Analyze ab test results notebook

March 30, 2021

0.1 Analyze A/B Test Results

You may either submit your notebook through the workspace here, or you may work from your local machine and submit through the next page. Either way assure that your code passes the project RUBRIC. Please save regularly.

This project will assure you have mastered the subjects covered in the statistics lessons. The hope is to have this project be as comprehensive of these topics as possible. Good luck!

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Introduction

A/B tests are very commonly performed by data analysts and data scientists. It is important that you get some practice working with the difficulties of these

For this project, you will be working to understand the results of an A/B test run by an e-commerce website. Your goal is to work through this notebook to help the company understand if they should implement the new page, keep the old page, or perhaps run the experiment longer to make their decision.

As you work through this notebook, follow along in the classroom and answer the corresponding quiz questions associated with each question. The labels for each classroom concept are provided for each question. This will assure you are on the right track as you work through the project, and you can feel more confident in your final submission meeting the criteria. As a final check, assure you meet all the criteria on the RUBRIC.

Part I - Probability

To get started, let's import our libraries.

```
[1]: import pandas as pd
import numpy as np
import random
import matplotlib.pyplot as plt
%matplotlib inline
```

```
#We are setting the seed to assure you get the same answers on quizzes as we_set up
random.seed(42)
```

- 1. Now, read in the ab_data.csv data. Store it in df. Use your dataframe to answer the questions in Quiz 1 of the classroom.
 - a. Read in the dataset and take a look at the top few rows here:

```
[2]: df = pd.read_csv("ab_data.csv")
    df.head()
```

```
[2]:
        user id
                                  timestamp
                                                  group landing_page
                                                                      converted
         851104
                 2017-01-21 22:11:48.556739
                                                control
                                                            old_page
                 2017-01-12 08:01:45.159739
     1
         804228
                                                control
                                                            old_page
                                                                               0
     2
         661590 2017-01-11 16:55:06.154213 treatment
                                                            new_page
                                                                               0
         853541 2017-01-08 18:28:03.143765 treatment
     3
                                                            new_page
                                                                               0
         864975 2017-01-21 01:52:26.210827
                                                control
                                                            old_page
                                                                               1
```

- b. Use the cell below to find the number of rows in the dataset.
- [3]: df.shape
- [3]: (294478, 5)
 - c. The number of unique users in the dataset.
- [4]: df.user_id.nunique()
- [4]: 290584
 - d. The proportion of users converted.
- [5]: df.query('converted == 1').shape[0] / df.shape[0]
- [5]: 0.11965919355605512
- [6]: df.query('converted == 1').user_id.nunique() / df.user_id.nunique()
- [6]: 0.12104245244060237
 - e. The number of times the new_page and treatment don't match.
- [7]: df.query('group == "treatment" and landing_page != "new_page"').shape[0] + df.

 →query('group == "control" and landing_page == "new_page"').shape[0]
- [7]: 3893
 - f. Do any of the rows have missing values?
- [8]: df.info()

- 2. For the rows where **treatment** does not match with **new_page** or **control** does not match with **old_page**, we cannot be sure if this row truly received the new or old page. Use **Quiz 2** in the classroom to figure out how we should handle these rows.
 - a. Now use the answer to the quiz to create a new dataset that meets the specifications from the quiz. Store your new dataframe in df2.

```
[9]: df_tempa = df.query('group == "treatment" and landing_page == "new_page"')
    df_tempb = df.query('group == "control" and landing_page == "old_page"')
    df2 = pd.DataFrame()
    df2 = df2.append([df_tempa, df_tempb])
    df2.shape
```

[9]: (290585, 5)

```
[10]: # Double Check all of the correct rows were removed - this should be 0

df2[((df2['group'] == 'treatment') == (df2['landing_page'] == 'new_page')) == 

→False].shape[0]
```

- [10]: 0
 - 3. Use df2 and the cells below to answer questions for Quiz3 in the classroom.
 - a. How many unique **user** ids are in df2?

```
[11]: df2.user_id.nunique()
```

- [11]: 290584
 - b. There is one **user_id** repeated in **df2**. What is it?

```
[12]: df2[df2['user_id'].duplicated()].user_id
```

