

Statistics I: Technology help — Python

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Description: Python code for an introductory applied statistics course. Work in progress.

Module 1: Data and decisions

The `pandas` library is one of the workhorses for data analysis:

```
In [1]: import pandas as pd      # imports the pandas library and assigns it the alias pd
```

The command `dir(pd)` will list all of the attributes of the `pandas` library. To get detailed information on an attribute of a library (for instance, on the `read_csv` attribute), use:

`help(pd.read_csv)`. Alternatively, enter `pandas` & documentation in a search engine to get to the on-line help (<https://pandas.pydata.org/docs/>).

To import a text file with data in Python, you first have to find the path to the file. The data file `students.csv` used in the example is stored on my web site: <https://luc-hens.github.io/students.csv>. Usually, you will have your data file stored on your computer. In that case you have to give the path to that file, which looks something like: `/Users/luchens/Documents/Data/students.csv` (This is for macOS; possibly Windows uses backslashes: `\` (check)).

It is good practice to not have spaces in a csv file. The `pandas` command `read_csv` has an option `skipinitialspace=True` ("Skip spaces after delimiter") but that does not solve all the problems caused by blank spaces in a csv file.

```
In [2]: import pandas as pd
df = pd.read_csv('https://luc-hens.github.io/students.csv', sep=',', skipinitialspace=True)
print(df)
display(df)
```

	case	sex	height	weight	major
0	1	Female	172	63	Business
1	2	Female	170	70	International Affairs
2	3	Female	170	52	Other
3	4	Female	171	52	Communications
4	5	Male	186	90	Business
5	6	Male	183	79	Business
6	7	Male	170	66	Communications
7	8	Female	169	56	Business
8	9	Male	175	75	International Affairs
9	10	Female	175	65	Communications
10	11	Male	195	94	Business
11	12	Female	176	51	International Affairs
12	13	Male	188	76	International Affairs

13	14	Male	192	82	Business
14	15	Male	172	70	International Affairs
15	16	Female	169	53	Business
16	17	Female	172	52	International Affairs
17	18	Male	178	85	Business
18	19	Female	177	59	Communications
19	20	Male	178	72	International Affairs
20	21	Female	160	54	Business
21	22	Female	175	54	International Affairs
22	23	Male	190	70	International Affairs
23	24	Male	178	85	Business
24	25	Female	163	55	Business
25	26	Female	161	59	Business
26	27	Female	162	44	Communications
27	28	Female	170	54	Business
28	29	Female	154	52	Business
29	30	Female	170	65	Business

	case	sex	height	weight	major
0	1	Female	172	63	Business
1	2	Female	170	70	International Affairs
2	3	Female	170	52	Other
3	4	Female	171	52	Communications
4	5	Male	186	90	Business
5	6	Male	183	79	Business
6	7	Male	170	66	Communications
7	8	Female	169	56	Business
8	9	Male	175	75	International Affairs
9	10	Female	175	65	Communications
10	11	Male	195	94	Business
11	12	Female	176	51	International Affairs
12	13	Male	188	76	International Affairs
13	14	Male	192	82	Business
14	15	Male	172	70	International Affairs
15	16	Female	169	53	Business
16	17	Female	172	52	International Affairs
17	18	Male	178	85	Business
18	19	Female	177	59	Communications
19	20	Male	178	72	International Affairs
20	21	Female	160	54	Business
21	22	Female	175	54	International Affairs
22	23	Male	190	70	International Affairs
23	24	Male	178	85	Business
24	25	Female	163	55	Business
25	26	Female	161	59	Business
26	27	Female	162	44	Communications
27	28	Female	170	54	Business
28	29	Female	154	52	Business
29	30	Female	170	65	Business

To display just the first couple of lines of the data frame called df:

In [3]: `df.head()`

Out[3]:

	case	sex	height	weight	major
0	1	Female	172	63	Business
1	2	Female	170	70	International Affairs
2	3	Female	170	52	Other
3	4	Female	171	52	Communications
4	5	Male	186	90	Business

To display the last couple of lines of the data frame called df:

In [4]:

```
df.tail()
```

Out[4]:

	case	sex	height	weight	major
25	26	Female	161	59	Business
26	27	Female	162	44	Communications
27	28	Female	170	54	Business
28	29	Female	154	52	Business
29	30	Female	170	65	Business

To display the 10 first lines of the data frame:

In [5]:

```
df.head(10)
```

Out[5]:

	case	sex	height	weight	major
0	1	Female	172	63	Business
1	2	Female	170	70	International Affairs
2	3	Female	170	52	Other
3	4	Female	171	52	Communications
4	5	Male	186	90	Business
5	6	Male	183	79	Business
6	7	Male	170	66	Communications
7	8	Female	169	56	Business
8	9	Male	175	75	International Affairs
9	10	Female	175	65	Communications

Show the column names (variable names) (this is useful to check whether there are no blank spaces in the variable names):

In [6]:

```
list(df)
```

Out[6]: ['case', 'sex', 'height', 'weight', 'major']

Inspect the data types in the dataframe called df:

```
In [7]: df.dtypes
```

```
Out[7]: case      int64  
sex      object  
height   int64  
weight   int64  
major    object  
dtype: object
```

To display just one of the variables:

```
In [8]: print(df.height)
```

```
0      172  
1      170  
2      170  
3      171  
4      186  
5      183  
6      170  
7      169  
8      175  
9      175  
10     195  
11     176  
12     188  
13     192  
14     172  
15     169  
16     172  
17     178  
18     177  
19     178  
20     160  
21     175  
22     190  
23     178  
24     163  
25     161  
26     162  
27     170  
28     154  
29     170  
Name: height, dtype: int64
```

```
In [ ]:
```

