In [1]:

from IPython.display import Image

1.Summary

Thank you for taking the time to complete Quora Data Challenge! We're excited about you continuing the interview process with us.

The data challenge simulates a typical experiment analysis you would be expected to work on at Quora. In this challenge, we will provide several datasets and ask you to analyze the results and provide product recommendations for a particular experiment.

The Google form is not timed but we expect you to spend around 1-2 hours answering the questions. While working on these problems, please document your process and code in any tools you are comfortable with. They don't have to be very organized but should demonstrate your thinking. We will ask you to submit that at the end. Feel free to refer to any online resources if necessary.

Suppose you are a Data Scientist on the Mobile team at Quora. The team has just introduced a new UI design to the Quora app. The goal of the new design is to increase user engagement (measured by minutes spent on site). The team ran an A/B test to evaluate the change. Using the data, help the team understand the impact of the UI change better.

In []:

https://www.zhihu.com/question/20458233/answer/1724891055

A/B Testing isn't an useful testing as new experiences. There will be two trends, first is that people like old way and old experience, thats called change aversion. Second is thay they like new staff, so they don't attract by the change due to its convienent, instead they just explore the new feature when they first met new things and next time they will back to normal when they have explored everything. Thats called novelty effect.

what is going to be the plateaued experience so that i can actually make a robust decision.

Table Schema

t1_user_active_min.csv This table contains active minutes data logged after experiment started(2019-02-06 to 2019-07-05). Each row represents the total number of minutes spent on site for each user on a date. If a user never visited the site for a given date, there wouldn't be data for that uid on that date. (for 50000 user, there are 3376 users never logged in Feb to May time period)

- uid: user ID
- dt: date when corresponding active minutes are registered
- active_mins: number of minutes spent on site for the date

t2_user_variant.csv This table contains users' treatment assignment. Each row represents the assignment information for a unique user.

- uid: user ID
- variant_number: the experiment variant user is in. 0 for control, 1 for treatment
- dt: date when user entered the experiment, should be '2019-02-06' for all users
- signup_date: the date string that user signed up on

t3_user_active_min_pre.csv This table contains active minutes data before the experiment started. It has a similar format as t1, except the dt range can extend before the experiment start date. ('2018-08-10' to '2019-02-05')

- uid: user ID
- dt: date when corresponding active minutes are registered
- active_mins: number of minutes spent on site for the date

t4_user_attributes.csv This table contains data about some user attributes. Each row represents attributes of a unique user.

- uid: user ID
- user_type: segment that a user belongs to, measured by activity level of the user. Can be 'new_user', 'non_reader', 'reader' or 'contributor'
- gender: user gender. Can be 'male', 'female' or 'unknown'

```
In [ ]:

In [ ]:
```

In order to measure the experiment effect on user engagement, let's say you want to perform a t-test of the difference in minutes spent on site, what would be your unit of analysis (i.e. when conducting a t-test, we calculate means and variances of two samples. What does each data point represent in the samples?)? *

- Total minutes per user in each group (number of data points in each group = number of users in that group)
- Total minutes per user per day in each group (number of data points in each group = number of users in that group x number of days of the experiment)
- Total minutes per group per day (number of data points in each group = number of days of the experiment)

Why do you think the other choices are not correct? Hint: think of the key assumptions of running a t-test.

Total minutes per user per day in each group regard each datapoint as in individual independent happening

we want to increase 'user' spend time increase or not.

Why do you think the other choices are not correct? Hint: think of the key assumptions of running a t-test. *

For the t-test assumption, if the distribution are not normal distribution, we need to follow the assumption: the population variance are equal. For second part, the variance are different because each user login days are not equal. For thrid option, the two part have different users login per day. So their population variance won't be equal.

