.6	98.6
	365	14	1.4	1.4	100.0
	Total	1030	100.0	100.0	

Step	Action
	<p data-bbox="370 306 1068 342">g. This is followed by a histogram of the variable.</p> <div data-bbox="321 396 1459 1152"> <p data-bbox="911 401 1036 428">Histogram</p>  <p data-bbox="1317 470 1451 527">Mean = 45.66 Std. Dev. = 63.17 N = 1,030</p> </div> <p data-bbox="370 1222 1500 1367">h. The histogram includes a curved line that represents a normal distribution of the variable. It appears that this variable is not normally distributed. See the descriptive information and discuss the information as it relates to the normality of the data.</p> <p data-bbox="418 1442 626 1474">Think on this:</p> <p data-bbox="418 1478 1036 1514">What do you notice about the Age variable?</p> <p data-bbox="418 1518 1403 1549">How does this relate to the descriptive statistics we discussed earlier?</p> <hr data-bbox="418 1577 1446 1583"/> <hr data-bbox="418 1650 1446 1656"/>

Step	Action
	<p>i. Use the dialog recall button in the toolbar to bring up the Frequencies dialog again.</p>  <p>j. Move the Age variable out of the Variable(s) box; and add all the other variables. Make sure to uncheck the box in the lower left.</p> <p>Think on this: Do you know why this is a good thing to do? Leave it checked if you need some insight.</p> 

Step	Action																																																																																																																																	
	<p>k. Click OK. You see the frequency statistics for each variable.</p> <p>➔ Frequencies</p> <table><tr><th colspan="2"></th><th colspan="8">Statistics</th></tr><tr><th colspan="2"></th><th>cement_amt</th><th>slag_amt</th><th>ash_amt</th><th>water_amt</th><th>plast_amt</th><th>coarse_agg_amt</th><th>fine_agg_amt</th><th>strength</th></tr><tr><td rowspan="2">N</td><td>Valid</td><td>1030</td><td>1030</td><td>1030</td><td>1030</td><td>1030</td><td>1030</td><td>1030</td><td>1030</td></tr><tr><td>Missing</td><td>0</td><td>0</td><td>0</td><td>0</td><td>0</td><td>0</td><td>0</td><td>0</td></tr><tr><td colspan="2">Mean</td><td>281.168</td><td>73.896</td><td>54.188</td><td>181.567</td><td>6.205</td><td>972.919</td><td>773.580</td><td>35.8180</td></tr><tr><td colspan="2">Std. Error of Mean</td><td>3.2563</td><td>2.6884</td><td>1.9941</td><td>.6654</td><td>.1861</td><td>2.4227</td><td>2.4982</td><td>.52053</td></tr><tr><td colspan="2">Median</td><td>272.900</td><td>22.000</td><td>.000</td><td>185.000</td><td>6.400</td><td>968.000</td><td>779.500</td><td>34.4450</td></tr><tr><td colspan="2">Mode</td><td>362.6^a</td><td>.0</td><td>.0</td><td>192.0</td><td>.0</td><td>932.0</td><td>594.0^a</td><td>33.40</td></tr><tr><td colspan="2">Std. Deviation</td><td>104.5064</td><td>86.2793</td><td>63.9970</td><td>21.3542</td><td>5.9738</td><td>77.7540</td><td>80.1760</td><td>16.70574</td></tr><tr><td colspan="2">Skewness</td><td>.509</td><td>.801</td><td>.537</td><td>.075</td><td>.907</td><td>-.040</td><td>-.253</td><td>.417</td></tr><tr><td colspan="2">Std. Error of Skewness</td><td>.076</td><td>.076</td><td>.076</td><td>.076</td><td>.076</td><td>.076</td><td>.076</td><td>.076</td></tr><tr><td colspan="2">Kurtosis</td><td>-.521</td><td>-.508</td><td>-1.329</td><td>.122</td><td>1.411</td><td>-.599</td><td>-.102</td><td>-.314</td></tr><tr><td colspan="2">Std. Error of Kurtosis</td><td>.152</td><td>.152</td><td>.152</td><td>.152</td><td>.152</td><td>.152</td><td>.152</td><td>.152</td></tr></table> <p>a. Multiple modes exist. The smallest value is shown</p>			Statistics										cement_amt	slag_amt	ash_amt	water_amt	plast_amt	coarse_agg_amt	fine_agg_amt	strength	N	Valid	1030	1030	1030	1030	1030	1030	1030	1030	Missing	0	0	0	0	0	0	0	0	Mean		281.168	73.896	54.188	181.567	6.205	972.919	773.580	35.8180	Std. Error of Mean		3.2563	2.6884	1.9941	.6654	.1861	2.4227	2.4982	.52053	Median		272.900	22.000	.000	185.000	6.400	968.000	779.500	34.4450	Mode		362.6 ^a	.0	.0	192.0	.0	932.0	594.0 ^a	33.40	Std. Deviation		104.5064	86.2793	63.9970	21.3542	5.9738	77.7540	80.1760	16.70574	Skewness		.509	.801	.537	.075	.907	-.040	-.253	.417	Std. Error of Skewness		.076	.076	.076	.076	.076	.076	.076	.076	Kurtosis		-.521	-.508	-1.329	.122	1.411	-.599	-.102	-.314	Std. Error of Kurtosis		.152	.152	.152	.152	.152	.152	.152	.152
		Statistics																																																																																																																																
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N	Valid	1030	1030	1030	1030	1030	1030	1030	1030																																																																																																																									
	Missing	0	0	0	0	0	0	0	0																																																																																																																									
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Skewness		.509	.801	.537	.075	.907	-.040	-.253	.417																																																																																																																									
Std. Error of Skewness		.076	.076	.076	.076	.076	.076	.076	.076																																																																																																																									
Kurtosis		-.521	-.508	-1.329	.122	1.411	-.599	-.102	-.314																																																																																																																									
Std. Error of Kurtosis		.152	.152	.152	.152	.152	.152	.152	.152																																																																																																																									
	<p>Followed by a histogram for each variable as a visual overview. Here is an example.</p> <p>Histogram</p> 																																																																																																																																	

Step	Action
4	<p><u>Explore correlations</u></p> <p>Next, explore the relationships between different variables.</p> <p>a. In the Analyze menu, go down to the Correlate entry and click on Bivariate.</p>  <p>b. Move all the variables to the Variables box and select the options as shown:</p> 

Step

Action

Click **OK**. You will see the following correlation table.

➔ Correlations

Correlations

		cement_amt	slag_amt	ash_amt	water_amt	plast_amt	coarse_agg_amt	fine_agg_amt	age	strength
cement_amt	Pearson Correlation	1	-.275**	-.397**	-.082**	.092**	-.109**	-.223**	.082**	.498**
	Sig. (2-tailed)		.000	.000	.009	.003	.000	.000	.009	.000
	N	1030	1030	1030	1030	1030	1030	1030	1030	1030
slag_amt	Pearson Correlation	-.275**	1	-.324**	.107**	.043	-.284**	-.282**	-.044	.135**
	Sig. (2-tailed)	.000		.000	.001	.165	.000	.000	.156	.000
	N	1030	1030	1030	1030	1030	1030	1030	1030	1030
ash_amt	Pearson Correlation	-.397**	-.324**	1	-.257**	.378**	-.010	.079*	-.154**	-.106**
	Sig. (2-tailed)	.000	.000		.000	.000	.750	.011	.000	.001
	N	1030	1030	1030	1030	1030	1030	1030	1030	1030
water_amt	Pearson Correlation	-.082**	.107**	-.257**	1	-.658**	-.182**	-.451**	.278**	-.290**
	Sig. (2-tailed)	.009	.001	.000		.000	.000	.000	.000	.000
	N	1030	1030	1030	1030	1030	1030	1030	1030	1030
plast_amt	Pearson Correlation	.092**	.043	.378**	-.658**	1	-.266**	.223**	-.193**	.366**
	Sig. (2-tailed)	.003	.165	.000	.000		.000	.000	.000	.000
	N	1030	1030	1030	1030	1030	1030	1030	1030	1030
coarse_agg_amt	Pearson Correlation	-.109**	-.284**	-.010	-.182**	-.266**	1	-.178**	-.003	-.165**
	Sig. (2-tailed)	.000	.000	.750	.000	.000		.000	.923	.000
	N	1030	1030	1030	1030	1030	1030	1030	1030	1030
fine_agg_amt	Pearson Correlation	-.223**	-.282**	.079*	-.451**	.223**	-.178**	1	-.156**	-.167**
	Sig. (2-tailed)	.000	.000	.011	.000	.000	.000		.000	.000
	N	1030	1030	1030	1030	1030	1030	1030	1030	1030
age	Pearson Correlation	.082**	-.044	-.154**	.278**	-.193**	-.003	-.156**	1	.329**
	Sig. (2-tailed)	.009	.156	.000	.000	.000	.923	.000		.000
	N	1030	1030	1030	1030	1030	1030	1030	1030	1030
strength	Pearson Correlation	.498**	.135**	-.106**	-.290**	.366**	-.165**	-.167**	.329**	1
	Sig. (2-tailed)	.000	.000	.001	.000	.000	.000	.000	.000	
	N	1030	1030	1030	1030	1030	1030	1030	1030	1030