Module 2: Displaying and describing categorical data

Bar chart and pie chart

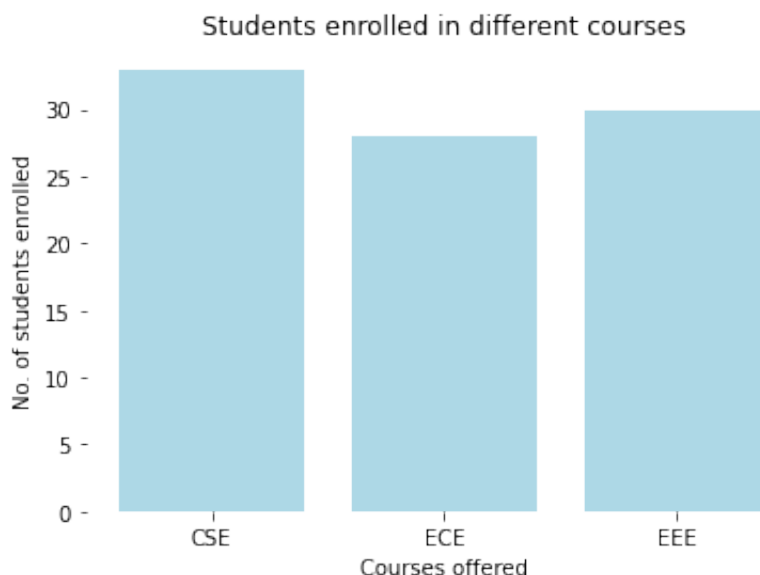
Generate a **bar chart** showing enrolment in three classes with course codes CSE (33 students), ECE (28 students), EEE (30 students):

(Documentation: https://matplotlib.org/stable/api/_as_gen/matplotlib.pyplot.bar.html)

(Example is taken from <https://www.analyticsvidhya.com/blog/2021/08/understanding-bar-plots-in-python-beginners-guide-to-data-visualization/>)

In [9]:

```
import numpy as np
import matplotlib.pyplot as plt
# Dataset generation
data_dict = {'CSE':33, 'ECE':28, 'EEE':30}
courses = list(data_dict.keys())
values = list(data_dict.values())
fig = plt.figure() # add option `figsize = (10, 5)` to control size
# Bar plot
plt.box(False) # get rid of the box
plt.bar(courses, values, color='lightblue')
plt.xlabel("Courses offered")
plt.ylabel("No. of students enrolled")
plt.title("Students enrolled in different courses")
plt.show()
```

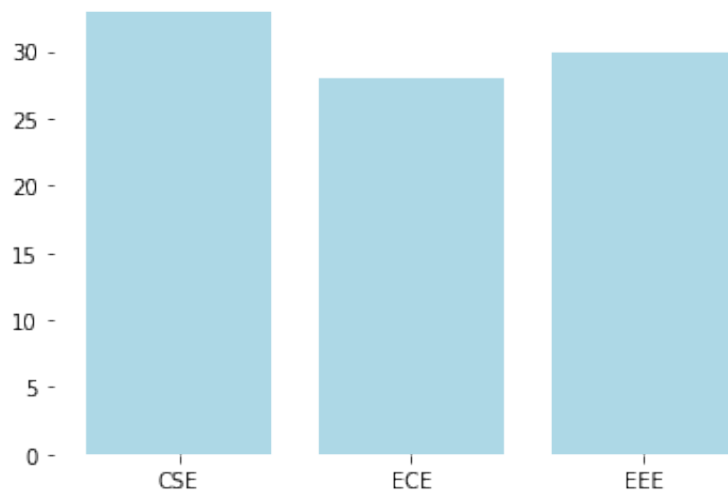


Save a plot to a file (.png or .pdf):

In [10]:

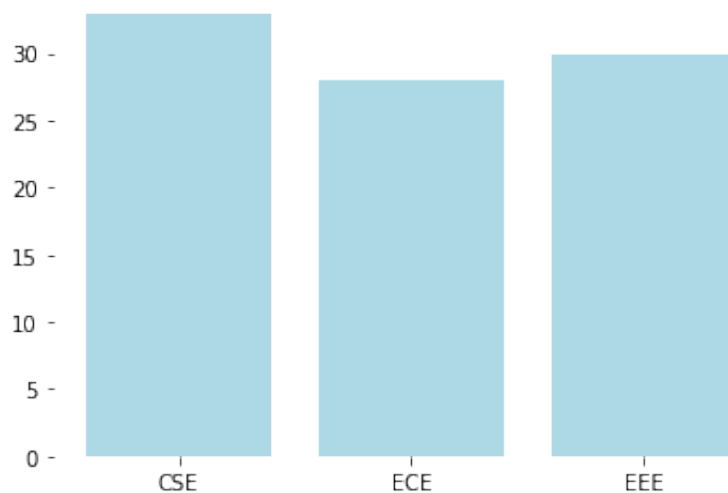
```
fig = plt.figure()
plt.box(False) # get rid of the box
plt.bar(courses, values, color='lightblue')
fig.savefig('saved_figure-1000dpi.png', dpi = 1000, transparent=True)
# the plot is saved to the current working directory (cwd)
# to find out what the current working directory (cwd) is:
import os
print('The file is saved to the current working directory: ',os.getcwd())
```

The file is saved to the current working directory: /Users/luchens/Documents/jupyter-notebooks



To save the plot as a .pdf:

