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# 1 ANOVA - Lab

## 1.1 Introduction

In this lab, you'll get some brief practice generating an ANOVA table (AOV) and interpreting its output. You'll also perform some investigations to compare the method to the t-tests you previously employed to conduct hypothesis testing.

# 1.2 Objectives

In this lab you will:

- Use ANOVA for testing multiple pairwise comparisons
- Interpret results of an ANOVA and compare them to a t-test

#### 1.3 Load the data

Start by loading in the data stored in the file 'ToothGrowth.csv':

```
[1]: # Your code here
import pandas as pd
import numpy as np

df = pd.read_csv('ToothGrowth.csv')
```

#### 1.4 Generate the ANOVA table

Now generate an ANOVA table in order to analyze the influence of the medication and dosage:

```
[2]: # Your code here
import statsmodels.api as sm
from statsmodels.formula.api import ols

formula = 'len ~ C(supp) + C(dose)'
lm = ols(formula, df).fit()
table = sm.stats.anova_lm(lm, typ=2)
print(table)
```

```
sum_sq df F PR(>F)
C(supp) 205.350000 1.0 14.016638 4.292793e-04
```

```
C(dose) 2426.434333 2.0 82.810935 1.871163e-17
Residual 820.425000 56.0 NaN NaN
```

# 1.5 Interpret the output

Make a brief comment regarding the statistics and the effect of supplement and dosage on tooth length:

```
[3]: # Your comment here

# As we can see, All of values in the far right column are significantly
# smaller than 0.05, so we can reject the null hypothesis.

# Both dose and supplement type are impactful. At first glance, dosage seems
# to be the more impactful of the two.
```

# 1.6 Compare to t-tests

Now that you've had a chance to generate an ANOVA table, its interesting to compare the results to those from the t-tests you were working with earlier. With that, start by breaking the data into two samples: those given the OJ supplement, and those given the VC supplement. Afterward, you'll conduct a t-test to compare the tooth length of these two different samples:

```
[4]: df.head()
[4]:
         len supp
                   dose
     0
         4.2
               VC
                     0.5
     1
        11.5
               VC
                     0.5
         7.3
     2
               VC
                     0.5
     3
         5.8
               VC
                     0.5
         6.4
               VC
                     0.5
[5]: # Your code here
     import scipy.stats as stats
     VC = df[df["supp"] == "VC"]["len"]
     OJ = df[df["supp"] == "OJ"]["len"]
     print(len(VC))
     print(VC.std())
     print("\n")
     print(len(OJ))
     print(OJ.std())
```

30

8.266028664664638

30

#### 6.605561049722362

Now run a t-test between these two groups and print the associated two-sided p-value:

```
[6]: # Calculate the 2-sided p-value for a t-test comparing the two supplement groups
results = stats.ttest_ind(VC, OJ, equal_var=False)
results.pvalue
```

[6]: 0.06063450788093387

# 1.7 A 2-Category ANOVA F-test is equivalent to a 2-tailed t-test!

Now, recalculate an ANOVA F-test with only the supplement variable. An ANOVA F-test between two categories is the same as performing a 2-tailed t-test! So, the p-value in the table should be identical to your calculation above.

Note: there may be a small fractional difference (>0.001) between the two values due to a rounding error between implementations.

```
[7]: # Your code here; conduct an ANOVA F-test of the oj and vc supplement groups.

formula = 'len ~ C(supp)'
lm = ols(formula, df).fit()
table = sm.stats.anova_lm(lm, typ=2)
print(table)

# Compare the p-value to that of the t-test above.
# They should match (there may be a tiny fractional difference due to # rounding errors in varying implementations)
```

```
sum_sq df F PR(>F)
C(supp) 205.350000 1.0 3.668253 0.060393
Residual 3246.859333 58.0 NaN NaN
```