Please enter the lower bound of your confidence interval (e.g. if the confidence Interval is [-1.00, 2.00], enter -1.00) = -38.591763263087756

Please enter the upper bound of your confidence interval (e.g. if the confidence Interval is [-1.00, 2.00], enter 2.00) = 38.22913569692153

In	[]:	
In	[]:	

T-Test Assumptions

- 1. The first assumption made regarding t-tests concerns the scale of measurement. The assumption for a t-test is that the scale of measurement applied to the data collected follows a continuous or ordinal scale, such as the scores for an IQ test.
- 2. The second assumption made is that of a simple random sample, that the data is collected from a representative, randomly selected portion of the total population.
- 3. The third assumption is the data, when plotted, results in a normal distribution, bell-shaped distribution curve. When a normal distribution is assumed, one can specify a level of probability (alpha level, level of significance, p) as a criterion for acceptance. In most cases, a 5% value can be assumed.
- 4. The fourth assumption is a reasonably large sample size is used. A larger sample size means the distribution of results should approach a normal bell-shaped curve.
- 5. The final assumption is homogeneity of variance. Homogeneous, or equal, variance exists when the standard deviations of samples are approximately equal.

```
In [ ]:
```

N > 100 T-distribution equal to normal distribution

Initial hypothesis: change the UI from A to B will increase customer active minutes on this page

Which metric to use?

- Total minutes per group? No, Total minues from Group A or Group B can not real reflct in impact
- Total minutes per user? No, We have two group, we need caculate active minutes by group classification
- Total minutes per user in each group? We split group into control group and experiment group. If we groupby 'uid', we got 40000 control group and 10000 experiment. The data points will be 40000: 10000
- Total minutes per user per day in each group? Further more, the data point will be each raw data point without any preprocess

If we envolve all user pre-experiment data, we might need to caculate the difference between different time and ratio of increase/all data point.

(unique user who increase time spend) / (unique users who accessd in this group)

```
In []:

In []:
```

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3.Preprocess

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3.1 Hyperparameters

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```
In [1]:
    class HyperParameters(object):
        """:arg
        This class will be used to transmit hyperparameters between class.parameters
        Most of class can inherit this class and its hyperparameters

### def __init__(self):
        """:arg

#### we use this to test whether other class read HyperParameters() successfully
        self.TEST = 1

#### you can change this root path in this class and import_data() function will s
        self.ROOTPATH = 'D:\\OneDrive\\03_Academic\\23_Github\\44_Quora_Data_Challenge\
#### confidence inveral value
        self.ALPHA = 0.95
```

3.2 Import Data

```
In [2]:
         from class 31 hyperparameters import HyperParameters
         import pandas as pd
         # measure running time
         from time import time
         import os
         class ImportData(HyperParameters):
             """:arg
             ....
             def __init__(self):
                 """:arg
                 HyperParameters.__init__(self)
             def import data(self):
                 """:arg
                 Use this function to complete import data process
                 print("*" * 50, "Start import data", "*" * 50)
                 start time = time()
                 path t1 = os.path.join(self.ROOTPATH, 't1 user active min.csv')
                 path_t2 = os.path.join(self.ROOTPATH, 't2_user_variant.csv')
                 path_t3 = os.path.join(self.ROOTPATH, 't3_user_active_min_pre.csv')
                 path t4 = os.path.join(self.ROOTPATH, '9999')
                 df_t1 = pd.read_csv(path_t1)
                 df t2 = pd.read csv(path t2)
                 df_t3 = pd.read_csv(path_t3)
                 df_t4 = pd.read_csv(path_t4)
                 return (df_t1, df_t2, df_t3, df_t4)
```

3.3.EDA

```
In [3]: from class_31_hyperparameters import HyperParameters

# use sns for visulization
import seaborn as sns
# numpy
import numpy as np
# solve statistical problem
from scipy import stats
```

```
class EDA(HyperParameters):
   """:arg
   Preproceess EDA
   ....
   def __init__(self):
       """:arg
       HyperParameters. init (self)
   def outlier eda(self, df):
       """:arg
       We use several way to detect and remove outliers
       Args:
       df:DataFrame
           This can be data_t1 or data_t3. These two dataframe need ourlier detect and
        .....
       # first we need to make sure this is outlier or error exist
       # we use boxplot() to demenstrate outliers
       sns.boxplot(x=df['active_mins'])
       # use z-score to eliminate outliers, the output is for indivudal data point
       z_score = np.abs(stats.zscore(df['active_mins']))
       # set a z-score threshold, any greater than this value will be eliminate
       z threshold = 3
       print(z threshold)
       # filter that ooutlier rows in dataframe
       index_outlier = np.where(z_score > z_threshold)
       # print out result
       print('We have {} data points are outliers'.format(len(index_outlier[0])))
       # drop these rows by index
       df = df.drop(index_outlier[0])
        return df
```

3.4 Join Data