** . Correlation is significant at the 0.01 level (2-tailed).

* . Correlation is significant at the 0.05 level (2-tailed).

Scan the rows labelled ‘Pearson Correlation’.

What is the largest positive correlation coefficient? _____

What is the largest negative correlation coefficient? _____

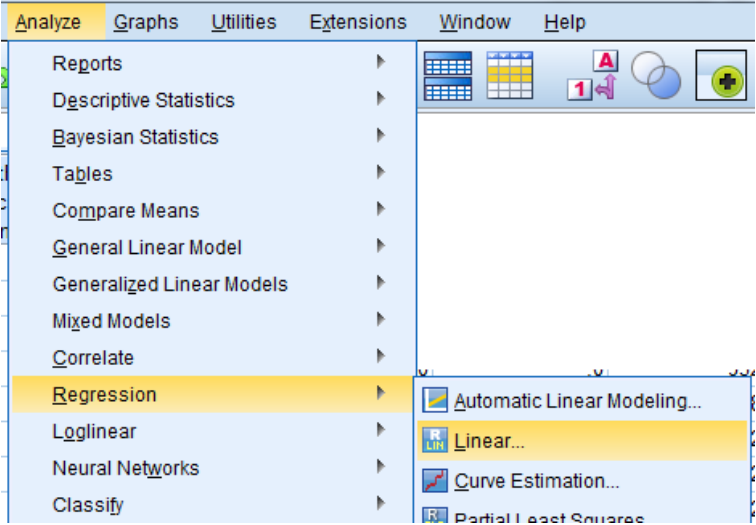
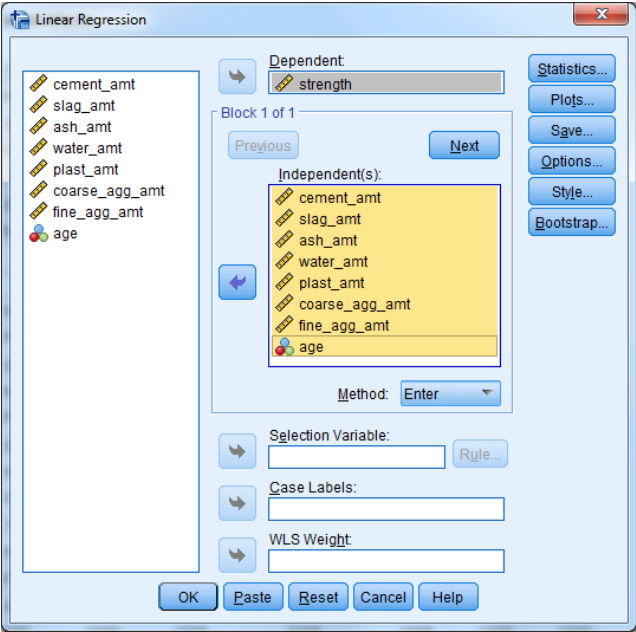
What are some of the significant correlations? _____

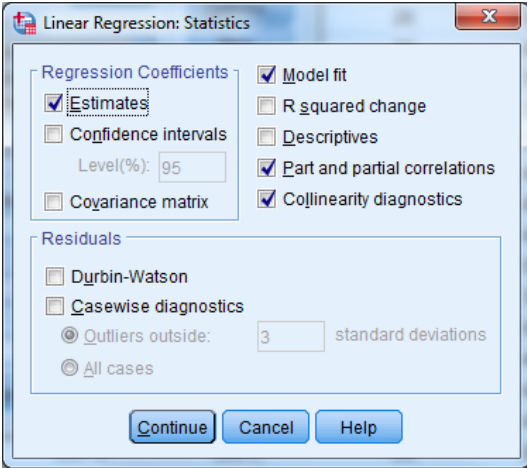
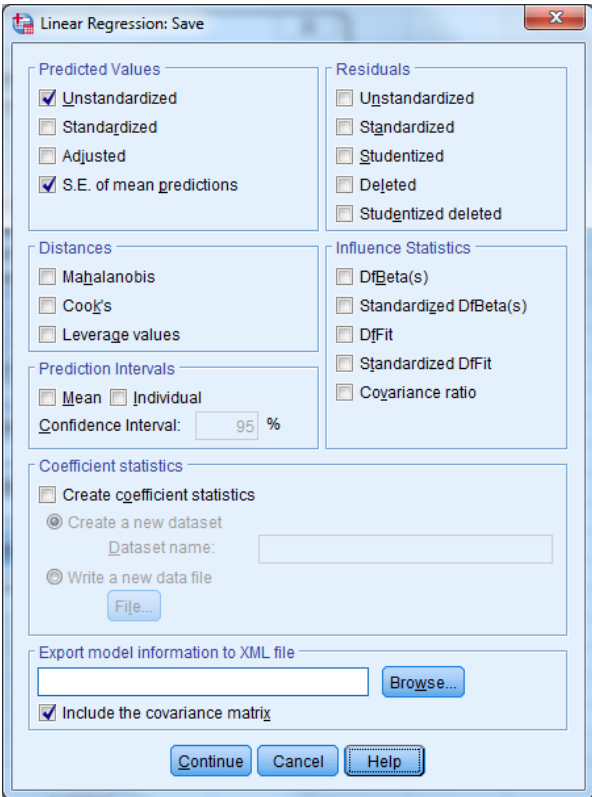
Do any of them surprise you? _____

5

Create model and measure performance

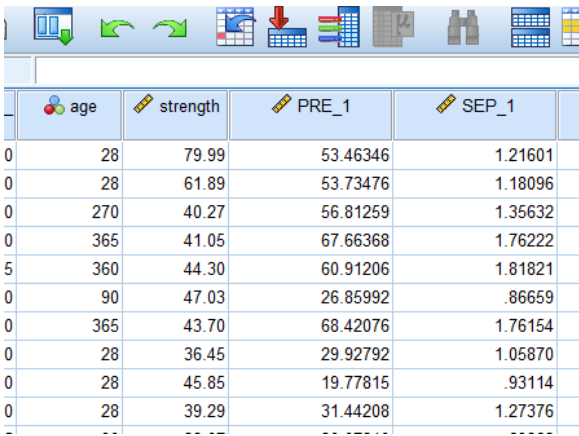
Creating a model of the data (a mathematical representation of what goes on in the data) is a goal in data science. A model can be used to gain insights from the data, but is more often viewed as a means to determine future outcomes. For example, you can get an idea of what the compressive strength of concrete would be if you varied the amount of one or more ingredients.

