This Analysis covers hypothesis formulation, testing, and interpreation. Data is received from Kaggle (https://www.kaggle.com/zhangluyuan/ab-testing? select=ab data.csv) - created to test the effectivness of 2 designs of a website page.

Here are the 5 steps of this Analysis:

```
##### 1. Experiment Definition
##### 2. Data collection and preparation
##### 3. Visualization
##### 4. Hypothesis testing
##### 5. Conclusions
```

1. EXPERIMENT DEFINITION

The data is related with an online e-commerce business, they are doing some changes in the website and want to see if the new website helps to convert customer to purchaser. Current conversion rate is 13%, if there is an increase 2%, then the effectivness of new webpage will be justified. So, with new website 15% conversion rate needs to be created. So, we are trying to test if the new webpage is better than the old one, but we don't know either it is better or worst. So, we are doing tow-tailed test:

```
H0: p = p0
H1: p! = p0
```

p & p0 are the conversion rate of the new and old design. Our confidence level is 95%: a = 0.05

a: "if the probability of observing a result as extreme or more (p-value) is lower than α , then we reject the Null hypothesis". a = 0.05 means 5% probability with confidence of (1 - a) of 95%.

Define variables

Control group - group of customers shown the old design.

Treatment group - group of customers shown the new design.

Our dependent variable (i.e., our target measurement) is the conversion_rate, which is a binary variable: 0 means user did not buy the product and 1 means user bought the product.

sample size for each group is calculated from the "Power Analysis", which depends on: (a) Power of test (1-B) -- probability of finding a statistical difference between the groups in test set when a difference is actually present. This is usually set at 0.8 by convention. (b) Alpha value (a) - The critical value set at 0.05. (c) Effect size -- How big of a difference we expect there to be between teh conversion rates.

we need a difference of 2%, so we need to use 13% and 15% to calculate the effect size we want. Rest Python will caclculate.

```
In [2]:
         # Packages imports
         import numpy as np
         import pandas as pd
```

```
import scipy.stats as stats
import statsmodels.stats.api as sms
import matplotlib as mpl
import matplotlib.pyplot as plt
import seaborn as sns
from math import ceil
%matplotlib inline
```

```
In [3]:
         # plot styling preferences
         plt.style.use('seaborn-whitegrid')
         font = {'family': 'Helvetica', 'weight':'bold', 'size': 14}
         mpl.rc('font', **font)
         # effect size calculation based on our expected rates
         effect size = sms.proportion effectsize(0.13, 0.15)
         # calculating sample size needed
         required n = sms.NormalIndPower().solve power(effect size, power=0.8, alpha=0.05, ratio
         # Rounding up to next whole number
         required_n = ceil(required_n)
         print(required n)
```

Above result suggests, we need at least 4720 observations for each group.

The meaning of power = 0.8 suggests if actual difference in conversion rates (13% vs. 15%) exists between our designs, there is about 80% chance to detect it statistically significant in our test with the sample size we calculated.

2. Data collection and preparation

4720

data is downloaded from Kaggle. Python is used to prepare and analyze the data. Each group will consists of 4720 random rows.

```
In [5]:
          df = pd.read_csv('ab_data.csv')
          df.head()
             user_id
                                                         landing_page converted
Out[5]:
                                   timestamp
                                                  group
         0 851104 2017-01-21 22:11:48.556739
                                                  control
                                                              old_page
                                                                                0
            804228 2017-01-12 08:01:45.159739
                                                  control
                                                              old_page
            661590 2017-01-11 16:55:06.154213 treatment
                                                             new_page
            853541 2017-01-08 18:28:03.143765 treatment
                                                             new_page
            864975 2017-01-21 01:52:26.210827
                                                              old_page
                                                  control
                                                                                1
In [6]:
          df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
```

