

02_ttestsAssignemnt

April 1, 2022

```
[ ]: # one sample t-test
# import libraries

import seaborn as sns
import pandas as pd
from scipy.stats import ttest_1samp

# load dataset

df = sns.load_dataset('penguins')
df
```

```
[ ]:   species    island  bill_length_mm  bill_depth_mm  flipper_length_mm  \
0   Adelie  Torgersen         39.1           18.7             181.0
1   Adelie  Torgersen         39.5           17.4             186.0
2   Adelie  Torgersen         40.3           18.0             195.0
3   Adelie  Torgersen          NaN           NaN              NaN
4   Adelie  Torgersen         36.7           19.3             193.0
..   ...      ...
339  Gentoo   Biscoe          NaN           NaN              NaN
340  Gentoo   Biscoe         46.8           14.3             215.0
341  Gentoo   Biscoe         50.4           15.7             222.0
342  Gentoo   Biscoe         45.2           14.8             212.0
343  Gentoo   Biscoe         49.9           16.1             213.0

      body_mass_g    sex
0         3750.0  Male
1         3800.0 Female
2         3250.0 Female
3           NaN   NaN
4         3450.0 Female
..          ...    ...
339         NaN   NaN
340        4850.0 Female
341        5750.0  Male
342        5200.0 Female
343        5400.0  Male
```

[344 rows x 7 columns]

```
[ ]: df1 = df[['species', 'bill_length_mm', 'bill_depth_mm']]
df1.head()
```

```
[ ]:   species  bill_length_mm  bill_depth_mm
0  Adelie         39.1         18.7
1  Adelie         39.5         17.4
2  Adelie         40.3         18.0
3  Adelie          NaN          NaN
4  Adelie         36.7         19.3
```

```
[ ]: df1.describe()
```

```
[ ]:   bill_length_mm  bill_depth_mm
count      342.000000      342.000000
mean       43.921930      17.151170
std         5.459584       1.974793
min        32.100000      13.100000
25%        39.225000      15.600000
50%        44.450000      17.300000
75%        48.500000      18.700000
max        59.600000      21.500000
```

```
[ ]: # check the age and compare witht a known value of 45 years
```

```
ttest_1samp(df1['bill_length_mm'], 50)

stat, p = ttest_1samp(df1['bill_length_mm'], 50)

print('stat=%.3f, p=%.3f' % (stat, p))

# make a conditional arguement for ease
if p > 0.05:
    print('Probably the same distribution')
else:
    print('Probably different Distribution')
```

```
stat=nan, p=nan
Probably different Distribution
```

```
[ ]: # we will compare
```

```
#splitting dataset
df_G= df1.loc[df1['species']== 'Gentoo']
df_A= df1.loc[df1['species']== 'Adeli']
```

```
# library
from scipy.stats import ttest_ind
stat, p = ttest_ind(df_G['bill_length_mm'], df_A['bill_length_mm'])

print('stat=%.3f, p=%.3f' % (stat, p))

# make a conditional argument for ease
if p > 0.05:
    print('Probably the same distribution')
else:
    print('Probably different Distribution')
```

```
stat=nan, p=nan
Probably different Distribution
```

```
[ ]: df_A.describe
```

```
[ ]: <bound method NDFrame.describe of Empty DataFrame
Columns: [species, bill_length_mm, bill_depth_mm]
Index: []>
```

```
[ ]: df_G.info
```

```
[ ]: <bound method DataFrame.info of
```

	species	bill_length_mm	bill_depth_mm
220	Gentoo	46.1	13.2
221	Gentoo	50.0	16.3
222	Gentoo	48.7	14.1
223	Gentoo	50.0	15.2
224	Gentoo	47.6	14.5
..
339	Gentoo	NaN	NaN
340	Gentoo	46.8	14.3
341	Gentoo	50.4	15.7
342	Gentoo	45.2	14.8
343	Gentoo	49.9	16.1

```
[124 rows x 3 columns]>
```

```
[ ]: df_G= df1.loc[df1['species']=='Gentoo']
df_G.head()
```

```
[ ]:
```

	species	bill_length_mm	bill_depth_mm
220	Gentoo	46.1	13.2
221	Gentoo	50.0	16.3
222	Gentoo	48.7	14.1
223	Gentoo	50.0	15.2

224	Gentoo	47.6	14.5
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