Step	Action
	<p>In this exercise, a multiple linear regression model will be built and evaluated.</p> <p>a. In the Analyze menu, hover over the Regression select then click on Linear in the sub-menu.</p>  <p>b. In the Linear Regression dialog, move 'Cement compressive strength' to the Dependent box and all the other variables to the Independent(s) box as shown.</p> 

Step	Action
	<p>c. Click Statistics and make the selections as shown below.</p>  <p>d. Click Continue.</p> <p>e. Click the Save button and make the selections as shown.</p> 

Step	Action																																						
	<p>These selections add variables to the dataset to determine how well the model is performing. Click Continue.</p> <p>f. Click OK; then, review the results.</p> <p>g. Scroll down to the Model Summary. The R Square number indicates that about 61% of the variability in Compressive strength is explained by the model.</p> <div><p>Model Summary^b</p><table><tr><th>Model</th><th>R</th><th>R Square</th><th>Adjusted R Square</th><th>Std. Error of the Estimate</th></tr><tr><td>1</td><td>.785^a</td><td>.616</td><td>.613</td><td>10.39914</td></tr></table><p>a. Predictors: (Constant), age, coarse_agg_amt, cement_amt, fine_agg_amt, plast_amt, ash_amt, water_amt, slag_amt</p><p>b. Dependent Variable: strength</p></div> <p>h. The next table in the Output window shows a significant F statistic, which means the model is better than guessing.</p> <div><p>ANOVA^a</p><table><tr><th>Model</th><th></th><th>Sum of Squares</th><th>df</th><th>Mean Square</th><th>F</th><th>Sig.</th></tr><tr><td>1</td><td>Regression</td><td>176762.034</td><td>8</td><td>22095.254</td><td>204.317</td><td>.000^b</td></tr><tr><td></td><td>Residual</td><td>110413.153</td><td>1021</td><td>108.142</td><td></td><td></td></tr><tr><td></td><td>Total</td><td>287175.187</td><td>1029</td><td></td><td></td><td></td></tr></table><p>a. Dependent Variable: strength</p><p>b. Predictors: (Constant), age, coarse_agg_amt, cement_amt, fine_agg_amt, plast_amt, ash_amt, water_amt, slag_amt</p></div> <p>i. The model as a whole appears to be accurate at predicting the concrete strength. However, there are a few variables that don't have a strong influence on the output.</p>	Model	R	R Square	Adjusted R Square	Std. Error of the Estimate	1	.785 ^a	.616	.613	10.39914	Model		Sum of Squares	df	Mean Square	F	Sig.	1	Regression	176762.034	8	22095.254	204.317	.000 ^b		Residual	110413.153	1021	108.142				Total	287175.187	1029			
Model	R	R Square	Adjusted R Square	Std. Error of the Estimate																																			
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	Total	287175.187	1029																																				

Step	Action
	<div><div><div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div><div></div></div><div><div></div><div></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Step	Action																		
	<table border="1"> <caption>Collinearity Statistics</caption> <thead> <tr> <th>Tolerance</th><th>VIF</th></tr> </thead> <tbody> <tr><td>.134</td><td>7.489</td></tr> <tr><td>.137</td><td>7.277</td></tr> <tr><td>.162</td><td>6.171</td></tr> <tr><td>.143</td><td>7.005</td></tr> <tr><td>.337</td><td>2.965</td></tr> <tr><td>.197</td><td>5.076</td></tr> <tr><td>.143</td><td>7.005</td></tr> <tr><td>.894</td><td>1.118</td></tr> </tbody> </table> <p>Low tolerances, such as you see here, indicate high multicollinearity of the variables. Its associated Variable Inflation Factor (VIF) is considered problematic when it gets higher than 2. You see in the list that, except for Age, all are much higher than 2. So, our model might not be as useful as it could be.</p> <p>k. Next, look at what the model actually did with the data. By saving the predicted values, they will be seen as a new variable in the dataset.</p>  <p>PRE_1 is the variable where the model wrote the predicted value. That is, the model was built, then the data was scored by the model. SEP_1 is the Standard Error of the prediction. Comparing PRE_1 to Concrete compressive strength (the dependent variable) you begin to see that there are some big differences.</p>	Tolerance	VIF	.134	7.489	.137	7.277	.162	6.171	.143	7.005	.337	2.965	.197	5.076	.143	7.005	.894	1.118
Tolerance	VIF																		
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Step	Action
	<p data-bbox="404 306 613 338">Think on this:</p> <p data-bbox="404 344 1195 375">What further actions could be taken to refine the model?</p> <hr data-bbox="404 407 1414 411"/> <hr data-bbox="404 480 1414 485"/>

Module 2: Lab Summary

In this module, SPSS Statistics was used to examine the characteristics of individual variables within a dataset. An analysis was performed to better understand the shape and size of the data and discover relationships among the variables. Further evaluation was completed by using regression models in order to describe the relationships of the variables. Finally, the regression model's performance was measured for its predictive accuracy.

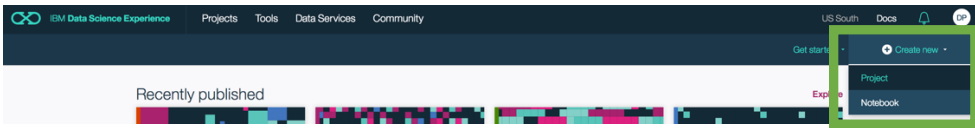
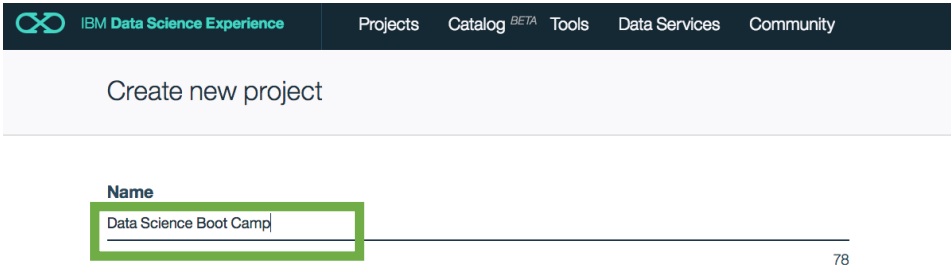

Module 3: Machine Learning Lab I


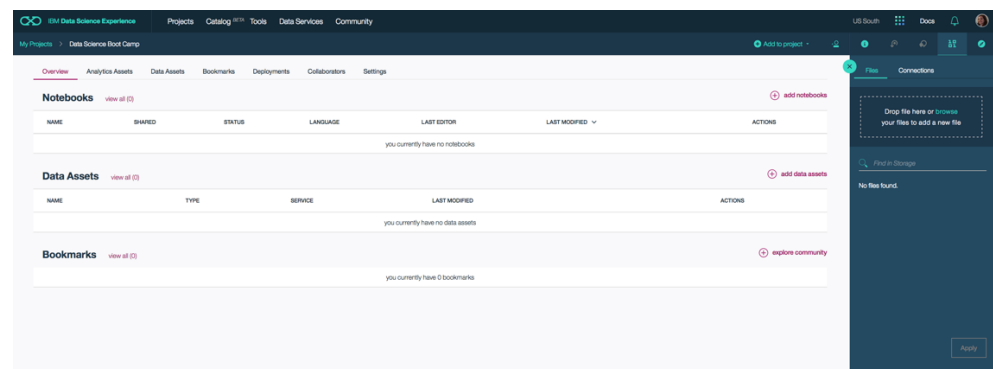
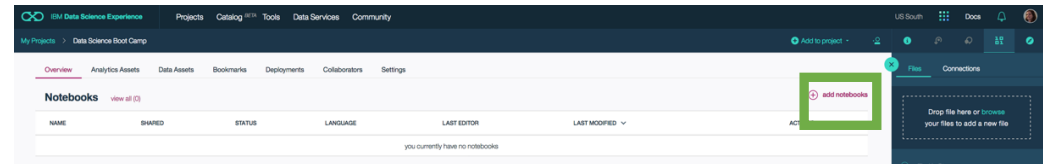
Purpose:	<p>This lab introduces the tools used for a Machine Learning project written in R. After completing the lab, you should be able to:</p> <ul style="list-style-type: none">• Pull data from a GitHub repository into a Jupyter notebook• Perform an exploratory analysis of a dataset in IBM's Data Science Experience (DSX)• Create training and testing datasets
Tasks:	<p>Tasks you will complete in this lab exercise include:</p> <ul style="list-style-type: none">• Install and load R libraries• Exploration and Analysis<ul style="list-style-type: none">○ Analyze the data using visualizations○ Test the data for correlations○ Create training and testing datasets