[12]: 2893 773192 Name: user_id, dtype: int64

c. What is the row information for the repeat **user** id?

```
[13]: df2[df2['user_id'].duplicated(keep=False)]
```

```
[13]:
            user_id
                                        timestamp
                                                        group landing_page
                                                                             converted
                      2017-01-09 05:37:58.781806
      1899
             773192
                                                   treatment
                                                                  new_page
                                                                                     0
      2893
             773192
                      2017-01-14 02:55:59.590927
                                                                                     0
                                                   treatment
                                                                  new_page
```

d. Remove **one** of the rows with a duplicate **user** id, but keep your dataframe as df2.

```
[14]: df2.drop_duplicates(subset='user_id', inplace=True) df2.shape
```

- [14]: (290584, 5)
 - 4. Use df2 in the cells below to answer the quiz questions related to Quiz 4 in the classroom.
 - a. What is the probability of an individual converting regardless of the page they receive?

```
[15]: df2.converted.mean()
```

- [15]: 0.11959708724499628
 - b. Given that an individual was in the control group, what is the probability they converted?

```
[16]: df2.query('group == "control"').converted.mean()
```

- [16]: 0.1203863045004612
 - c. Given that an individual was in the treatment group, what is the probability they converted?

```
[17]: df2.query('group == "treatment"').converted.mean()
```

- [17]: 0.11880806551510564
 - d. What is the probability that an individual received the new page?

```
[18]: df2.query('landing_page == "new_page"').shape[0] / df2.shape[0]
```

- [18]: 0.5000619442226688
 - e. Consider your results from parts (a) through (d) above, and explain below whether you think there is sufficient evidence to conclude that the new treatment page leads to more conversions.

It appears that the new page's conversion is no better than the old page's conversion. Therefore at this point there is no sufficient evidence to reject the old page (null hypothesis).

```
### Part II - A/B Test
```

Notice that because of the time stamp associated with each event, you could technically run a hypothesis test continuously as each observation was observed.

However, then the hard question is do you stop as soon as one page is considered significantly better than another or does it need to happen consistently for a certain amount of time? How long do you run to render a decision that neither page is better than another?

These questions are the difficult parts associated with A/B tests in general.

1. For now, consider you need to make the decision just based on all the data provided. If you want to assume that the old page is better unless the new page proves to be definitely better at a Type I error rate of 5%, what should your null and alternative hypotheses be? You can state your hypothesis in terms of words or in terms of p_{old} and p_{new} , which are the converted rates for the old and new pages.

$$H_0: p_{new} - p_{old} \le 0$$
$$H_1: p_{new} - p_{old} > 0$$

2. Assume under the null hypothesis, p_{new} and p_{old} both have "true" success rates equal to the **converted** success rate regardless of page - that is p_{new} and p_{old} are equal. Furthermore, assume they are equal to the **converted** rate in **ab_data.csv** regardless of the page.

Use a sample size for each page equal to the ones in ab data.csv.

Perform the sampling distribution for the difference in **converted** between the two pages over 10,000 iterations of calculating an estimate from the null.

Use the cells below to provide the necessary parts of this simulation. If this doesn't make complete sense right now, don't worry - you are going to work through the problems below to complete this problem. You can use **Quiz 5** in the classroom to make sure you are on the right track.

- [19]: df2.head(1)
- [19]: user_id timestamp group landing_page converted 2 661590 2017-01-11 16:55:06.154213 treatment new_page 0
 - a. What is the **conversion rate** for p_{new} under the null?
- [20]: df2.converted.mean()
- [20]: 0.11959708724499628
 - b. What is the **conversion rate** for p_{old} under the null?
- [21]: df2.converted.mean()
- [21]: 0.11959708724499628
 - c. What is n_{new} , the number of individuals in the treatment group?
- [22]: n_new = df2.query('group == "treatment"').shape[0]
 n_new
- [22]: 145310
 - d. What is n_{old} , the number of individuals in the control group?
- [23]: n_old = df2.query('group == "control"').shape[0]
 n_old

- [23]: 145274
 - e. Simulate n_{new} transactions with a conversion rate of p_{new} under the null. Store these n_{new} 1's and 0's in new_page_converted.