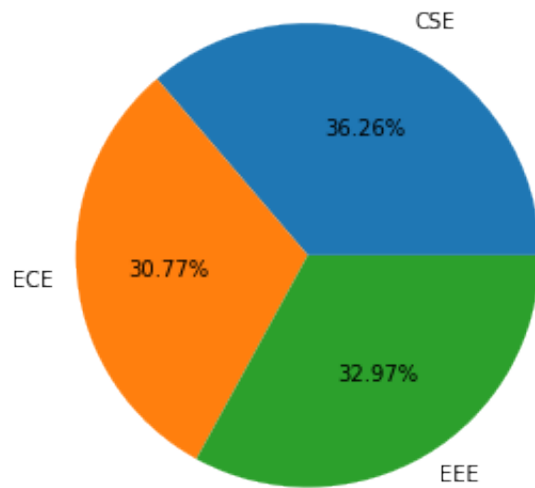
```
In [11]: fig = plt.figure()
plt.box(False)           # get rid of the box
plt.bar(courses, values, color='lightblue')
fig.savefig('saved_figure-1000dpi.pdf', dpi = 1000, transparent=True)
```



Generate a **pie chart** for the same data (pie charts are usually a poor way to display data):

(documentation: https://matplotlib.org/stable/api/_as_gen/matplotlib.pyplot.pie.html)

```
In [12]: import numpy as np
import matplotlib.pyplot as plt
# Dataset generation
data_dict = {'CSE':33, 'ECE':28, 'EEE':30}
courses = list(data_dict.keys())
values = list(data_dict.values())
fig = plt.figure(figsize = (10, 5))
# Pie chart:
plt.pie(values, labels=courses, autopct='%1.2f%%') # option: autopct to sl
plt.show()
```



Generate a bar chart and a pie chart from an imported data set (students.csv):

Use `value_counts()` to count the how many times each of the values of the variable 'sex' in the dataframe 'df' occurs:

```
In [13]: a = df.sex.value_counts()
         print(a)
```

```
Female    18
Male      12
Name: sex, dtype: int64
```

```
In [14]: a.values
```

```
Out[14]: array([18, 12])
```

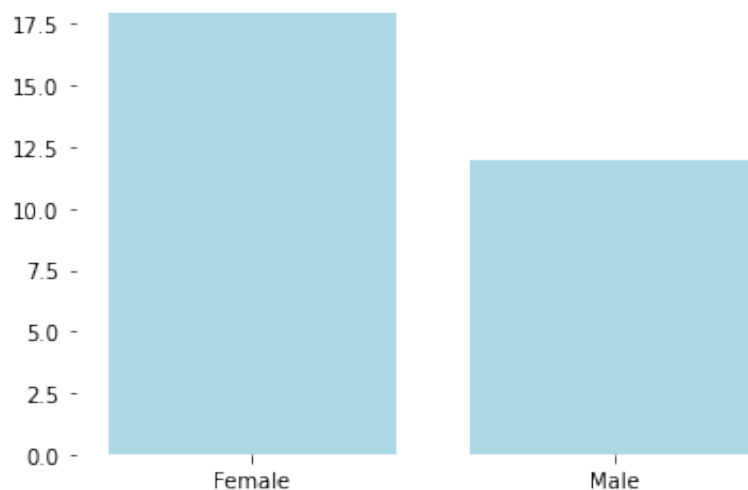
```
In [15]: a.index
```

```
Out[15]: Index(['Female', 'Male'], dtype='object')
```

Then use `a` as input for the `bar()` function:

```
In [16]: plt.box(False)                                # get rid of the box
         plt.bar(a.index, a.values, color='lightblue')
```

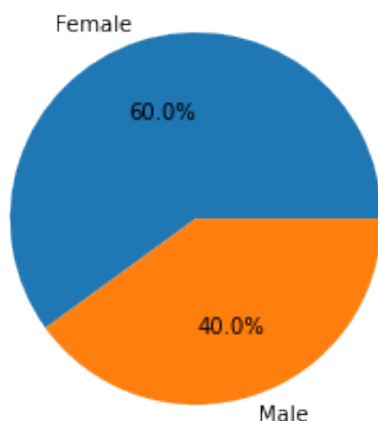

Out[16]: <BarContainer object of 2 artists>



Or use `a` as input for the `pie()` function:

```
In [17]: plt.pie(a.values, labels=a.index, autopct='%1.1f%%')
```

```
Out[17]: ([<matplotlib.patches.Wedge at 0x7f7a29d107c0>,
<matplotlib.patches.Wedge at 0x7f7a29d10f40>],
[Text(-0.3399187721714582, 1.046162142464278, 'Female'),
Text(0.3399188701202255, -1.0461621106387813, 'Male')],
[Text(-0.18541023936624992, 0.5706338958896061, '60.0%'),
Text(0.18541029279285026, -0.5706338785302443, '40.0%')])
```



Contingency table

To create a contingency table use `pandas.crosstab(index, columns)`. For an example see: <https://www.statology.org/contingency-table-python/> Here is how to create a contingency table for the categorical variables (sex, major) from the students data file stored in the dataframe called `df`:

```
In [18]: import pandas as pd
pd.crosstab(index=df['major'], columns=df['sex'], margins=True)
```

Out[18]:

	sex	Female	Male	All
major				
Business		9	6	15
Communications		4	1	5
International Affairs		4	5	9
Other		1	0	1
All		18	12	30

To express all frequencies as relative frequencies, divide by number of observations:

```
In [19]: n = len(df)      # number of observations
pd.crosstab(index=df['major'], columns=df['sex'], margins=True)/n
```

Out[19]:

	sex	Female	Male	All
major				
Business		0.300000	0.200000	0.500000
Communications		0.133333	0.033333	0.166667
International Affairs		0.133333	0.166667	0.300000
Other		0.033333	0.000000	0.033333
All		0.600000	0.400000	1.000000

... and multiply by 100 (percent) to get percentages:

```
In [20]: 100*pd.crosstab(index=df['major'], columns=df['sex'], margins=True)/n
```

Out[20]:

	sex	Female	Male	All
major				
Business		30.000000	20.000000	50.000000
Communications		13.333333	3.333333	16.666667
International Affairs		13.333333	16.666667	30.000000
Other		3.333333	0.000000	3.333333
All		60.000000	40.000000	100.000000

Summary statistics (qualitative variables only):

```
In [21]: df.describe(include=['object'])
```

```
Out[21]:
```

	sex	major
count	30	30
unique	2	4
top	Female	Business
freq	18	15

Summary statistics of one categorical variable (in this case: sex)

```
In [22]: df['sex'].describe()
```

```
Out[22]: count          30
unique          2
top            Female
freq           18
Name: sex, dtype: object
```

Module 3: Displaying and describing quantitative variables

Descriptive statistics (mean, median, standard deviation,...)