Module 3: Lab Workflow Overview

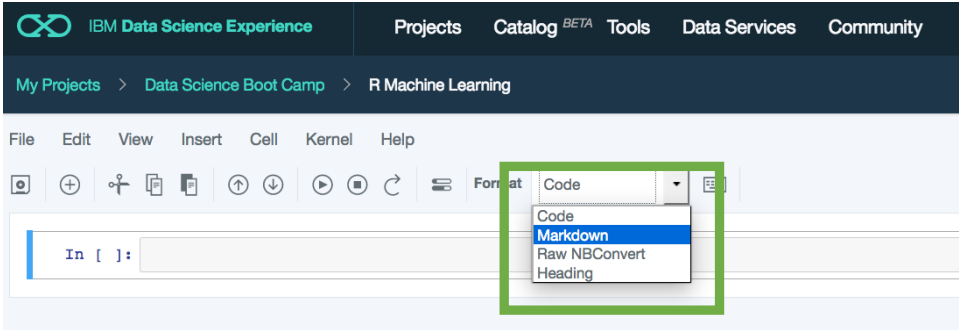
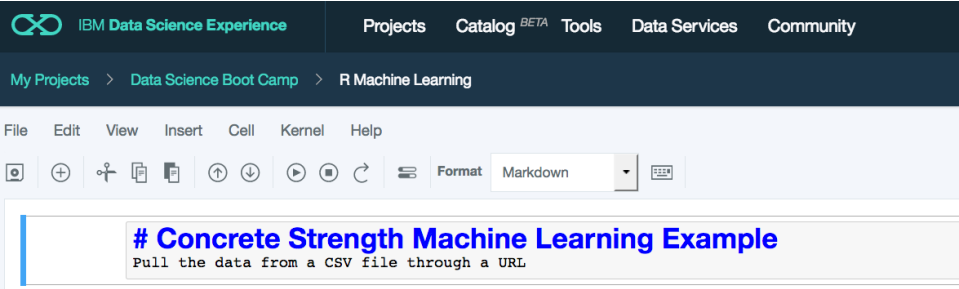
- 1 • Open DSX and Create a new project
- 2 • Create a notebook
- 3 • Add a markdown title
- 4 • Install necessary libraries
- 5 • Load libraries
- 6 • Pull data from GitHub
- 7 • Set the seed
- 8 • Check the data was loaded and the size
- 9 • Examine the descriptive statistics
- 10 • Observe the distribution of each variable
- 11 • Determine the correlation value
- 12 • Create training and testing datasets

Module 3: Lab Instructions

Step	Action
1	<p><u>Open DSX and Create a new project</u></p> <ol style="list-style-type: none"> Navigate to https://datascience.ibm.com/ Login to DSX On the top right side, click Create New and select project  <ol style="list-style-type: none"> Type the Project Name Data Science Boot Camp  <ol style="list-style-type: none"> Type the Description Boot Camp  <ol style="list-style-type: none"> Ensure the defaults are selected as follows: Select your Spark Service DSX-Spark Select Object Storage(Swift API) Select Target Object Storage Instance DSX-ObjectStorage Default Target Container DataScienceBootCamp

Step	Action
	<div data-bbox="324 268 1442 604"> <p>Spark Service</p> <p>DSX-Spark ▼</p> <p>⚠ If you associate the same Spark service with multiple projects, the Spark history server will display job history information for all the projects.</p> <p>Storage Type</p> <p><input checked="" type="radio"/> Object Storage (Swift API) <input type="radio"/> Object Storage (S3 API)</p> <p>Target Object Storage Instance</p> <p>DSX-ObjectStorage ▼</p> <p>Target Container <input checked="" type="checkbox"/></p> <p>DataScienceBootcamp</p> <p>Cancel Create </p> </div> <p data-bbox="357 640 592 682">g. Click Create</p> <div data-bbox="357 709 1344 1075">  </div>
2	<p><u>Create a notebook</u></p> <p>a. Click Add Notebooks</p> <div data-bbox="357 1255 1393 1417">  </div> <p>b. Type Notebook Name R Machine Learning</p> <div data-bbox="357 1522 1364 1795"> <p>Create Notebook</p> <p>Blank From File From URL</p> <p>Name*</p> <p>R Machine Learning</p> <p>32 Characters Remaining</p> </div>

Step	Action
	<p>c. Type the Description Machine Learning notebook in R</p> <div data-bbox="386 363 1010 541"> <p>Description</p> <div>Machine learning notebook in R</div> <div>470 Characters Remaining</div> </div> <p>d. Select R for the language</p> <div data-bbox="394 615 950 703"> <p>Language*</p> <p><input type="radio"/> Python 2 <input checked="" type="radio"/> R <input type="radio"/> Scala <input type="radio"/> Python 3.5 Experimental</p> </div> <p>e. Select 2.1 for the Spark version</p> <div data-bbox="386 800 628 867"> <p>Spark version*</p> <p><input checked="" type="radio"/> 2.1 <input type="radio"/> 2.0 <input type="radio"/> 1.6</p> </div> <p>f. Select the Spark Service DSX-Spark</p> <div data-bbox="363 961 1133 1039"> <p>Spark Service*</p> <p>DSX-Spark</p> </div> <p>g. Click Create Notebook</p> <div data-bbox="363 1113 1323 1192"> <p>Cancel Create Notebook</p> </div>

Step	Action
	<p>a. Select the format to be Markdown</p>  <p>b. Type in the cell:</p> <pre># Concrete Strength Machine Learning Example Pull the data from a CSV file through a URL</pre> <p>c. Click run (play button) or you can use the shortcut “shift + enter” to execute the cell.</p> 
4	<p><u>Install necessary libraries</u></p> <p>Many R functions come in packages, which are free libraries of code written by R's active user community. There are thousands of helpful R packages but this lab will only require the following:</p> <p><i>Corrplot</i>: graphical display of a correlation matrix, confidence interval</p> <p><i>Psych</i>: basic descriptive statistics useful for psychometrics</p> <p><i>Caret</i>: set of functions that streamline the process for creating predictive models</p> <p><i>MASS</i>: functions and datasets to support Venables and Ripley, “Modern Applied Statistics with S”</p>

Step	Action
	<p><i>Relaimpo</i>: provides several metrics for assessing relative importance in linear models where they can be printed, plotted and bootstrapped</p> <p>a. Enter the code:</p> <pre># Install Libraries install.packages("corrplot") install.packages("psych") install.packages("caret") install.packages("MASS") install.packages("relaimpo")</pre> <p>b. Run cell</p> <pre>In [1]: # Install Libraries install.packages("corrplot") install.packages("psych") install.packages("caret") install.packages("MASS") install.packages("relaimpo")</pre> <p>Note: Installing the libraries may take some time. Once installed, a red box will appear with an installation confirmation. This is normal and informational only. A similar red box will appear in the next step as well and is normal when loading libraries.</p>
5	<p><u>Load libraries</u></p> <p>Loading libraries gives you access to the functions that they contain. By using libraries, programmers can focus on the task at hand and not worry about developing functions that the user community has already developed.</p> <p>a. Enter the code:</p> <pre># Load libraries library(corrplot) library(psych) library(caret) library(MASS) library(relaimpo)</pre> <p>Note: A red box will appear. This is normal and informational only.</p>