```
In [4]:
    from class_31_hyperparameters import HyperParameters
    import pandas as pd
    # test normal distribution
    from scipy import stats

class JoinData(HyperParameters):
    """:arg
    """
```

```
def __init__(self):
    """:arg
   HyperParameters.__init__(self)
def join_data_uid(self, df_t1, df_t2, df_t3, df_t4):
    """:arg
   We use function join to dataframe
    # outer join df_t1 and df_t2 by uid
   df_t1t2 = pd.merge(df_t1, df_t2, on=['uid'], how='outer')
    # clean data drop NaN by row
    df_clean = df_t1t2.dropna(axis=0)
    # print clearn process
    print("Before {}, After {}".format(df_t1t2.shape[0], df_clean.shape[0]))
    # split into control group and treatment group and aggregate by user
    df_control = df_clean[df_clean['variant_number'] == 0].groupby(['uid'], as_inde
    # unit is total time spend per user per group, sometimes average or middle migh
    df treat = df clean[df clean['variant number'] == 1].groupby(['uid'], as index=
    return df clean, df control, df treat
def join_data_group(self, df_t1, df_t2, df_t3, df_t4):
    """:arg
    We use function join to dataframe
    0.00
   # outer join df_t1 and df_t2 by uid
    df t1t2 = pd.merge(df t1, df t2, on=['uid'], how='outer')
    # clean data drop NaN by row
   df clean = df t1t2.dropna(axis=0)
    # print clearn process
    print("Before {}, After {}".format(df_t1t2.shape[0], df_clean.shape[0]))
    # split into control group and treatment group and aggregate by user
    df_control = df_clean[df_clean['variant_number'] == 0].groupby(['dt_x'], as_ind
    # unit is total time spend per user per group, sometimes average or middle migh
    df_treat = df_clean[df_clean['variant_number'] == 1].groupby(['dt_x'], as_index
    return df_clean, df_control, df_treat
def join_data_dt(self, df_t1, df_t2, df_t3, df_t4):
    """:arg
   I choose the unit is 'Total minutes per user per day in each group.
    First, we split data into control (not chagne) and treatment (changed) group
   Second, we aggregate data records (88197 / 34175) in each group by date(['dt_x'
   This meaning is, in this group, everyday the total minutes from user log in act
   Typically, we have different log in user number in this whole 150 experiment da
   Third, we divide everyday total mintues by the number of user who log in this d
   They we can test them with normal distribution. If you want to calculate t-test
    be normal distribution (stats.shapiro(df))
    # outer join df_t1 and df_t2 by uid
    df_t1t2 = pd.merge(df_t1, df_t2, on=['uid'], how='outer')
   # clean data drop NaN by row
```

```
df clean = df t1t2.dropna(axis=0)
# print clearn process
print("Before {}, After {}".format(df_t1t2.shape[0], df_clean.shape[0]))
# split into control group and treatment group and aggregate by user
df_dt_control = df_clean[df_clean['variant_number'] == 0].groupby(['dt_x'], as_
# unit is total time spend per user per group, sometimes average or middle migh
df dt treat = df clean[df clean['variant number'] == 1].groupby(['dt x'], as in
# we caculate how many user log in each day
df_usr_control = df_clean[df_clean['variant_number']==0].groupby(['dt_x'], as_i
# caculate how many user login each day in treatment group
df usr treat = df clean[df clean['variant number']==1].groupby(['dt x'], as ind
# (total minutes per day) / (number of users who login in each day)
df_control = df_dt_control['active_mins'].divide(df_usr_control['uid'])
df treat = df dt treat['active mins'].divide(df usr treat['uid'])
print('For df control, its normal distribution test is {}'.format(stats.shapiro
print('For df treat, its normal distribution test is {}'.format(stats.shapiro(d
return df_clean, df_control, df_treat
```

3.5 Hypothesis