Out[11]:

user id

```
An Introduction to AB Test in Python
         RangeIndex: 294478 entries, 0 to 294477
         Data columns (total 5 columns):
              Column
                            Non-Null Count
                                             Dtype
                            -----
              ----
              user_id 294478 non-null int64
          0
                          294478 non-null object
          1
            timestamp
          2
              group
                            294478 non-null object
              landing_page 294478 non-null
                                            object
              converted
                            294478 non-null int64
         dtypes: int64(2), object(3)
         memory usage: 11.2+ MB
 In [7]:
          ## To make sure all the control group are seeing the old page and viceversa
          pd.crosstab(df['group'], df['landing_page'])
 Out[7]: landing_page new_page old_page
               group
                          1928
                                 145274
              control
                        145311
                                   1965
            treatment
         For this analysis we only need the group and converted columns. But, before
        sampling, let's make sure there are no users that have been sampled multiple
        times.
 In [8]:
          session_counts = df['user_id'].value_counts(ascending = False)
          multi_users = session_counts[session_counts > 1].count()
          print(f'There are {multi users} that appear multiple times in the dataset')
         There are 3894 that appear multiple times in the dataset
In [10]:
          # Lets remove the repetition from the DataFrame to avoid samping the same users twice.
          users_to_drop = session_counts[session_counts > 1].index
          df = df[~df['user id'].isin(users to drop)]
          print(f'The updated dataset now has {df.shape[0]} entries')
         The updated dataset now has 286690 entries
In [11]:
          # Lets sample our data from each group using DataFrame.sample(), which performs Simple
```

```
# random_state = 22 suggests the results are reproducible.
control_sample = df[df['group'] == 'control'].sample(n=required_n, random_state = 22)
treatment sample = df[df['group'] == 'treatment'].sample(n=required n, random state = 2
ab_test = pd.concat([control_sample, treatment_sample], axis = 0)
ab test.reset index(drop = True, inplace = True)
ab_test
```

group landing_page converted

timestamp

| converted | landing_page | group | timestamp | user_id | |
|-----------|--------------|-----------|----------------------------|---------|------|
| 0 | old_page | control | 2017-01-21 03:43:17.188315 | 763854 | 0 |
| 0 | old_page | control | 2017-01-18 06:38:13.079449 | 690555 | 1 |
| 0 | old_page | control | 2017-01-06 21:13:40.044766 | 861520 | 2 |
| 0 | old_page | control | 2017-01-05 16:42:36.995204 | 630778 | 3 |
| 0 | old_page | control | 2017-01-04 15:31:21.676130 | 656634 | 4 |
| | | | | | ••• |
| 0 | new_page | treatment | 2017-01-14 22:02:29.922674 | 908512 | 9435 |
| 0 | new_page | treatment | 2017-01-05 00:57:16.167151 | 873211 | 9436 |
| 0 | new_page | treatment | 2017-01-20 18:56:58.167809 | 631276 | 9437 |
| 0 | new_page | treatment | 2017-01-03 08:10:57.768806 | 662301 | 9438 |
| 1 | new_page | treatment | 2017-01-19 10:56:01.648653 | 944623 | 9439 |
| | | | | | |

9440 rows × 5 columns

```
In [12]:
          ab test.info()
         <class 'pandas.core.frame.DataFrame'>
         RangeIndex: 9440 entries, 0 to 9439
         Data columns (total 5 columns):
            Column
                         Non-Null Count Dtype
            user id
                           9440 non-null int64
          0
                           9440 non-null object
          1
              timestamp
                           9440 non-null
                                           object
              group
              landing_page 9440 non-null
                                           object
              converted
                           9440 non-null
                                           int64
         dtypes: int64(2), object(3)
         memory usage: 368.9+ KB
In [13]:
          ab_test['group'].value_counts()
Out[13]: control
                      4720
                      4720
         treatment
         Name: group, dtype: int64
```

3. Visualising the results

```
In [14]:
          conversion_rates = ab_test.groupby('group')['converted']
          # Std. deviation of the proportion
          std_p = lambda x: np.std(x, ddof = 0)
          # Std. error of the proportion (std / sqrt(n))
          se_p = lambda x: stats.sem(x, ddof = 0)
          conversion_rates = conversion_rates.agg([np.mean, std_p, se_p])
          conversion_rates.columns = ['conversion_rate', 'std_deviation', 'std_error']
```

```
conversion rates.style.format('{:.3f}')
```