Step	Action
	<p>b. Run cell</p> <pre data-bbox="500 411 878 590">In [2]: # Load libraries library(corrplot) library(psych) library(caret) library(MASS) library(relaimpo)</pre>
6	<p><u>Pull data from GitHub</u></p> <p>Data can be brought into DSX in multiple ways. For this lab, you will pull a data file from GitHub related to concrete strength. The data will be stored as a data frame named concrete. A data frame is an in-memory storage format that is representative of the csv data, and accessed via a variable name.</p> <p>a. Enter the code:</p> <pre data-bbox="418 982 1528 1056">concrete <- read.csv(url("https://raw.githubusercontent.com/team-wolfpack/DS-Boot-Camp/master/data/Concrete_Data.csv"))</pre> <p>b. Run cell</p> <pre data-bbox="418 1146 1471 1167">In [3]: concrete <- read.csv(url("https://raw.githubusercontent.com/team-wolfpack/DS-Boot-Camp/master/data/Concrete_Data.csv"))</pre>
7	<p><u>Set the seed</u></p> <p>Generally, in statistics, samples are chosen at random. A random number generator is used to select the samples and is based off of a seed value. The seed is explicitly set so results are reproducible. To ensure everyone retrieves the same results in this lab, the seed value was randomly chosen as 3482.</p> <p>a. Enter the code</p> <pre data-bbox="483 1591 987 1665"># Set seed to ensure reproducibility set.seed(3482)</pre> <p>b. Run cell</p> <pre data-bbox="483 1759 1154 1812">In [4]: # Set seed to ensure reproducibility set.seed(3482)</pre>

Step	Action
8	<p><u>Check the data was loaded and the size</u></p> <p>To check that data is present, the head command is used to retrieve the first few rows of data from the data frame specified.</p> <p>a. Enter the code</p> <pre># Ensure the data was loaded head(concrete)</pre> <p>Note: You can access documentation pages for R functions, datasets and other objects directly by entering the command ?function. Example: ?head If you want more information on the commands used in this lab, you should access them to better understand the code being entered.</p> <p>b. Run cell</p> <pre>In [5]: # Ensure the data was loaded head(concrete)</pre> <p>c. For the observed greatest strength, what are the values for cement_amt_____ and water_amt_____?</p> <p>d. Enter the code</p> <pre># Determine the size of data that was loaded dim(concrete)</pre> <p>e. Run cell</p> <pre>In [6]: # Determine the size of data that was loaded dim(concrete)</pre> <p>f. How many rows and columns are in the dataset? Rows: _____ Columns: _____</p> <p>Hint: You can use the help operator to learn about the dim() function output in order to answer this question. See Note in Step 8a.</p>

Step	Action
9	<p><u>Examine the descriptive statistics</u></p> <p>Descriptive statistics are used to describe or summarize features in a collection of data. These are broken down into some well-known components such as mean, median and standard deviation.</p> <p>a. Enter the code</p> <pre># Examine the descriptive statistics of the dataset con_desc <- describe(concrete)</pre> <p>b. Run cell</p> <pre>In [7]: # Examine the descriptive statistics of the dataset con_desc <- describe(concrete)</pre> <p>c. Enter the code</p> <pre>con_desc</pre> <p>d. Run cell</p> <pre>In [8]: con_desc</pre> <p>e. Write down the median for plast_amt: _____</p>
10	<p><u>Observe the distribution of each variable</u></p> <p>Histograms allow for the distribution of values to be seen very quickly. Values are counted in bins that consist of ranges of values. The taller the bar, the larger the count of values for the range is.</p> <p>a. Enter the code</p> <pre># Display histograms to show how the data is distributed multi.hist(concrete, bcol = "gray", dcol = c("blue", "red"), dlty = c("dotted", "solid"), main = c("Histogram, Density, Normal"))</pre> <p>b. Run cell</p>

Step	Action
	<pre>In [9]: # Display histograms to show how the data is distributed multi.hist(concrete, bcol = "gray", dcol = c("blue", "red"), dty = c("dotted", "solid"), main = c("Histogram, Density, Normal"))</pre>
11	<p><u>Determine the correlation value</u></p> <p>Correlation shows how two variables relate to each other. The value of the correlation represents a percentage of change that is related between the variables. If the correlation is positive, both variables move in the same direction. Negative correlation means the variables move in opposite directions. A correlation of 1 indicates both variables move the same amount together. If correlation is zero, there is no relationship in how the variables move together.</p> <ol style="list-style-type: none"> Enter the code <pre>concrete_cor <- cor(concrete) concrete_cor</pre> Run cell <pre>In [10]: concrete_cor <- cor(concrete) concrete_cor</pre> Write down the correlation for water_amt and plast_amt: _____ Enter code <pre>corrplot(concrete_cor, method="number", type="upper")</pre> Run cell <pre>In [11]: corrplot(concrete_cor, method="number", type="upper")</pre>
12	<p><u>Create training and testing datasets</u></p> <p>In this section, the model will have the ability to adaptively resample the tuning parameter in order to concentrate on values that will provide the optimal settings.</p> <p>First, the data will be split. 80% to train the model and 20% to test it.</p> <p>“One of the first decisions to make when modeling is to decide which samples will be used to evaluate performance. Ideally, the model should be evaluated on samples that were not used to build or fine-tune the model, so that they provide an unbiased sense of model effectiveness. When a large amount of data is at hand, a set of samples can be set aside to evaluate the final model. The “training” data set is the general term for</p>

Step	Action
	<p>the samples used to create the model, while the “test” or “validation” data set is used to qualify performance.” ⁽¹⁾</p> <p>In most cases, the training and test samples are desired to be as homogenous as possible. Random sampling methods can be used to create similar datasets.</p> <p>Example:</p> <p>“Assume that we need to estimate average number of votes for each candidate in an election. Assume that country has 3 towns: Town A has 1 million factory workers; Town B has 2 million office workers and Town C has 3 million retirees. We can choose to get a random sample of size 60 over entire population but there is some chance that the random sample turns out to be not well balanced across these towns and hence is biased causing a significant error in estimation. Instead if we choose to take a random sample of 10, 20 and 30 from Town A, B and C respectively then we can produce a smaller error in estimation for the same total size of sample.” ⁽²⁾</p> <p>a. Enter the code</p> <pre>trainIndex <- createDataPartition(concrete\$strength, p=0.8, list=FALSE,times=1)</pre> <p>b. Run cell</p> <pre>In [12]: trainIndex <- createDataPartition(concrete\$strength, p=0.8, list=FALSE,times=1)</pre> <p>The split data will be labeled, train and test.</p> <p>c. Enter the code</p> <pre>train <- concrete[trainIndex,] test <- concrete[-trainIndex,]</pre> <p>d. Run cell</p> <pre>In [13]: train <- concrete[trainIndex,] test <- concrete[-trainIndex,]</pre> <p>The data in train can be viewed.</p>

Step	Action
	<p>e. Enter the code</p> <pre>describe(train)</pre> <p>f. Run cell</p> <pre>In [14]: describe(train)</pre> <p>g. Write down the median for plast_amt _____</p> <p><u>Examine Training Dataset</u></p> <p>Note: We want to ensure that the splitting of the data did not result in different profiles for the training and testing data. The function used does its best to ensure the resulting datasets have a similar profile, but the best practice is to check.</p> <p>h. Enter the code</p> <pre>train_cor <- cor(train) train_cor</pre> <p>i. Run cell</p> <pre>In [15]: train_cor <- cor(train) train_cor</pre> <p>j. Write down the correlation for water_amt and plast_amt: _____</p> <p>Plot the correlations for train.</p> <p>k. Enter the code</p> <pre>corrplot(train_cor, method="number", type="upper")</pre> <p>l. Run cell</p> <pre>In [16]: corrplot(train_cor, method="number", type="upper")</pre> <p><u>Examine Testing Dataset</u></p> <p>The data in test can be viewed.</p>

Step	Action
	<p>m. Enter the code</p> <pre>describe(test)</pre> <p>n. Run cell</p> <pre>In [17]: describe(test)</pre> <p>o. Write down the median for plast_amt: _____</p> <p>Find the correlations for test.</p> <p>p. Enter the code</p> <pre>test_cor <- cor(test) test_cor</pre> <p>q. Run cell</p> <pre>In [18]: test_cor <- cor(test) test_cor</pre> <p>r. Write down the correlation for water_amt and plast_amt: _____</p> <p>Plot the correlations for test.</p> <p>s. Enter the code</p> <pre>corrplot(test_cor, method="number", type="upper")</pre> <p>t. Run cell</p> <pre>In [19]: corrplot(test_cor, method="number", type="upper")</pre>

Note: At this time, the training and test data is ready for Module 4: Approaches to Machine Learning.