```
In [21]:
          from class 31 hyperparameters import HyperParameters
          import numpy as np
          import seaborn as sns
          # this model will output degree of freedom
          import statsmodels.stats.weightstats as st
          from scipy import stats
          class Hypothesis(HyperParameters):
              We use this class to calculate statical data and judge reject or accept hypothesis
              0.00
              def __init__(self):
                  """:arg
                  HyperParameters.__init__(self)
              def ttest ind(self, df control, df treat):
                   """:arg
                  Args:
                  df control:Series
```

```
print('Distribution of control group')
                 sns.boxplot(x = df control)
                 sns.distplot(df_control)
                 # caculate t-test for tow independent dataset
                 # because we caculate with total minutes, so the user rate 40000:10000 is imbal
                 # we need divide df_control with 4 to achieve same total minutes for 10000 user
                 # t value is t statical
                 t value, p two, degree free = st.ttest ind(x1 = df control
                                                             , x2 = df_treat
                                                             , usevar='unequal')
                 # use a diiffernt function for t-test
                 t_stat, p_value = stats.ttest_ind(df_control, df_treat, equal_var=False)
                 # default confidence interval is 95%, so one side alpha = (1-95\%)/2 = 0.025
                 if (p two < 1-self.ALPHA):</pre>
                     print('P-value {} < {}'.format(p_two, 1-self.ALPHA))</pre>
                     print('Reject H0 (control and treatment dont have siginificent differnet) h
                     print('Accept H1 (control and treatment have siginificent different) hypoth
                 else:
                     print('P-value {} > {}'.format(p two, 1 - self.ALPHA))
                     print('Accept H0 (control and treatment dont have siginificent differnet) h
                     print('Reject H1 (control and treatment have siginificent different) hypoth
                 # start cacualte confidence interval (right tail=0.025) (two tail=0.05)
                 # we have 150-200 data point, so when check t-distribution table,
                 t ci = stats.t.ppf(q = self.ALPHA, df=150)
                 # sample number
                 control_num = df_control.shape[0]
                 treat_num = df_treat.shape[0]
                 # cacualte stand diviation
                 control std = df control.std()
                 treat_std = df_treat.std()
                 # cacualte se
                 se = np.sqrt( np.square(control_std)/control_num + np.square(treat_std)/treat_n
                 sample_mean = df_control.mean() - df_treat.mean()
                 # calcuate confidenal interval
                 upper_bound = sample_mean - t_ci * se
                 lower_bound = sample_mean + t_ci * se
                 return (t_value, p_two, degree_free, t_stat, p_value, upper_bound, lower_bound)
In [ ]:
In [ ]:
In [ ]:
In [ ]:
```

7 Main Function

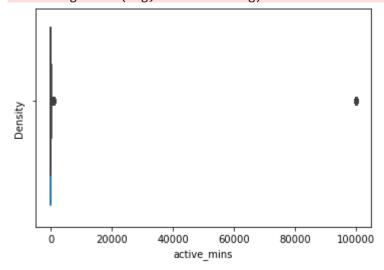
Click this Link back to Top