Summary statistics (all variables) using the `describe()` function from pandas:

```
In [23]: df.describe(include='all')
```

```
Out[23]:
```

	case	sex	height	weight	major
count	30.000000	30	30.000000	30.000000	30
unique	NaN	2	NaN	NaN	4
top	NaN	Female	NaN	NaN	Business
freq	NaN	18	NaN	NaN	15
mean	15.500000	NaN	174.033333	65.133333	NaN
std	8.803408	NaN	9.625696	13.356474	NaN
min	1.000000	NaN	154.000000	44.000000	NaN
25%	8.250000	NaN	170.000000	54.000000	NaN
50%	15.500000	NaN	172.000000	64.000000	NaN
75%	22.750000	NaN	178.000000	74.250000	NaN
max	30.000000	NaN	195.000000	94.000000	NaN

Summary statistics of one variable (in this case: height) using the `describe()` function from pandas

```
In [24]: df['height'].describe()
```

```
Out[24]: count      30.000000
mean       174.033333
std        9.625696
min        154.000000
25%        170.000000
50%        172.000000
75%        178.000000
max        195.000000
Name: height, dtype: float64
```

Summary statistics (quantitative variables only) using the `describe()` function from pandas:

```
In [25]: df.describe()
```

```
Out[25]:
```

	case	height	weight
count	30.000000	30.000000	30.000000
mean	15.500000	174.033333	65.133333
std	8.803408	9.625696	13.356474
min	1.000000	154.000000	44.000000
25%	8.250000	170.000000	54.000000
50%	15.500000	172.000000	64.000000
75%	22.750000	178.000000	74.250000
max	30.000000	195.000000	94.000000

Mean and standard deviation of one of the variables ('height') from the 'df' dataframe:

```
In [26]: import pandas as pd
df.height.mean()                                # height.mean() does not work: NameError
```

```
Out[26]: 174.03333333333333
```

```
In [27]: df.height.std()
```

```
Out[27]: 9.625695974240289
```

Other descriptive statistics:

```
In [28]: df.height.median()    # similarly: min() ; max() ; sum() ; count() ; quantile()
```

```
Out[28]: 172.0
```

```
In [29]: df.height.quantile(q=0.50)    # the 50th percentile is the same as the median
```

```
Out[29]: 172.0
```

```
In [30]: df.height.quantile(q=0.25) # the 25th percentile (the first quartile)
```

```
Out[30]: 170.0
```

```
In [31]: df.height.quantile(q=0.75) # the 75th percentile (the third quartile)
```

```
Out[31]: 178.0
```

```
In [32]: df.height.quantile(q=0.75)-df.height.quantile(q=0.25) # the interquartile range
```

```
Out[32]: 8.0
```

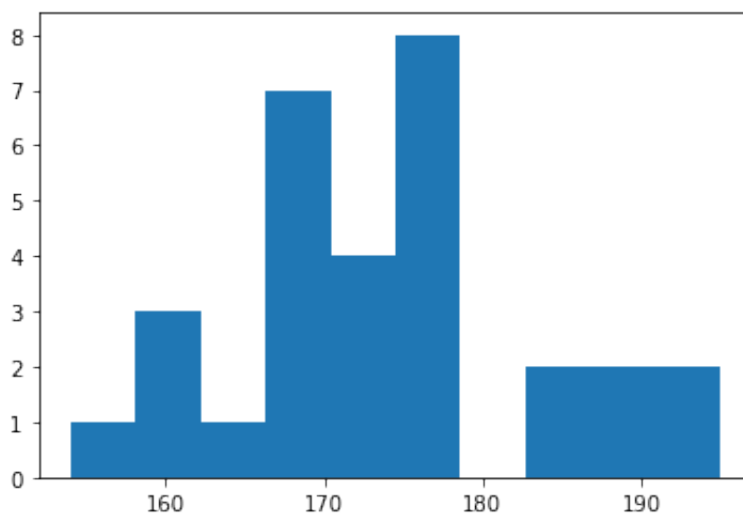
Histogram

To draw a **histogram** use `hist()` from the `matplotlib.pyplot` library.

Frequency histogram (vertical axis shows counts, absolute frequencies):

```
In [33]: import matplotlib.pyplot as plt
plt.hist(df.height) # the default is a frequency histogram: vertical axis is counts
```

```
Out[33]: (array([1., 3., 1., 7., 4., 8., 0., 2., 2., 2.]),
array([154. , 158.1, 162.2, 166.3, 170.4, 174.5, 178.6, 182.7, 186.8,
       190.9, 195. ]),
<BarContainer object of 10 artists>)
```



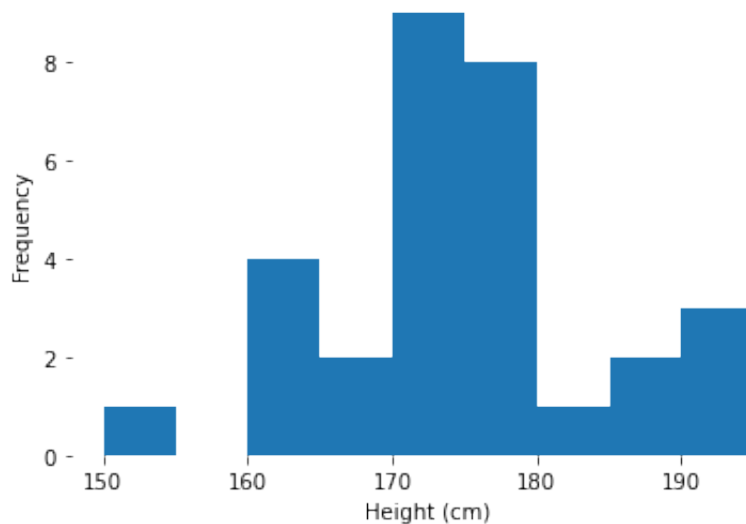
In the output, the first array gives the counts (absolute frequencies) for each of the classes (bins). The second array gives the edges of the bins (

https://matplotlib.org/stable/api/_as_gen/matplotlib.pyplot.hist.html).