Module 3: Lab Summary

Module 3 Lab started with IBM Data Science Experience (DSX). A project was created followed by an R notebook. A brief introduction to cell types was provided and a title was added to the notebook using Markdown. The required libraries, also called packages, were loaded. Then, a dataset was loaded from a CSV file stored on GitHub.

To ensure reproducibility, a seed value was set at beginning of the lab. A seed value is used for random number generation and utilized for the random selection process when the training and testing datasets are created.

The first few rows of the dataset were visually inspected along with the size. Descriptive statistics were leveraged to better understand the data. To describe the data, visualizations were implemented.

Variable relationships were measured to determine which were strongly correlated. Correlations range from 0 to 1, where 0 indicates there is no correlation and 1 means there is a strong correlation. Values between 0 and 1 mean there is some relationship between the two variables and high correlation generally starts at a value of 0.60.

The final step of this lab was to create the training and testing datasets which will be utilized in the Module 4 Lab.

References

- [1] Kuhn, M., & Johnson, K. (2013). Applied predictive modeling (pp. 389-400). New York: Springer.
- [2] Stratified Sampling. Wikipedia. https://en.wikipedia.org/wiki/Stratified_sampling


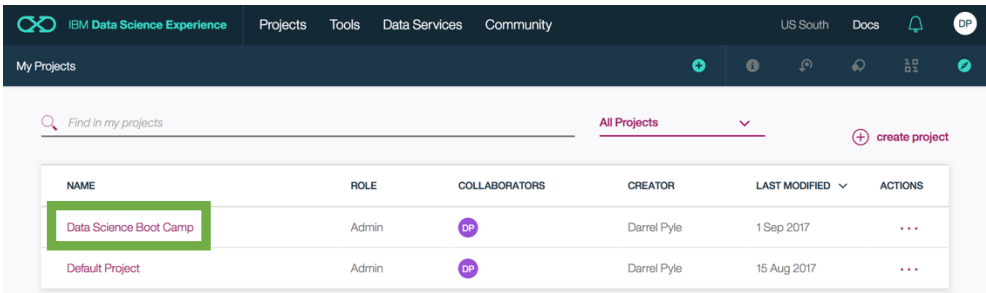
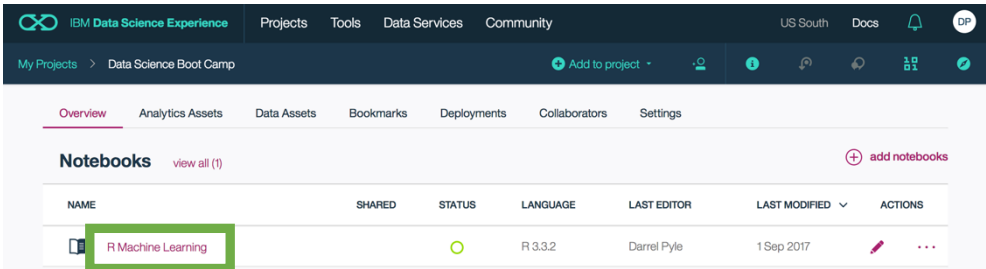
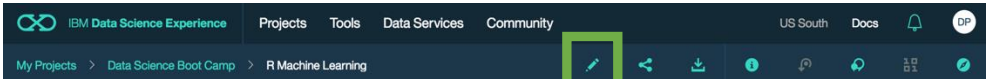
Module 4: Machine Learning Lab II

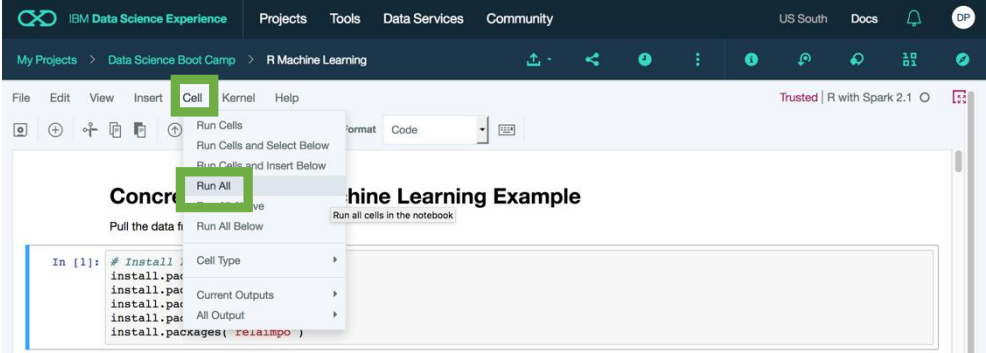

Purpose:	<p>This lab is a continuation of the Module 3 Lab. After completing the lab, you should be able to:</p> <ul style="list-style-type: none">• Utilize a training dataset to train models• Interpret and test the model• Evaluate model accuracy• Explain model results• Choose the best model
Tasks:	<p>Tasks you will complete in this lab exercise include:</p> <ul style="list-style-type: none">• Train multiple models• Test multiple models against a testing dataset• Compare the different models' performance

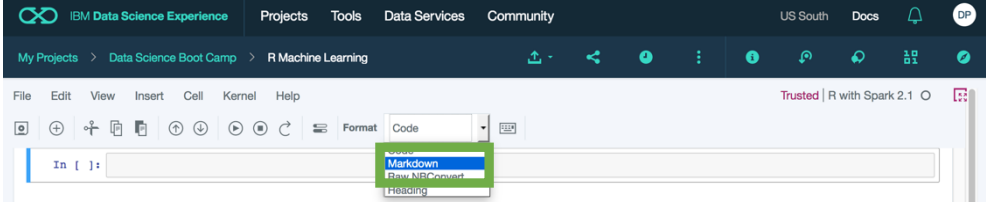


Module 4: Lab Workflow Overview

- 1 • Open and Run the Module 3 Notebook
- 2 • Create markdown title
- 3 • Create a multiple linear regression model
- 4 • View the generated coefficients
- 5 • Examine the confidence intervals
- 6 • Review residuals
- 7 • Review the Anova table
- 8 • Check covariance
- 9 • Examine model plots
- 10 • Check predictions
- 11 • Evaluate model accuracy
- 12 • Model using stepwise regression
- 13 • Check predictor importance
- 14 • Bootstrap measure of relative importance

Module 4: Lab Instructions

Step	Action
1	<p><u>Open and Run the Module 3 Notebook</u></p> <ol style="list-style-type: none"> Navigate to https://datascience.ibm.com/ Login to DSX In the top menu bar, click Projects and select View All Projects  <ol style="list-style-type: none"> Click Data Science Boot Camp  <ol style="list-style-type: none"> Click R Machine Learning to open the notebook that was created in Module 3  <ol style="list-style-type: none"> Click the edit (pencil) icon in the toolbar 