```
In [22]:
         from class 31 hyperparameters import HyperParameters
         from class_32_import_data import ImportData
         from class 33 eda import EDA
         from class 34 join data import JoinData
         from class 35 hypothesis import Hypothesis
          def main():
             """:arg
             All program running here by sequence
             1.import data
             2. Through EDA delete outliers
             3. Join data and split control and treatment
             class import = ImportData()
             (df_t1, df_t2, df_t3, df_t4) = class_import.import_data()
             class_eda = EDA()
             # clean df t1 dataframe by deleting outliers
             df_t1 = class_eda.outlier_eda(df_t1)
             class_join = JoinData()
             # outer join t1 table and t2 table
             df clean, df control, df treat = class join.join data dt(df t1, df t2, df t3, df t4
             class hypo = Hypothesis()
             (t_value, p_two, degree_free
              , t stat, p value, upper bound, lower bound) = class hypo.ttest ind(df control, df
             return (df_t1, df_t2, df_t3, df_t4, df_clean, df_control, df_treat
                     , t value, p two, degree free, t stat, p value, upper bound, lower bound)
          if __name__=="__main__":
             """:arg
             (df_t1, df_t2, df_t3, df_t4, df_clean, df_control, df_treat
              , t value, p two, degree free, t stat, p value, upper bound, lower bound) = main()
             print('OVER')
         *********
         We have 172 data points are outliers
         Before 1069597, After 1066230
         For df control, its normal distribution test is ShapiroResult(statistic=0.99359279870986
         94, pvalue=0.7472291588783264)
         For df treat, its normal distribution test is ShapiroResult(statistic=0.870302498340606
         7, pvalue=3.8012573644330416e-10)
         Distribution of control group
         P-value 1.0 > 0.050000000000000044
         Accept H0 (control and treatment dont have siginificent differnet) hypothesis
         Reject H1 (control and treatment have siginificent different) hypothesis
         OVER
```

D:\ProgramData\Anaconda3\envs\08 web mining\lib\site-packages\seaborn\distributions.py:2

551: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with si milar flexibility) or `histplot` (an axes-level function for histograms).

warnings.warn(msg, FutureWarning)



In []:

In [28]:

Out[28]:

df_t3

9.Test Code

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```
In [25]:
           df_t3.shape
          (1190093, 3)
Out[25]:
In [27]:
           df_t3.describe()
Out[27]:
                          uid
                                active_mins
          count 1.190093e+06
                             1.190093e+06
           mean 2.420589e+04 3.220315e+01
            std 1.417999e+04
                             1.181531e+03
                 0.000000e+00
                              1.000000e+00
           25%
                 1.197500e+04
                              2.000000e+00
           50%
                 2.393200e+04 4.000000e+00
           75%
                 3.619600e+04
                              1.400000e+01
            max 4.999900e+04 9.999900e+04
```

dt active_mins

uid

	uid	dt	active_mins
0	0	2018-09-24	3.0
1	0	2018-11-08	4.0
2	0	2018-11-24	3.0
3	0	2018-11-28	6.0
4	0	2018-12-02	6.0

1190088	49999	2018-09-15	5.0
1190089	49999	2018-09-26	8.0
1190090	49999	2018-10-20	29.0
1190091	49999	2018-12-14	3.0
1190092	49999	2019-01-28	32.0

1190093 rows × 3 columns

```
In [ ]:
 In [ ]:
In [23]:
           upper_bound
          -4.509893654334767
Out[23]:
In [24]:
          lower_bound
          -3.665273229338887
Out[24]:
 In [8]:
           df_control
                 19.893420
 Out[8]: 0
                 20.235027
                 20.529420
                 19.662953
                 20.119112
          145
                 19.716245
          146
                 18.977998
          147
                 19.840566
          148
                 19.897338
          149
                 19.860828
          Length: 150, dtype: float64
In [10]:
          df_control.describe()
```

```
150.000000
Out[10]: count
                    19.323468
          mean
                     0.646480
          std
          min
                    17.732331
          25%
                    18.895117
          50%
                    19.278251
          75%
                    19.836096
          max
                    21.135562
          dtype: float64
```

In [11]: df_treat.describe()

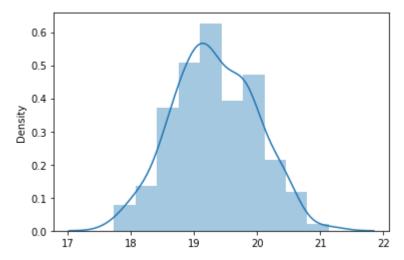
```
150.000000
          count
Out[11]:
                     23.411051
          mean
          std
                     3.057468
                     18.758485
          min
          25%
                     21.385178
          50%
                     22.300901
          75%
                     24.437592
                     32.389844
          max
          dtype: float64
```

In [13]: sns.distplot(df_control)

D:\ProgramData\Anaconda3\envs\08_web_mining\lib\site-packages\seaborn\distributions.py:2 551: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with si milar flexibility) or `histplot` (an axes-level function for histograms).

warnings.warn(msg, FutureWarning)