Get rid of the box, add labels to the axes, and let the bins start at 150, 155, 160,...:

```
In [34]: import matplotlib.pyplot as plt
plt.box(False) # get rid of the box
plt.xlabel('Height (cm)') # add label on x-axis
plt.ylabel('Frequency') # add label on y-axis
plt.hist(df.height, bins=[150, 155, 160, 165, 170, 175, 180, 185, 190, 195])
```

```
Out[34]: (array([1., 0., 4., 2., 9., 8., 1., 2., 3.]),
          array([150, 155, 160, 165, 170, 175, 180, 185, 190, 195]),
          <BarContainer object of 9 artists>)
```



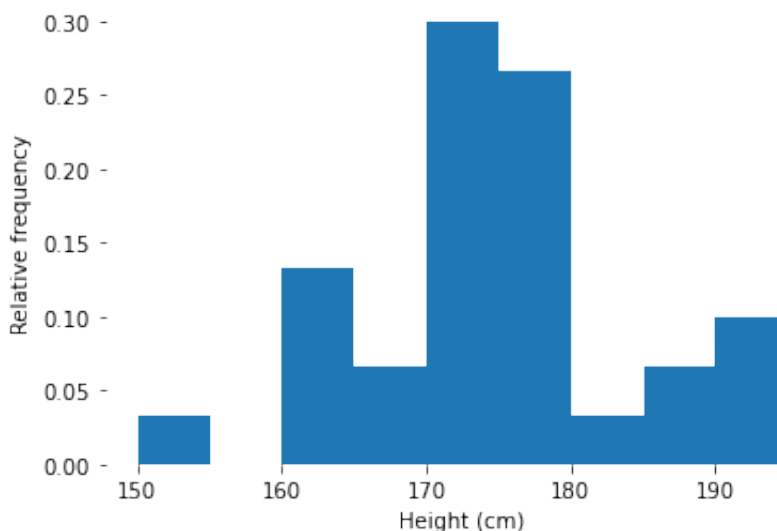
Relative frequency histogram (vertical axis shows relative frequencies):

```
In [35]: df.height.size # size gives the number of observations of the variable height
```

```
Out[35]: 30
```

```
In [36]: import matplotlib.pyplot as plt
import numpy as np
plt.box(False) # get rid of the box
plt.xlabel('Height (cm)') # add label on x-axis
plt.ylabel('Relative frequency') # add label on y-axis
plt.hist(df.height, weights=np.zeros_like(df.height) + 1. / df.height.size,
         bins=[150,155,160,165,170,175,180,185,190,195])
```

```
Out[36]: (array([0.03333333, 0.13333333, 0.06666667, 0.3, 0.26666667, 0.03333333, 0.06666667, 0.1, 0.1]),
          array([150, 155, 160, 165, 170, 175, 180, 185, 190, 195]),
          <BarContainer object of 9 artists>)
```



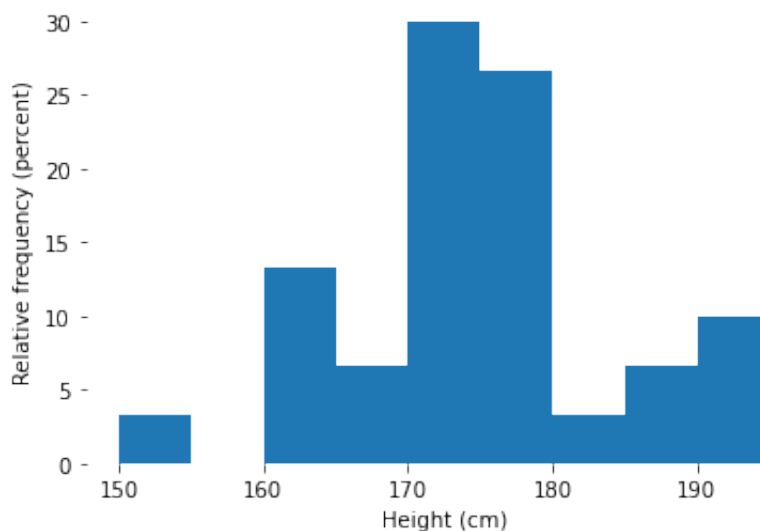
In the output, the first array gives the relative frequencies for each of the classes (bins). The second array gives the edges of the bins (

https://matplotlib.org/stable/api/_as_gen/matplotlib.pyplot.hist.html).