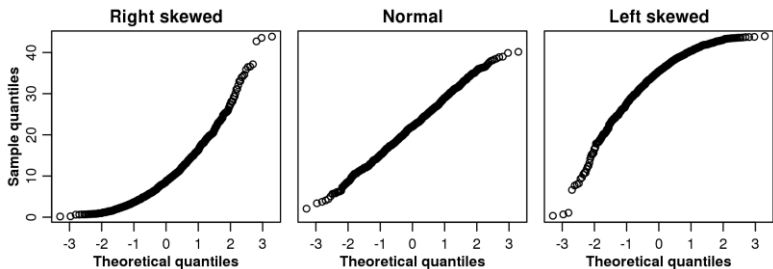
Step	Action
	<p data-bbox="370 310 1110 344">g. From the menu bar, select Cell and click Run All</p>  <p data-bbox="310 772 1461 991">By running all of the cells, the referenced libraries are installed and loaded into memory to be available for usage. If the libraries have already been installed, typically the rows with the <code>install.packages</code> statements would be commented out by placing <code>#</code> in front of each line or the rows would be deleted. This would reduce the script execution time. Below is an example of commenting out the <code>install.packages</code> lines.</p> <pre data-bbox="399 1031 1372 1136">In [1]: # Install Libraries # install.packages("corrplot") # install.packages("psych") # install.packages("caret") # install.packages("MASS") # install.packages("relaimpo")</pre> <p data-bbox="310 1173 1461 1318">Data is also loaded into the data frames and are made available to the script. All the remaining commands are also executed. This places the notebook at the same point as it was when Module 3 Lab was completed and it is ready to have additional script added to it.</p>
2	<p data-bbox="310 1360 643 1394"><u>Create markdown title</u></p> <p data-bbox="310 1432 1461 1503">Markdown will be used to place a title to separate the Module 4 Lab code from the code of the previous exercise.</p> <p data-bbox="370 1541 1352 1575">a. Scroll to the bottom of the notebook and click in the last empty cell</p> 

Step	Action
	<p>b. Change the cell to Markdown in the format dropdown in the toolbar</p>  <p>c. Type in the cell:</p> <pre> *** *** # Start of Module 4 ### Several models will be created and compared within this module. </pre> <p>d. Run the cell</p>  <p>e. What do the “***” characters create? _____</p>
3	<p><u>Create a multiple linear regression model</u></p> <p>Multiple Linear Regression determines the relationship between a dependent (target) variable and the remaining independent (attribute) variables. The relationship of a linear regression equation assumes that each independent variable affects the dependent variable in a linear and additive manner.</p> <p>a. Type in the cell:</p> <pre> # Use R to create a linear regression model fit <- lm(strength ~ age + fine_agg_amt + course_agg_amt + plast_amt + water_amt + ash_amt + slag_amt + cement_amt, data = train) </pre> <p>b. Run the cell</p> 

Step	Action
	<p>Strength (located before ~) is the dependent variable. The remaining variables (located after ~) are the independent variables. The multiple linear regression model is stored as a variable named fit.</p>
4	<p><u>View the generated coefficients</u></p> <p>Coefficients indicate the amount of impact an independent variable has on the dependent variable, the larger the coefficient, the greater the influence an independent variable has on the dependent variable. The sign of the coefficient indicates if the relationship of the independent variable to the dependent variable is negative or positive. The intercept indicates a base starting point for the dependent variable when the independent variables are not present.</p> <ol style="list-style-type: none"> Type in the cell: <pre># Examine the generated coefficients coefficients(fit)</pre> Run the cell <pre>In [21]: # Examine the generated coefficients coefficients(fit)</pre> The coefficient of course_agg_amt is: _____ Does course_agg_amt have a positive or negative relationship to strength? _____ Does this indicate that course_agg_amt's effect increases or decreases strength? _____
5	<p><u>Examine the confidence intervals</u></p> <p>A confidence interval indicates the range of values that may contain the mean of the variable being looked at for a given probability (level). We will look at the confidence intervals that have a 95% probability (specified with "level=" parameter) of containing each variable.</p> <ol style="list-style-type: none"> Enter the code: <pre># Examine the confidence intervals of the model confint(fit, level=0.95)</pre> Run cell

Step	Action
	<div data-bbox="492 275 1458 321" data-label="Code-Block"> <pre>In [22]: # Examine the confidence intervals of the model confint(fit, level=0.95)</pre> </div> <p data-bbox="370 359 1039 394">c. The 95% confidence interval for ash_amt is:</p> <p data-bbox="418 436 1029 472">_____ to _____</p> <p data-bbox="418 506 1446 579">The confidence interval means there is a 95% chance that the calculated interval contains the true mean (average) of ash_amt.</p>
6	<p data-bbox="310 621 570 657"><u>Review residuals</u></p> <p data-bbox="310 690 1458 909">In multiple linear regression, a residual is the vertical distance that a data point is from the line that is generated from the regression equation. The linear regression equation estimates the value of the dependent variable. The estimate is compared against the known data point to return the residual value. The closer the residual value is to zero, the closer the linear regression equation is to estimating the actual dependent value (i.e. the more accurate the model is).</p> <p data-bbox="578 947 1198 982" style="text-align: center;">residual = known value – predicted value</p> <p data-bbox="310 1020 1446 1129">When the residual value is negative, it indicates the predicted value is too large which is known as an overestimate. Underestimated predicted values result in a positive residual value.</p> <p data-bbox="310 1167 1406 1241">For this example, the residual of each data point is the difference between the regression value of strength and the known strength value for each data point.</p> <p data-bbox="370 1278 634 1314">a. Enter the code:</p> <div data-bbox="483 1318 805 1398" data-label="Code-Block"> <pre># Review residuals head(residuals(fit), 15)</pre> </div> <p data-bbox="370 1436 537 1472">b. Run cell</p> <div data-bbox="475 1476 1446 1522" data-label="Code-Block"> <pre>In [23]: # Review residuals head(residuals(fit), 15)</pre> </div> <p data-bbox="370 1560 1182 1596">c. Which data point has the smallest residual? _____</p> <p data-bbox="370 1633 1187 1669">d. What is the residual value of the data point? _____</p> <p data-bbox="370 1707 1422 1780">e. Did the regression equation overestimate or underestimate the strength value? _____</p>

Step	Action
7	<p><u>Review the Anova table</u></p> <p>An Anova table is another way to measure which variables have a significant impact on the dependent variable. To determine if a variable is significant, the p-value column is assessed and compared to the level of significance being used. Typically, a 5% level of significance is used which means the model explains 95% of the data. The p-value is compared to the level of significance (0.05) and if it is less than the level of significance, the variable is determined to have an impact on the dependent variable.</p> <p>The p-value for each variable is located in the “Pr(>F)” column corresponding to each variable in the Anova table.</p> <ol style="list-style-type: none"> Enter the code <pre># Review the anova table anova(fit)</pre> Run cell <pre>In [24]: # Review the anova table anova(fit)</pre> Which variable has the most significant impact (smallest p-value) on strength? _____
8	<p><u>Check covariance</u></p> <p>Covariance provides an indication of whether two variables increase or decrease together. If two variables increase or decrease together, the covariance will be positive. If one variable decreases while the other variable increases, then the covariance value will be negative. The magnitude of the covariance value indicates how far apart the values are from the mean; however, if the variables being compared are of different scales (such as feet compared to inches) the magnitude can be misleading. Typically, only the sign of covariance is used and the magnitude is ignored.</p> <ol style="list-style-type: none"> Enter the code <pre># Covariance matrix for model parameters vcov(fit)</pre>

Step	Action
	<p>b. Run cell</p> <pre data-bbox="487 384 1459 430">In [25]: # Covariance matrix for model parameters vcov(fit)</pre>
9	<p><u>Examine model plots</u></p> <p>The plots that will be examined are:</p> <p><u>Residuals vs Fitted</u>: Detect non-linearity, unequal error variance, and outliers. <u>Scale-location</u>: Shows equally spread residuals along the ranges of predictors <u>Normal Q-Q</u>: Check if residuals are normally distributed <u>Residuals vs Leverage</u>: Helps find influential data points</p> <div data-bbox="500 793 1268 1060">  </div> <p>The figure above is a set of Q-Q plots with various data distributions. Right skewed data means the majority of data has small values with few large values. When the majority of data has large values with only a few small values, the data is left skewed. Normal data has most of its values surrounding the mean with decreasing amounts both smaller and larger than the mean. The smaller or larger the values, the fewer values there are, overall the data value count follow the bell curve. The Q-Q plotting the residual and quantity of the residuals. When a Q-Q plot is normal, illustrated by the middle graph above, it means the model predicts values higher and lower than the actual value with equal frequency. If the Q-Q plot is not normal, then the model is not accounting for everything that it needs to</p> <p>a. Enter the code</p> <pre data-bbox="438 1581 1203 1661">layout(matrix(c(1,2,3,4),2,2)) # optional 4 graphs/page plot(fit)</pre> <p>b. Run the cell</p> <pre data-bbox="438 1749 1411 1795">In [26]: layout(matrix(c(1,2,3,4),2,2)) # optional 4 graphs/page plot(fit)</pre>