Out[13]: <AxesSubplot:ylabel='Density'>

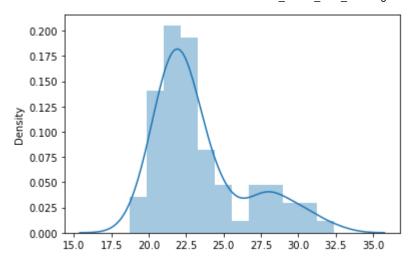


In [12]: sns.distplot(df_treat)

D:\ProgramData\Anaconda3\envs\08_web_mining\lib\site-packages\seaborn\distributions.py:2 551: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with si milar flexibility) or `histplot` (an axes-level function for histograms).

warnings.warn(msg, FutureWarning)

Out[12]: <AxesSubplot:ylabel='Density'>



```
In [9]:
           df_treat
                 28.474670
 Out[9]:
                 27.436468
                 27.181484
          3
                 26.838880
                 27.444191
                 21.384615
          145
          146
                 21.485224
          147
                 22.237192
          148
                 21.462128
          149
                 22.929389
          Length: 150, dtype: float64
 In [7]:
           t_value, p_two, degree_free
          (-16.019643876306674, 1.0, 162.29643381428605)
 In [ ]:
 In [ ]:
 In [ ]:
In [295...
           stats.t.ppf(q=0.975, df=25)
          2.059538552753294
Out[295...
In [297...
           stats.t.ppf(q=0.95, df=200)
Out[297... 1.652508100910269
In [255...
           # caculate confidence internal right_tail =0.025 two_tail=0.05
           t_ci = 1.971
```

```
# sample number
           control num = df 21.shape[0]
          treat_num = df_22.shape[0]
In [256...
          control_num
Out[256... 150
In [257...
          control_std = df_21.std()
          treat_std = df_22.std()
          se = np.sqrt( np.square(control_std)/control_num + np.square(treat_std)/treat_num)
           sample_mean = df_21.mean() - df_22.mean()
           sample_mean
          upper_bound = sample_mean - t_ci * se
           lower bound = sample mean + t ci * se
In [258...
          upper_bound, lower_bound
         (-0.03298816640548901, -0.022860283394956115)
Out[258...
 In [ ]:
 In [ ]:
In [133...
          # caculate confidence internal right tail =0.025 two tail=0.05
          t_ci = 1.971
          # sample number
          control num = df control.shape[0]
          treat_num = df_treat.shape[0]
In [134...
           control_std = df_control['active_mins'].std()
In [135...
          treat_std = df_treat['active_mins'].std()
In [136...
          se = np.sqrt( np.square(control_std)/control_num + np.square(treat_std)/treat_num)
In [141...
          df_control['active_mins'].mean(), df_treat['active_mins'].mean()
Out[141... (458.2211623246493, 458.4024761077324)
In [137...
           sample_mean = df_control['active_mins'].mean() - df_treat['active_mins'].mean()
           sample_mean
```

```
Out[137... -0.1813137830831124
In [138...
           upper_bound = sample_mean - t_ci * se
           lower_bound = sample_mean + t_ci * se
In [139...
           upper bound
Out[139...
         -38.591763263087756
In [140...
           lower bound
Out[140... 38.22913569692153
 In [ ]:
In [210...
          # caculate confidence internal right tail =0.025 two tail=0.05
          t_ci = 1.971
          t_ci = 2.60
          # sample number
          control num = df control.shape[0]/4
          treat_num = df_treat.shape[0]
          print(control_num, treat_num)
           control_std = (df_control['active_mins']/4).std()
          treat std = df treat['active mins'].std()
          se = np.sqrt( np.square(control_std)/control_num + np.square(treat_std)/treat_num)
           (df_control['active_mins']/4).mean(), df_treat['active_mins'].mean()
           sample_mean = (df_control['active_mins']/4).mean() - df_treat['active_mins'].mean()
           sample_mean
          upper_bound = sample_mean - t_ci * se
          lower_bound = sample_mean + t_ci * se
          upper_bound
          lower bound
          37.5 150
         2021.5178573516198
Out[210...
In [211...
          upper_bound
          -1138.0278573516218
Out[211...
In [212...
          1639.3343468615546
Out[212... 1639.3343468615546
```

```
In [205... (df_control['active_mins']/4).mean()
          28581.545
Out[205...
In [206...
           df_treat['active_mins'].mean()
Out[206...
          28139.8
In [213...
          607.6049451352387
Out[213...
In [214...
           control_std
          2650.8065794906283
Out[214...
In [215...
           (df_control['active_mins']/4).std()
          2650.8065794906283
Out[215...
In [216...
           (df_control['active_mins']).std()
          10603.226317962513
Out[216...
In [118...
           t_value, p_two, degree_free
          (-0.009303964709989936, 0.9925767506273644, 13920.469441655145)
Out[118...
 In [ ]:
