

# conclusions\_query

October 18, 2017

## 1 Drawing Conclusions Using Query

```
In [1]: # Load `winequality_edited.csv`  
import pandas as pd
```

```
df = pd.read_csv('winequality_edited.csv')  
df.head()
```

```
Out[1]:
```

	fixed_acidity	volatile_acidity	citric_acid	residual_sugar	chlorides	\
0	7.4	0.70	0.00	1.9	0.076	
1	7.8	0.88	0.00	2.6	0.098	
2	7.8	0.76	0.04	2.3	0.092	
3	11.2	0.28	0.56	1.9	0.075	
4	7.4	0.70	0.00	1.9	0.076	

  

	free_sulfur_dioxide	total_sulfur_dioxide	density	pH	sulphates	\
0	11.0	34.0	0.9978	3.51	0.56	
1	25.0	67.0	0.9968	3.20	0.68	
2	15.0	54.0	0.9970	3.26	0.65	
3	17.0	60.0	0.9980	3.16	0.58	
4	11.0	34.0	0.9978	3.51	0.56	

  

	alcohol	quality	color	acidity_levels
0	9.4	5	RED	low
1	9.8	5	RED	med-high
2	9.8	5	RED	med-low
3	9.8	6	RED	med-high
4	9.4	5	RED	low

### 1.0.1 Do wines with higher alcoholic content receive better ratings?

```
In [2]: # get the median amount of alcohol content  
alc_median = df["alcohol"].median()
```

```
In [3]: # select samples with alcohol content less than the median  
low_alcohol = df.query('alcohol < @alc_median')
```

```
# select samples with alcohol content greater than or equal to the median
high_alcohol = df.query('alcohol >= @alc_median')
```

```
# ensure these queries included each sample exactly once
```

```
num_samples = df.shape[0]
```

```
num_samples == low_alcohol['quality'].count() + high_alcohol['quality'].count() # should be
```

```
Out[3]: True
```

```
In [5]: # get mean quality rating for the low alcohol and high alcohol groups
```

```
qlty_mean_low_alcohol = low_alcohol["quality"].mean()
```

```
qlty_mean_high_alcohol = high_alcohol["quality"].mean()
```

```
print("mean quality, low alcohol: {}".format(qlty_mean_low_alcohol))
```

```
print("mean quality, high alcohol: {}".format(qlty_mean_high_alcohol))
```

```
mean quality, low alcohol: 5.475920679886686
```

```
mean quality, high alcohol: 6.146084337349397
```

## 1.0.2 Do sweeter wines receive better ratings?

```
In [6]: # get the median amount of residual sugar
```

```
resid_sugar_median = df["residual_sugar"].median()
```

```
print(resid_sugar_median)
```

```
3.0
```

```
In [7]: # select samples with residual sugar less than the median
```

```
low_sugar = df.query('residual_sugar < @resid_sugar_median')
```

```
# select samples with residual sugar greater than or equal to the median
```

```
high_sugar = df.query('residual_sugar >= @resid_sugar_median')
```

```
# ensure these queries included each sample exactly once
```

```
num_samples == low_sugar['quality'].count() + high_sugar['quality'].count() # should be
```

```
Out[7]: True
```

```
In [8]: # get mean quality rating for the low sugar and high sugar groups
```

```
qlty_mean_low_sugar = low_sugar["quality"].mean()
```

```
qlty_mean_high_sugar = high_sugar["quality"].mean()
```

```
print("mean quality, low sugar: {}".format(qlty_mean_low_sugar))
```

```
print("mean quality, high sugar: {}".format(qlty_mean_high_sugar))
```

```
mean quality, low sugar: 5.808800743724822
```

```
mean quality, high sugar: 5.82782874617737
```

```
In [ ]:
```