Step	Action
	<p>c. The Normal Q-Q plot shows if residuals are normally distributed. This is indicated if the data points line up along the diagonal dotted line in the plot. Do the residuals appear to be normally distributed? _____</p>
10	<p><u>Check predictions</u></p> <p>The model can be applied to a testing dataset to calculate predicted values. The predicted values can then be compared against the actual values.</p> <p>a. Enter the code</p> <pre># Use the model to predict the results when the linear regression is # applied to the test data # View the first 5 rows to verify there are some results conc_pred <- predict(fit,newdata=test, fit=TRUE) head(conc_pred)</pre> <p>b. Run the cell</p> <pre>In [27]: # Use the model to predict the results when the linear regression is applied to the test data # View the first 5 rows to verify there are some results conc_pred <- predict(fit,newdata=test, fit=TRUE) head(conc_pred)</pre> <p>c. What is the value predicted for row ID 10? _____</p> <p>d. Enter the code</p> <pre># Compare the predicted value to the test (actual) value test[1,]</pre> <p>e. Run the cell</p> <pre>In [28]: # Compare the predicted value to the test (actual) value test[1,]</pre> <p>f. What is the actual value of strength for row ID 10? _____</p> <p>g. What is the percent difference between the predicted and actual value? _____</p>

Step	Action
11	<p><u>Evaluate model accuracy</u></p> <p>The value of R-Squared can be used to get a feel for the accuracy of predictions. The value ranges from 0% to 100% expressed as a decimal value. Generally, the higher the value the more accurate the model is.</p> <p>a. Enter the code</p> <pre># Examine the accuracy of the results postResample(pred = conc_pred, obs = test\$strength)</pre> <p>b. Run the cell</p> <pre>In [28]: # Examine the accuracy of the results postResample(pred = conc_pred, obs = test\$strength)</pre> <p>c. What percentage of the data is explained by the model? _____</p>
12	<p><u>Model using stepwise regression</u></p> <p>Another modelling technique uses stepwise selection. Stepwise regression can function in one of three methods:</p> <p><u>Forward Selection</u>: starts with no variables and adds in variables until model improvements cannot be made</p> <p><u>Backward Elimination</u>: starts with all variables and removes variables until model improvements cannot be made</p> <p><u>Bidirectional Elimination</u>: combines both forward and backward methods until model improvements cannot be made</p> <p>The overall goal of stepwise regression is to optimize which variables are included in a model. For the lab, bidirectional elimination will be used.</p> <p>a. Enter the code</p> <pre># Stepwise Regression step <- stepAIC(fit, direction="both")</pre>

Step	Action
	<p><code>step\$anova # display results</code></p> <p>b. Run the cell</p> <pre>In [29]: # Stepwise Regression step <- stepAIC(fit, direction="both") step\$anova # display results</pre> <p>Next, examine the stepwise model.</p> <p>c. Enter the code</p> <p><code>step</code></p> <p>d. Run the cell</p> <pre>In [30]: step</pre> <p>e. The coefficients from the linear regression were:</p> <pre>(Intercept) -6.69842457185358 age 0.111665479837888 fine_agg_amt 0.0150081212898075 course_agg_amt 0.0136261119756282 plast_amt 0.297816548355918 water_amt -0.178034408123492 ash_amt 0.0775641746632639 slag_amt 0.0963375880785001 cement_amt 0.112662766607026</pre> <p>Does the stepwise regression model have different coefficients?</p> <p>_____</p> <p>If the coefficients are the same, that can be an indication that all variables are adding value to the model and removing any particular variable will reduce the accuracy of the model.</p>
13	<p><u>Check predictor importance</u></p> <p>Predictor importance can change based on when a variable is added to a model. Analyze how adding variables at different times affects the model. Four types of predictor importance will be examined:</p> <p><u>LMG</u>: utilizes R squared</p>

Step	Action
	<p><u>Last</u>: each variables contribution when included last <u>First</u>: each variables contribution when included first <u>Pratt</u>: utilizes the product of the standardized coefficient and the correlation</p> <p>a. Enter the code</p> <pre># Calculate Relative Importance for Each Predictor calc.relimp(fit,type=c("lmg","last","first","pratt"), rela=TRUE)</pre> <p>b. Run the cell</p> <pre>In [32]: # Calculate Relative Importance for Each Predictor calc.relimp(fit,type=c("lmg","last","first","pratt"), rela=TRUE)</pre> <p>c. Which type of predictor importance has the variable with the largest relative importance metric? _____</p> <p>d. Which variable has the largest importance? _____</p> <p>e. What is the value of the importance? _____</p>
14	<p><u>Bootstrap measure of relative importance</u></p> <p>Bootstrapping is any test or metric that relies on random sampling with replacement. Random sampling with replacement means that after each sample is taken, the data point is placed back into the dataset and is available once again for the random sample process. Random sampling can also be performed without replacement. In this scenario, when a sample is taken, it is removed from the dataset and not available for further sampling. For this lab, the “with replacement” option is being used. Specifying a parameter for the number of bootstrap runs (b) to execute allows for greater validation and potentially more accuracy.</p> <p>a. Enter the code</p> <pre># Bootstrap Measures of Relative Importance (100 samples) boot <- boot.relimp(fit, b = 100, diff = TRUE, rela = TRUE, rank = TRUE, type = c("lmg", "last", "first", "pratt")) booteval.relimp(boot) # print result</pre>

Step	Action
	<p>b. Run the cell</p> <pre data-bbox="391 310 1365 390">In [33]: # Bootstrap Measures of Relative Importance (100 samples) boot <- boot.relimp(fit, b = 100, diff = TRUE, rela = TRUE, rank = TRUE, type = c("lmg", "last", "first", "pratt")) booteval.relimp(boot) # print result</pre> <p>c. This produced similar output to the previous step to check the predictor relative importance. Is there anything that appears to be considerably different by using the bootstrap method? _____</p> <p>To make it easier to identify the most important predictors, plots can be used.</p> <p>d. Enter the code</p> <pre data-bbox="488 726 1203 762">plot(booteval.relimp(boot,sort=TRUE)) # plot result</pre> <p>e. Run the cell</p> <pre data-bbox="488 846 1459 884">In [34]: plot(booteval.relimp(boot,sort=TRUE)) # plot result</pre> <p>f. Which predictor is the most important for each predictor type? LMG: _____ Last: _____ First: _____ Pratt: _____</p>

Module 4: Lab Summary

At the end of Module 3's Lab, training and testing datasets were created and used for this lab. Training data is used to create and train various machine learning models. Testing data is used to validate a trained model. It contains the actual data values for the dependent variable (strength) in addition to all of the independent variables.

The model was run against the independent variables (inputs) and calculated a value for the dependent variable (output). Once complete, the calculated value was compared against the actual value. The closer the calculated value was to the true value, the more accurate the model. These concepts were supported by many of the activities in the Module 3 Lab.

A multiple linear regression model was created. With this kind of model, the data typically appears to have a linear relationship. In this case, each independent variable was multiplied by a coefficient and all the values were added together to arrive at a value for the dependent variable.

For this model, here is the specific equation generated (shown with rounded coefficients):

$$\begin{aligned} \text{strength} = & -6.6984 + 0.1117 \cdot \text{age} + 0.0150 \cdot \text{fine_agg_amt} + 0.0136 \cdot \text{course_agg_amt} \\ & + 0.2978 \cdot \text{plast_amt} - 0.1780 \cdot \text{water_amt} + 0.0776 \cdot \text{ash_amt} + 0.0963 \cdot \text{slag_amt} \\ & + 0.1127 \cdot \text{cement_amt} \end{aligned}$$

Finally, model accuracy and performance were measured using confidence intervals, residuals, Anova, covariance, and bootstrapping.