### 1.8 Run multiple t-tests

While the 2-category ANOVA test is identical to a 2-tailed t-test, performing multiple t-tests leads to the multiple comparisons problem. To investigate this, look at the various sample groups you could create from the 2 features:

```
[9]: for group in df.groupby(['supp', 'dose'])['len']:
    group_name = group[0]
    data = group[1]
    print(group_name)
```

```
('OJ', 0.5)
('OJ', 1.0)
('OJ', 2.0)
```

```
('VC', 0.5)
      ('VC', 1.0)
      ('VC', 2.0)
[39]: | 1 = list(df.groupby(['supp', 'dose'])['len'])
      pd.DataFrame(1)
[39]:
                   0
                                                                              1
          (0J, 0.5)
                       30
                              15.2
      31
             21.5
      32
             17.6
      33
               9.7
      34...
          (0J, 1.0)
                              19.7
                       40
      41
             23.3
      42
             23.6
      43
             26.4
      44...
          (0J, 2.0)
                              25.5
                       50
      51
             26.4
      52
             22.4
      53
             24.5
      54...
          (VC, 0.5)
                              4.2
      3
                       0
      1
            11.5
      2
             7.3
      3
             5.8
      4
          (VC, 1.0)
                       10
                              16.5
      4
      11
             16.5
      12
             15.2
      13
             17.3
      14...
          (VC, 2.0)
                              23.6
      5
                       20
      21
             18.5
      22
             33.9
      23
             25.5
      24...
```

While bad practice, examine the effects of calculating multiple t-tests with the various combinations of these. To do this, generate all combinations of the above groups. For each pairwise combination, calculate the p-value of a 2-sided t-test. Print the group combinations and their associated p-value for the two-sided t-test.

```
[12]: # Your code here; reuse your t-test code above to calculate the p-value for # a 2-sided t-test # for all combinations of the supplement-dose groups listed above. # (Since there isn't a control group, compare each group to every other group.)
```

```
### From GitHub Solution
from itertools import combinations

groups = [group[0] for group in df.groupby(['supp', 'dose'])['len']]
combos = combinations(groups, 2)
for combo in combos:
    supp1 = combo[0][0]
    dose1 = combo[0][1]
    supp2 = combo[1][0]
    dose2 = combo[1][0]
    dose2 = combo[1][1]
    sample1 = df[(df.supp == supp1) & (df.dose == dose1)]['len']
    sample2 = df[(df.supp == supp2) & (df.dose == dose2)]['len']
    p = stats.ttest_ind(sample1, sample2, equal_var=False)[1]
    print(combo, p)

# Note that while ANOVA also concluded that all factors were significant,
# these p-values are substantially lower.
```

```
(('OJ', 0.5), ('OJ', 1.0)) 8.784919055161479e-05

(('OJ', 0.5), ('OJ', 2.0)) 1.3237838776972294e-06

(('OJ', 0.5), ('VC', 0.5)) 0.006358606764096813

(('OJ', 0.5), ('VC', 1.0)) 0.04601033257637553

(('OJ', 0.5), ('VC', 2.0)) 7.196253524006043e-06

(('OJ', 1.0), ('OJ', 2.0)) 0.039195142046244004

(('OJ', 1.0), ('VC', 0.5)) 3.6552067303259103e-08

(('OJ', 1.0), ('VC', 1.0)) 0.001038375872299884

(('OJ', 1.0), ('VC', 2.0)) 0.09652612338267014

(('OJ', 2.0), ('VC', 0.5)) 1.3621396478988818e-11

(('OJ', 2.0), ('VC', 0.5)) 1.3621396478988818e-07

(('OJ', 2.0), ('VC', 1.0)) 2.3610742020468435e-07

(('OJ', 2.0), ('VC', 1.0)) 6.811017702865016e-07

(('VC', 0.5), ('VC', 2.0)) 4.6815774144921145e-08

(('VC', 1.0), ('VC', 2.0)) 9.155603056638692e-05
```

# 1.9 Summary

In this lesson, you implemented the ANOVA technique to generalize testing methods to multiple groups and factors.