To get a relative frequency histogram with relative frequencies expressed as percentages, multiply the weights by 100:

```
In [37]: import matplotlib.pyplot as plt
import numpy as np
df.height.size # size gives the number of observations of the variable height
plt.box(False) # get rid of the box
plt.xlabel('Height (cm)') # add label on x-axis
plt.ylabel('Relative frequency (percent)') # add label on y-axis
plt.hist(df.height, weights=100*(np.zeros_like(df.height) + 1. / df.height.size),
        bins=[150,155,160,165,170,175,180,185,190,195])
```

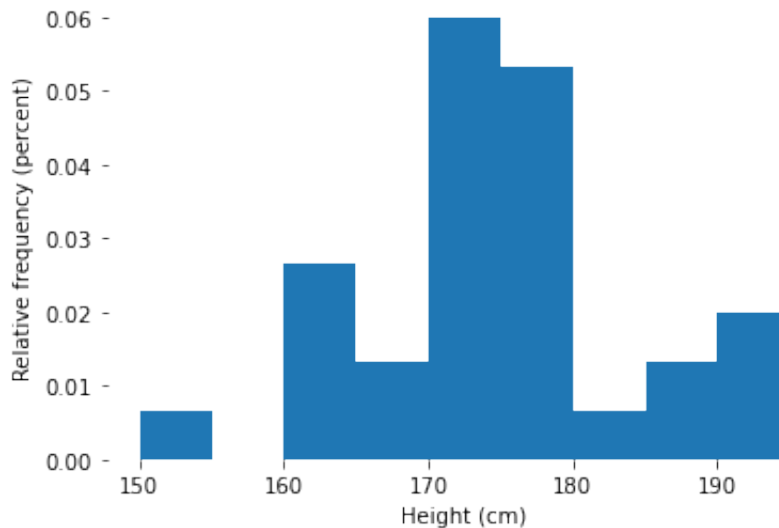
```
Out[37]: (array([ 3.33333333,  0.          , 13.33333333,  6.66666667, 30.          ,
        26.66666667,  3.33333333,  6.66666667, 10.          ]),
array([150, 155, 160, 165, 170, 175, 180, 185, 190, 195]),
<BarContainer object of 9 artists>)
```



Density histogram (vertical axis shows densities):

```
In [38]: import matplotlib.pyplot as plt
df.height.size # size gives the number of observations of the variable height
plt.box(False) # get rid of the box
plt.xlabel('Height (cm)') # add label on x-axis
plt.ylabel('Relative frequency (percent)') # add label on y-axis
plt.hist(df.height, density=True, bins=[150,155,160,165,170,175,180,185,190,195])
```

```
Out[38]: (array([0.00666667, 0.00666667, 0.02666667, 0.01333333, 0.06666667, 0.05333333, 0.00666667, 0.01333333, 0.02666667]),
          array([150, 155, 160, 165, 170, 175, 180, 185, 190, 195]),
          <BarContainer object of 9 artists>)
```



In the output, the first array gives the densities for each of the classes (bins). The second array gives the edges of the bins (

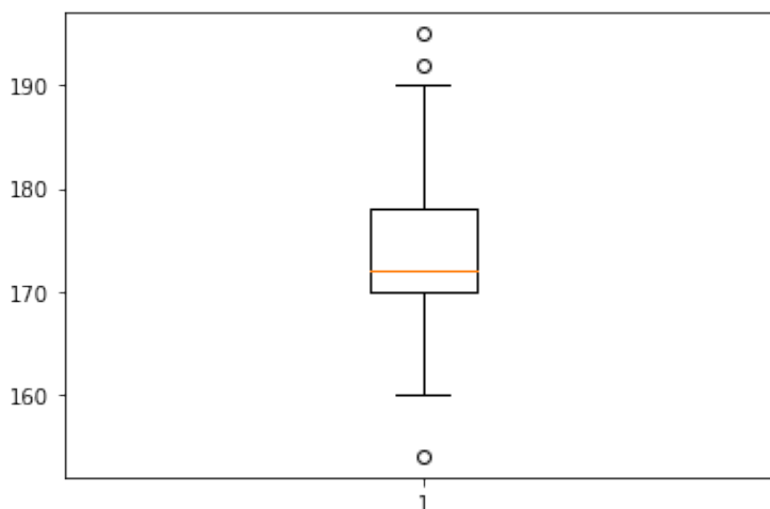
https://matplotlib.org/stable/api/_as_gen/matplotlib.pyplot.hist.html).

Box plot

Use `boxplot()` from `matplotlib`:

```
In [39]: import matplotlib.pyplot as plt
         plt.boxplot(df.height)
```

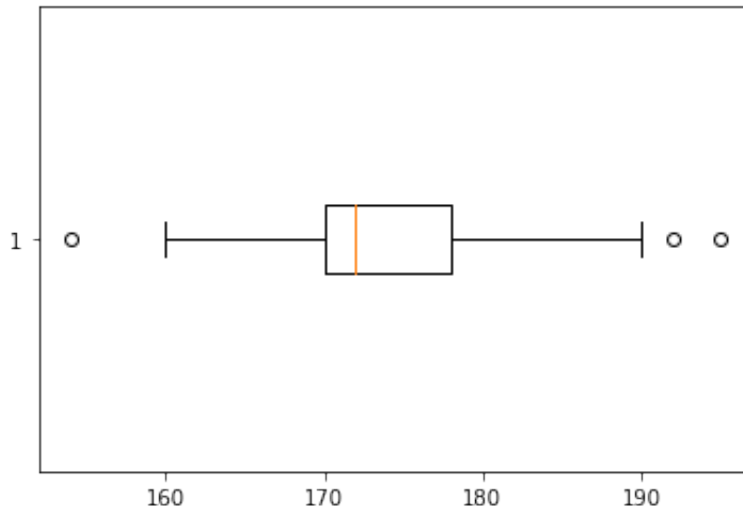
```
Out[39]: {'whiskers': [<matplotlib.lines.Line2D at 0x7f7a485d3ac0>,
                     <matplotlib.lines.Line2D at 0x7f7a485d3d90>],
          'caps': [<matplotlib.lines.Line2D at 0x7f7a389b3160>,
                  <matplotlib.lines.Line2D at 0x7f7a389b34c0>],
          'boxes': [<matplotlib.lines.Line2D at 0x7f7a485d3760>],
          'medians': [<matplotlib.lines.Line2D at 0x7f7a389b3850>],
          'fliers': [<matplotlib.lines.Line2D at 0x7f7a389b3be0>],
          'means': []}
```