Out[14]: conversion rate std deviation std error

| group | | | |
|-----------|-------|-------|-------|
| control | 0.123 | 0.329 | 0.005 |
| treatment | 0.126 | 0.331 | 0.005 |

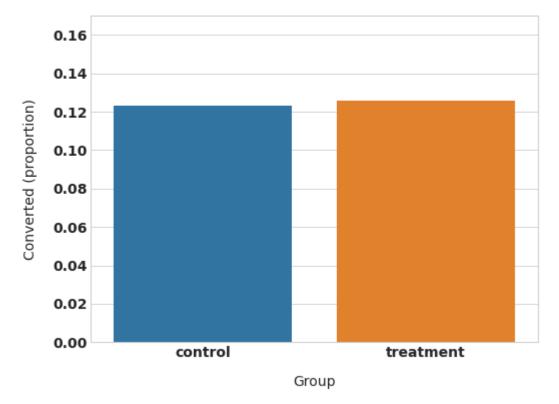
Above table shows conversion_rate for both group is close 12.3% vs 12.6%.

```
In [15]:
          plt.figure(figsize = (8,6))
          sns.barplot(x = ab_test['group'], y = ab_test['converted'], ci = False)
          plt.ylim(0, 0.17)
          plt.title('Conversion rate by group', pad=20)
          plt.xlabel('Group', labelpad=15)
          plt.ylabel('Converted (proportion)', labelpad=15)
```

Out[15]: Text(0, 0.5, 'Converted (proportion)')

```
findfont: Font family ['Helvetica'] not found. Falling back to DejaVu Sans. findfont: Font family ['Helvetica'] not found. Falling back to DejaVu Sans.
findfont: Font family ['Helvetica'] not found. Falling back to DejaVu Sans.
```

Conversion rate by group



conversion rate of control group is slightly lower than treatment group. but, conversion rate of the control group is even lower than the average rate (12.3%) vs. 13%). This might suggests that there is some variation in results when sampling from a population.

lets check if the treatment group's higher value is statistically significant.

4. Hypothesis testing

Due to large sample size, we can use the normal approximation for the calculation of the p-value (i.e., z-test).

we can use the statsmodels.stats.proportion module to get the p-value and confidence intervals.

```
In [16]:
          from statsmodels.stats.proportion import proportions_ztest, proportion_confint
          control_results = ab_test[ab_test['group'] == 'control']['converted']
          treatment results = ab test[ab test['group'] == 'treatment']['converted']
          n_con = control_results.count()
          n treat = treatment results.count()
          successes = [control results.sum(), treatment results.sum()]
          nobs = [n con, n treat]
          z stat, pval = proportions ztest(successes, nobs=nobs)
          (lower con, lower treat), (upper con, upper treat) = proportion confint(successes, nobs
          print(f'z statistic: {z_stat:.2f}')
          print(f'p-value: {pval:.3f}')
          print(f'ci 95% for control group: [{lower con:.3f}, {upper con:.3f}]')
          print(f'ci 95% for treatment group: [{lower treat:.3f}, {upper treat:.3f}]')
         z statistic: -0.34
         p-value: 0.732
         ci 95% for control group: [0.114, 0.133]
         ci 95% for treatment group: [0.116, 0.135]
```

5. Conclusions

p-value of 0.732 suggests our result is statistically insignificant, that means we cannot reject the Null Hypothesis H0, which suggests that the new design did not perform significantly different than the old one.

Additionally, confidence interval for the treatment group is of 11.6% - 13.5%, which means it includes the baseline value of 13% conversion rate, but doesn't include target value of 15% (the 2% uplift).

Above result suggests that the new design doesn't increase the conversion rate, so new design is not an improvement over the old one.