- [24]: 0.12054228889959397
 - f. Simulate n_{old} transactions with a conversion rate of p_{old} under the null. Store these n_{old} 1's and 0's in old_page_converted.

```
[25]: old_page_converted = np.random.choice([0, 1], n_old, p=[1-df2.converted.mean(), u →df2.converted.mean()]) old_page_converted.mean()
```

- [25]: 0.11931935514957942
 - g. Find p_{new} p_{old} for your simulated values from part (e) and (f).
- [26]: new_page_converted.mean() old_page_converted.mean()
- [26]: 0.001222933750014546
 - h. Create $10,000~p_{new}$ p_{old} values using the same simulation process you used in parts (a) through (g) above. Store all 10,000 values in a NumPy array called **p_diffs**.

```
[27]: # Setting single probability value for null testing purpose
    p = df2.converted.mean()

# Simulating binomial distribution taking given p into account
    newp_conv = np.random.binomial(n_new, p, int(1e4))/n_new
    oldp_conv = np.random.binomial(n_old, p, int(1e4))/n_old

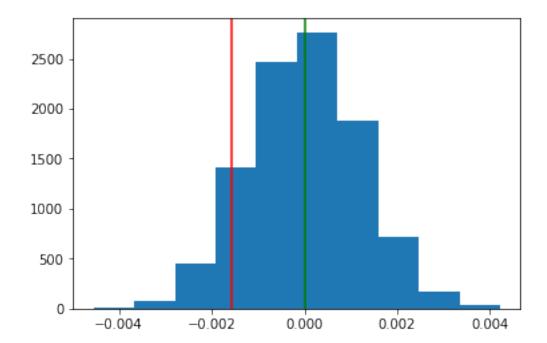
# Calculating mean conversion differences
    p_diffs = newp_conv - oldp_conv
    p_diffs.mean()
```

- [27]: 1.2451989387239457e-05
 - i. Plot a histogram of the **p_diffs**. Does this plot look like what you expected? Use the matching problem in the classroom to assure you fully understand what was computed here.

```
# Displaying observed difference and standard deviation for future reference
obs_diff, p_diffs.std()
```

[28]: (-0.0015782389853555567, 0.0012032803783433657)

```
[29]: # Plotting the histogram
plt.hist(p_diffs)
plt.axvline(obs_diff, color='r')
plt.axvline(p_diffs.mean(), color='g');
```



j. What proportion of the $\mathbf{p_diffs}$ are greater than the actual difference observed in $\mathbf{ab_data.csv?}$

```
[30]: pval1 = (p_diffs > obs_diff).mean()
pval1
```

[30]: 0.90600000000000003

k. Please explain using the vocabulary you've learned in this course what you just computed in part **j.** What is this value called in scientific studies? What does this value mean in terms of whether or not there is a difference between the new and old pages?

Part j. outcome is a **p-value**.

P-value is the probability of observing a statistic (here: mean difference in conversion rate) if the null hypothesis is true.

In this case, a **very high p-value of roughly 0.91** (meaning 91% probability) suggests we should definitely **not reject the null hypothesis**.

l. We could also use a built-in to achieve similar results. Though using the built-in might be easier to code, the above portions are a walkthrough of the ideas that are critical to correctly thinking about statistical significance. Fill in the below to calculate the number of conversions for each page, as well as the number of individuals who received each page. Let n_old and n_new refer the the number of rows associated with the old page and new pages, respectively.

```
[31]: df2.head(1)
[31]:
         user_id
                                   timestamp
                                                   group landing_page
                                                                       converted
      2
          661590 2017-01-11 16:55:06.154213 treatment
                                                             new page
[32]: convert old = df2.query('group == "control"').converted
[33]: # importing statsmodels
      import statsmodels.api as sm
      # Assigning values for z-test: old page and new page convertion proportions
      convert_old = df2.query('group == "control" and converted == 1').shape[0]
      convert_new = df2.query('group == "treatment" and converted == 1').shape[0]
      n_old = df2.query('group == "control"').shape[0]
      n_new = df2.query('group == "treatment"').shape[0]
     /opt/conda/lib/python3.6/site-packages/statsmodels/compat/pandas.py:56:
     FutureWarning: The pandas.core.datetools module is deprecated and will be
     removed in a future version. Please use the pandas.tseries module instead.
       from pandas.core import datetools
       m. Now use stats.proportions_ztest to compute your test statistic and p-value. Here is a
          helpful link on using the built in.
```

```
[34]: from statsmodels.stats.proportion import proportions_ztest

zscore, pval2 = proportions_ztest([convert_new, convert_old], [n_new, n_old],

alternative='larger')
zscore, pval2
```

[34]: (-1.3109241984234394, 0.90505831275902449)

```
[35]: # Compare ztest results to earlier calculations
print(zscore * p_diffs.std() - obs_diff )
print(pval2 - pval1)
```

8.29619897127e-07 -0.000941687240976

n. What do the z-score and p-value you computed in the previous question mean for the conversion rates of the old and new pages? Do they agree with the findings in parts **j.** and **k.**?