Rotate the boxplot to get a horizontal orientation:


```
In [40]: import matplotlib.pyplot as plt
plt.boxplot(df.height,vert=False)
```

```
Out[40]: {'whiskers': [<matplotlib.lines.Line2D at 0x7f7a29e19a90>,
<matplotlib.lines.Line2D at 0x7f7a29e19e20>],
'caps': [<matplotlib.lines.Line2D at 0x7f7a29e271f0>,
<matplotlib.lines.Line2D at 0x7f7a29e27580>],
'boxes': [<matplotlib.lines.Line2D at 0x7f7a29e19700>],
'medians': [<matplotlib.lines.Line2D at 0x7f7a29e278b0>],
'fliers': [<matplotlib.lines.Line2D at 0x7f7a29e27c40>],
'means': []}
```

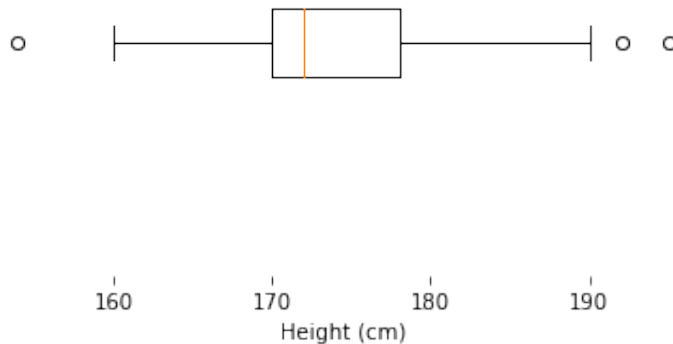


Make the box plot prettier (get rid of the box, label the axis, add a title, get rid of the "1" tick):

```
In [41]: import matplotlib.pyplot as plt
plt.box(False) # get rid of the box
plt.title("Box plot of the heights of 30 students") # add title
plt.xlabel("Height (cm)") # add label to x-axis
plt.ylabel("")
plt.tick_params( # this block removes the "1" tick
    axis='y', # changes apply to the y-axis
    which='both', # both major and minor ticks are affected
    left=False, # ticks along the left edge are off (for x axis t
    right=False, # ticks along the right edge are off (for x axis t
    labelleft=False) # labels along the left edge are off (for x axis t
plt.boxplot(df.height,vert=False)
```

```
Out[41]: {'whiskers': [<matplotlib.lines.Line2D at 0x7f7a599e4fa0>,
<matplotlib.lines.Line2D at 0x7f7a5978c310>],
'caps': [<matplotlib.lines.Line2D at 0x7f7a5978c6a0>,
<matplotlib.lines.Line2D at 0x7f7a5978ca30>],
'boxes': [<matplotlib.lines.Line2D at 0x7f7a599e4c10>],
'medians': [<matplotlib.lines.Line2D at 0x7f7a5978cdc0>],
'fliers': [<matplotlib.lines.Line2D at 0x7f7a5979b190>],
'means': []}
```

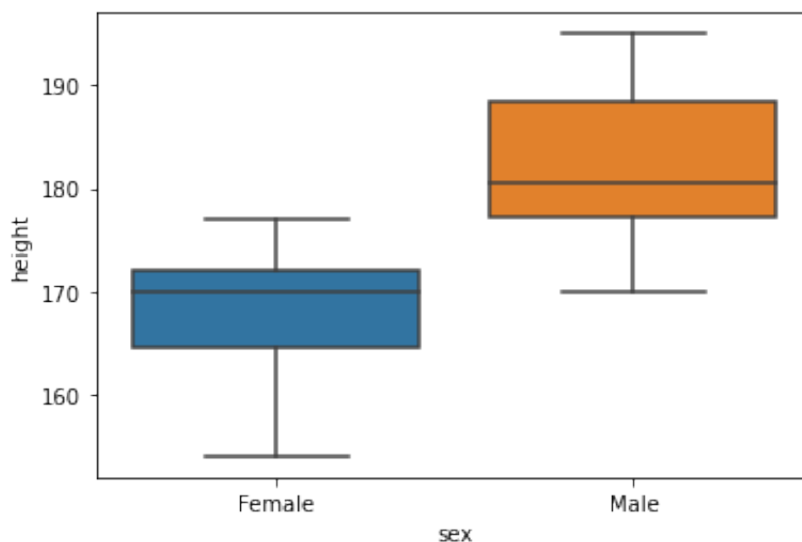
Box plot of the heights of 30 students



To compare the heights of men and women, use **side-by-side boxplots** (from the seaborn library):

```
In [42]: import seaborn as sns
sns.boxplot(data=df, x="sex", y='height')
```

```
Out[42]: <AxesSubplot:xlabel='sex', ylabel='height'>
```



```
In [43]: df.sex.count()
```

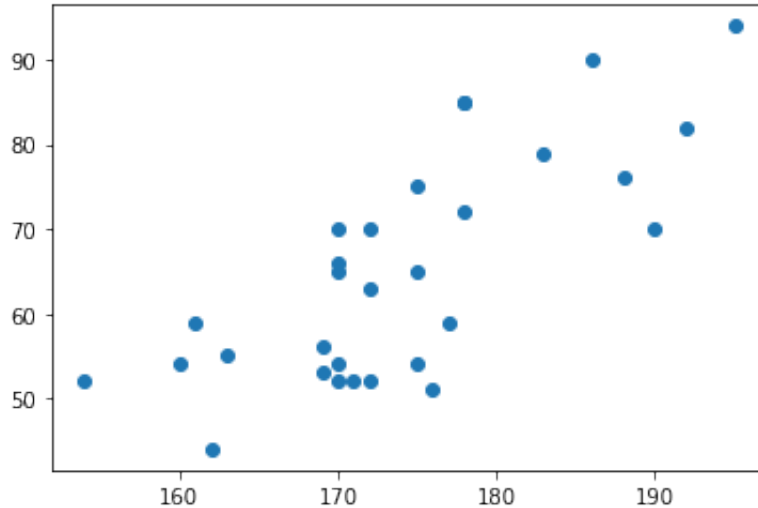
```
Out[43]: 30
```

Module 4: Correlation and Regression

Make a **scatter plot** of heights (horizontal axis) and weights (vertical axis) using `scatter()` from `matplotlib.pyplot` :

```
In [44]: import numpy as np
import matplotlib.pyplot as plt
plt.scatter(df.height,df.weight)
```