 In [ ]:
In [573...
           df_treat.shape
          (9208, 2)
Out[573...
In [20]:
           p_value
          3.0324185230356597e-35
Out[20]:
In [19]:
           stats.ttest_ind_from_stats(mean1=df_control.mean()
                                        , std1=df_control.std()
                                        , nobs1=150
                                        , mean2=df_treat.mean()
                                        , std2=df_treat.std()
```

```
, nobs2=150
                                                                                                            , equal_var=False)
Out[19]: Ttest_indResult(statistic=-16.019643876306688, pvalue=3.03241852303566e-35)
In [15]:
                              import numpy as np
                             from scipy.stats import ttest_ind
                              from scipy.stats import t
                              import pandas as pd
                             def welch ttest(x1, x2,alternative):
                                         n1 = x1.size
                                         n2 = x2.size
                                         m1 = np.mean(x1)
                                         m2 = np.mean(x2)
                                         v1 = np.var(x1, ddof=1)
                                         v2 = np.var(x2, ddof=1)
                                         pooled se = np.sqrt(v1 / n1 + v2 / n2)
                                         delta = m1-m2
                                         tstat = delta / pooled se
                                         df = (v1 / n1 + v2 / n2)**2 / (v1**2 / (n1**2 * (n1 - 1)) + v2**2 / (n2**2 * (n2 - 1)) + v2**2 / (n2 - 1) + 
                                         # two side t-test
                                         p = 2 * t.cdf(-abs(tstat), df)
                                         # upper and Lower bounds
                                         lb = delta - t.ppf(0.975,df)*pooled_se
                                         ub = delta + t.ppf(0.975,df)*pooled se
                                         return pd.DataFrame(np.array([tstat,df,p,delta,lb,ub]).reshape(1,-1),
                                                                                                      columns=['T statistic','df','pvalue 2 sided','Difference in me
In [16]:
                             welch ttest(df control,df treat,"equal")
Out[16]:
                                      T statistic
                                                                                    df pvalue 2 sided Difference in mean
                                                                                                                                                                                                    lb
                                                                                                                                                                                                                             ub
                           0 -16.019644 162.296434
                                                                                                 3.032419e-35
                                                                                                                                                          -4.087583 -4.591446 -3.583721
In [17]:
                              upper_bound
                            -4.509893654334767
Out[17]:
In [18]:
                              lower_bound
Out[18]: -3.665273229338887
   In [ ]:
```

```
stats.kstest(df_61['active_mins'], 'norm')
In [487...
Out[487... KstestResult(statistic=0.9616591299099563, pvalue=0.0)
In [492...
           df 52.head(2)
Out[492...
                    uid
                                dt active_mins
          533200 23941 2019-02-20
                                           1.0
          264373 11922 2019-04-09
                                           1.0
In [493...
           df_62 = df_52.groupby(['dt'], as_index=False)['active_mins'].sum()
In [494...
           stats.kstest(df_62['active_mins'], 'norm')
Out[494...
          KstestResult(statistic=1.0, pvalue=0.0)
In [331...
           st.kstest(df_42, 'norm')
Out[331... KstestResult(statistic=0.8678963361749233, pvalue=0.0)
In [134...
           st.shapiro(dist_10)
          ShapiroResult(statistic=0.9979808926582336, pvalue=0.2747354805469513)
Out[134...
In [129...
           st.shapiro(df_11)
Out[129... ShapiroResult(statistic=0.0052002668380737305, pvalue=0.0)
 In [ ]:
 In [ ]:
 In [ ]:
```