The **z-score** (stat) of -1.31 tells that the observed statistic is 1.31 standard deviations smaller than the mean - in this case we are testing the null hypothesis, so the mean is 0.

The p-value (pval) denotes probability for the statistic or a more extreme value to be observed if the null hypothesis is true.

The values correspond with those in parts j. and k., therefore again we fail to reject the null hypothesis.

Part III - A regression approach

- 1. In this final part, you will see that the result you achieved in the A/B test in Part II above can also be achieved by performing regression.
 - a. Since each row is either a conversion or no conversion, what type of regression should you be performing in this case?

Logistic Regression

b. The goal is to use **statsmodels** to fit the regression model you specified in part **a.** to see if there is a significant difference in conversion based on which page a customer receives. However, you first need to create in df2 a column for the intercept, and create a dummy variable column for which page each user received. Add an **intercept** column, as well as an **ab_page** column, which is 1 when an individual receives the **treatment** and 0 if **control**.

```
[36]:
     import statsmodels.api as sm
      df2['intercept'] = 1
[37]:
     df2.head(1)
[37]:
         user_id
                                    timestamp
                                                    group landing_page
                                                                         converted
      2
          661590
                  2017-01-11 16:55:06.154213 treatment
                                                              new_page
         intercept
      2
                 1
[38]: df2['ab_page'] = 0
      df2.loc[df2.group == 'treatment', 'ab_page'] = 1
[39]:
     df2.head(1)
[39]:
         user_id
                                    timestamp
                                                    group landing_page
                                                                         converted
                  2017-01-11 16:55:06.154213 treatment
                                                                                 0
                                                              new_page
         intercept
                    ab_page
      2
                 1
```

c. Use **statsmodels** to instantiate your regression model on the two columns you created in part b., then fit the model using the two columns you created in part b. to predict whether or not an individual converts.

```
[40]: log_mdl = sm.Logit(df2.converted, df2[['intercept', 'ab_page']])
results = log_mdl.fit()
```

Optimization terminated successfully.

Current function value: 0.366118

Iterations 6

d. Provide the summary of your model below, and use it as necessary to answer the following questions.

```
[41]: # Fixing deprecated function
from scipy import stats
stats.chisqprob = lambda chisq, df: stats.chi2.sf(chisq, df)

# Showing results
results.summary()
```

[41]: <class 'statsmodels.iolib.summary.Summary'>

Logit Regression Results

=========	=======				========	=======	========	
Dep. Variable	:	converted			Observations:		290584	
Model:		Logit			esiduals:		290582	
Method:		MLE			odel:		1	
Date:	Fr	Fri, 19 Mar 2021		Pseu	do R-squ.:	8.077e-06		
Time:		22:15:20		Log-	Likelihood:	-1.0639e+05		
converged:		True		LL-N	ull:	-1.0639e+05		
				LLR	p-value:		0.1899	
	====== coef 	std eri	·===== ·	z	P> z	[0.025	0.975]	
intercept	-1.9888	0.008	3 -24	6.669	0.000	-2.005	-1.973	
ab_page	-0.0150	0.011	. -	1.311	0.190	-0.037	0.007	
========	=======			=====	========		========	

11 11 11

e. What is the p-value associated with **ab_page**? Why does it differ from the value you found in **Part II**? **Hint**: What are the null and alternative hypotheses associated with your regression model, and how do they compare to the null and alternative hypotheses in **Part II**?

P-value associated with ab page is **0.19**.

It differs from p-values in Part II because in this case the p-value comes from a two-tailed test. That is here, this is a **probability that the observed statistic** - the difference in means - **is not equal to zero**.

With p-value well over 5% (.05), version of the page does **not seem statistically significant** in relation to the conversion rate.

f. Now, you are considering other things that might influence whether or not an individual converts. Discuss why it is a good idea to consider other factors to add into your regression

model. Are there any disadvantages to adding additional terms into your regression model?

In this particular case introducing other factors into the regression model makes much sense simply because we haven't yet found the answer to the original question. That is, we can't conclude whether the new page is better than the old one.

Potential problems include: - R-squared value becomes less useful, - Multicollinearity (when explanatory variables are related to each other), - A linear relationship may not exist at all.

g. Now along with testing if the conversion rate changes for different pages, also add an effect based on which country a user lives in. You will need to read in the **countries.csv** dataset and merge together your datasets on the appropriate rows. Here are the docs for joining tables.

Does it appear that country had an impact on conversion? Don't forget to create dummy variables for these country columns - **Hint: You will need two columns for the three dummy variables.** Provide the statistical output as well as a written response to answer this question.

```
[42]: df_ctr = pd.read_csv("countries.csv")
      df_ctr.head(1)
[42]:
         user_id country
      0
          834778
                      UK
[43]: print(df_ctr.country.nunique())
      print(df_ctr.country.unique())
     3
     ['UK' 'US' 'CA']
[44]: # Joining dataframes
      df2 = df2.join(df_ctr.set_index('user_id'), on='user_id')
      df2.head(1)
                                                   group landing_page
[44]:
         user_id
                                    timestamp
                                                                        converted
          661590
                 2017-01-11 16:55:06.154213 treatment
                                                              new_page
                                                                                 0
      2
         intercept
                   ab_page country
      2
                                  US
                 1
                           1
[45]: # Creating dummy columns for countries
      df2[['CA', 'UK', 'US']] = pd.get_dummies(df2.country)
[46]: # Verifying
      df2.head()
[46]:
                                                                        converted
         user id
                                    timestamp
                                                   group landing page
      2
          661590
                  2017-01-11 16:55:06.154213
                                                                                 0
                                               treatment
                                                              new_page
      3
          853541
                  2017-01-08 18:28:03.143765
                                                                                 0
                                               treatment
                                                              new_page
          679687
                  2017-01-19 03:26:46.940749
                                               treatment
                                                              new_page
                                                                                 1
```

```
8
    817355 2017-01-04 17:58:08.979471 treatment
                                                          new_page
                                                                              1
    839785 2017-01-15 18:11:06.610965
9
                                          treatment
                                                          new_page
                                                                              1
   intercept ab_page country
                                 CA
                                     UK
                                          US
2
                             US
                                   0
                                       0
                                           1
            1
                     1
            1
                     1
3
                             US
                                  0
                                       0
                                           1
6
            1
                     1
                             CA
                                      0
                                           0
                                  1
8
            1
                     1
                             UK
                                   0
                                       1
                                           0
9
            1
                     1
                             CA
                                           0
                                   1
                                       0
```

```
[47]: # Fitting a model and displaying summary
log_mdl2 = sm.Logit(df2.converted, df2[['intercept', 'ab_page', 'CA', 'UK']])
results2 = log_mdl2.fit()
results2.summary()
```

Optimization terminated successfully.

Current function value: 0.366113

Iterations 6

[47]: <class 'statsmodels.iolib.summary.Summary'>

Logit Regression Results

______ No. Observations: Dep. Variable: 290584 converted Model: Logit Df Residuals: 290580 Method: MLE Df Model: 3 Date: Fri, 19 Mar 2021 Pseudo R-squ.: 2.323e-05 22:15:21 Log-Likelihood: Time: -1.0639e+05 LL-Null: -1.0639e+05 converged: True LLR p-value: 0.1760

	coef	std err	z	P> z	[0.025	0.975]
intercept	-1.9893	0.009	-223.763	0.000	-2.007	-1.972
ab_page	-0.0149	0.011	-1.307	0.191	-0.037	0.007
CA	-0.0408	0.027	-1.516	0.130	-0.093	0.012
UK	0.0099	0.013	0.743	0.457	-0.016	0.036
=========						
11 11 11						

```
[48]: # Exponentiating coefficients
np.exp(-0.0149), np.exp(-0.0408), np.exp(0.0099)
```

[48]: (0.9852104557227469, 0.96002111497165088, 1.0099491671175422)

The model suggests there is **no statistically significant impact of user's country** in relation to conversion rate. Values of exponentiated coefficients being **close to 1** means that with a change in country or page version, conversion is roughly **one times more likely**, holding all else constant.

h. Though you have now looked at the individual factors of country and page on conversion, we would now like to look at an interaction between page and country to see if there are significant effects on conversion. Create the necessary additional columns, and fit the new model.

Provide the summary results, and your conclusions based on the results.

```
[49]: # Creating interaction columns
     df2['ab page CA'] = df2.ab page * df2.US
     df2['ab_page_UK'] = df2.ab_page * df2.UK
     df2.head(3)
[50]:
[50]:
        user_id
                                                group landing_page
                                  timestamp
                                                                    converted
         661590 2017-01-11 16:55:06.154213 treatment
     2
                                                          new_page
                                                                            0
     3
         853541
                 2017-01-08 18:28:03.143765 treatment
                                                                            0
                                                          new_page
         679687 2017-01-19 03:26:46.940749 treatment
                                                          new_page
                                                                            1
        intercept ab_page country CA UK US
                                               ab_page_CA
                                                          ab page UK
     2
                1
                         1
                                US
                                     0
                                        0
                                            1
                                                        1
                                                                    0
                                                                    0
     3
                1
                         1
                                US
                                     0
                                         0
                                            1
                                                        1
                1
                         1
                                CA
                                     1
                                        0
                                            0
                                                        0
                                                                    0
[51]: log_mdl3 = sm.Logit(df2.converted, df2[['intercept', 'ab_page', 'CA', 'UK', L
      results3 = log_mdl3.fit()
     results3.summary()
```

Optimization terminated successfully.

Current function value: 0.366109

Iterations 6

CA

[51]: <class 'statsmodels.iolib.summary.Summary'>

-0.0175

0.038

Logit Regression Results

Dep. Variable	converted			No. C	bservations:	290584		
Model:		Logit			siduals:	290578		
Method:		MLE			del:	5		
Date:	Fr	Fri, 19 Mar 2021			lo R-squ.:	3.482e-05		
Time:		22:15:22		Log-Likelihood:		-1.0639e+05		
converged:		True		LL-Null:		-1.0639e+05		
				LLR p-value:		0.1920		
=========	=======	=======	=====		=========	=======	=======	
	coef	std err		Z	P> z	[0.025	0.975]	
intercept	 -1.9865	0.010	 -206	 3.344	0.000	-2.005	-1.968	
ab_page	-0.0674	0.052	-1	1.297	0.195	-0.169	0.034	

-0.465

0.642

-0.091

0.056

UK	-0.0057	0.019	-0.306	0.760	-0.043	0.031
ab_page_CA	0.0469	0.054	0.872	0.383	-0.059	0.152
ab_page_UK	0.0783	0.057	1.378	0.168	-0.033	0.190
========	========	=======		=======	========	
11 11 11						

Adding page-country interactions to the model provides **lower p-values** compared to when considering country influence independently (ab page CA, ab page UK).

However, these p-values are still higher than the .05 threshold of significance.

Therefore, using all above methods we can conclude the same:

There is no statistical nor practical significance in rejecting the null hypothesis and deploying the new page, based on the assessed data.

This might be subject to change if for example users are given more time for testing and getting accustomed with the new page layout.

Based on the above analysis I would not switch to the new page.

Finishing Up

Congratulations! You have reached the end of the A/B Test Results project! You should be very proud of all you have accomplished!

Tip: Once you are satisfied with your work here, check over your report to make sure that it is satisfies all the areas of the rubric (found on the project submission page at the end of the lesson). You should also probably remove all of the "Tips" like this one so that the presentation is as polished as possible.

0.3 Directions to Submit

Before you submit your project, you need to create a .html or .pdf version of this notebook in the workspace here. To do that, run the code cell below. If it worked correctly, you should get a return code of 0, and you should see the generated .html file in the workspace directory (click on the orange Jupyter icon in the upper left).

Alternatively, you can download this report as .html via the **File** > **Download as** submenu, and then manually upload it into the workspace directory by clicking on the orange Jupyter icon in the upper left, then using the Upload button.

Once you've done this, you can submit your project by clicking on the "Submit Project" button in the lower right here. This will create and submit a zip file with this .ipynb doc and the .html or .pdf version you created. Congratulations!

```
[52]: from subprocess import call call(['python', '-m', 'nbconvert', 'Analyze_ab_test_results_notebook.ipynb'])
```

[52]: 0