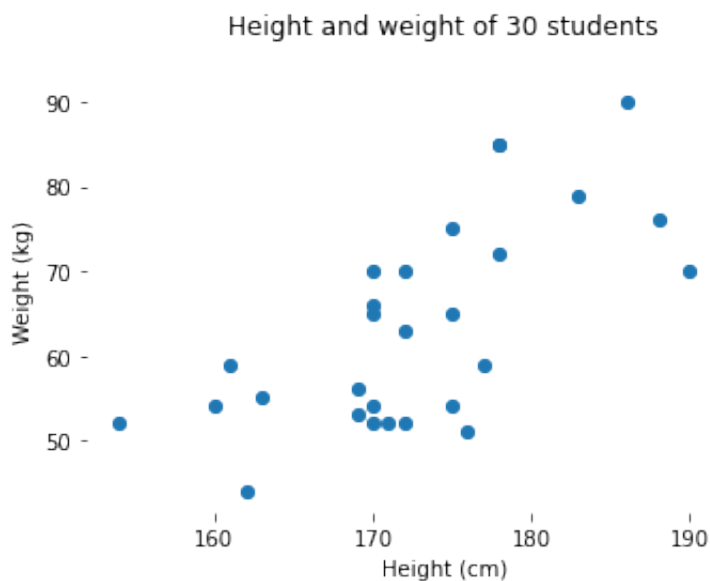
```
Out[44]: <matplotlib.collections.PathCollection at 0x7f7a4896a970>
```



Make the scatter plot prettier (get rid of the box, add labels to the axes):

```
In [45]: plt.box(False) # get rid of the box
plt.title("Height and weight of 30 students") # add title
plt.xlabel("Height (cm)") # add label to x-axis
plt.ylabel("Weight (kg)") # add label to y-axis
plt.scatter(df.height,df.weight)
```

```
Out[45]: <matplotlib.collections.PathCollection at 0x7f7a2a2bb1f0>
```



Correlation coefficient:

```
In [46]: # correlation between height and weight:
df.height.corr(df.weight)
```

```
Out[46]: 0.7525668130284301
```

```
In [47]: # correlation matrix between all quantitative variables of a data frame:
df.corr()
```

```
Out[47]:
```

	case	height	weight
case	1.000000	-0.320457	-0.254260
height	-0.320457	1.000000	0.752567
weight	-0.254260	0.752567	1.000000

To get *one* of the correlation coefficients, first convert to matrix:

```
In [48]: import numpy as np
corr_matrix = np.array(df.corr())
print(corr_matrix)

[[ 1.          -0.32045686 -0.25426022]
 [-0.32045686  1.          0.75256681]
 [-0.25426022  0.75256681  1.          ]]
```

```
In [49]: # extract the correlation between height and weight (caution: rows and columns)
corr_matrix[1][2]
```

```
Out[49]: 0.7525668130284303
```

To round numbers, use `round()` from the numpy library:

```
In [50]: import numpy as np
np.round(_,2) # in Python, _ is the output of the last cell; the second argument is the number of digits to round to
```

```
Out[50]: 0.75
```

Find the **line of best fit** (using the ordinary least squares method) using `statsmodels` :

```
In [51]: import statsmodels.api as sms
import statsmodels.formula.api as smf
# Fit regression model:
results = smf.ols('df.weight ~ df.height', data=df).fit()
# Inspect the results:
print(results.summary())
```

```

OLS Regression Results
=====
===
Dep. Variable:          df.weight    R-squared:          0.
566
Model:                  OLS          Adj. R-squared:       0.
551
Method:                 Least Squares    F-statistic:        36
.57
Date:                   Thu, 13 Jan 2022    Prob (F-statistic):   1.61e
-06
Time:                   16:36:42          Log-Likelihood:      -107
.29
No. Observations:      30              AIC:                21
8.6
Df Residuals:          28              BIC:                22
1.4
Df Model:              1
Covariance Type:       nonrobust
=====
===
               coef      std err          t      P>|t|      [0.025      0.9
75]
-----
---
Intercept    -116.6011     30.097     -3.874     0.001    -178.252    -54.
951
df.height     1.0443       0.173      6.047     0.000      0.691      1.
398
=====
===
Omnibus:          2.804    Durbin-Watson:      2.
486
Prob(Omnibus):    0.246    Jarque-Bera (JB):    1.
333
Skew:            0.007    Prob(JB):            0.
514
Kurtosis:        1.968    Cond. No.            3.21e
+03
=====
===

```

Notes:

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified.

[2] The condition number is large, 3.21e+03. This might indicate that there are strong multicollinearity or other numerical problems.

To get just the coefficients:

In [52]:

```
results.params
```

Out[52]:

```
Intercept    -116.601087
df.height     1.044251
dtype: float64
```

The intercept is the first coefficient:

```
In [53]: results.params[0]
```

```
Out[53]: -116.6010867272883
```

The slope coefficient is the second coefficient:

```
In [54]: results.params[1]
```

```
Out[54]: 1.044250641987867
```

To get the t -values of the coefficients (Statistics II):

```
In [55]: results.tvalues
```

```
Out[55]: Intercept    -3.874200  
df.height      6.047248  
dtype: float64
```

To get the p -values of the coefficients (Statistics II):

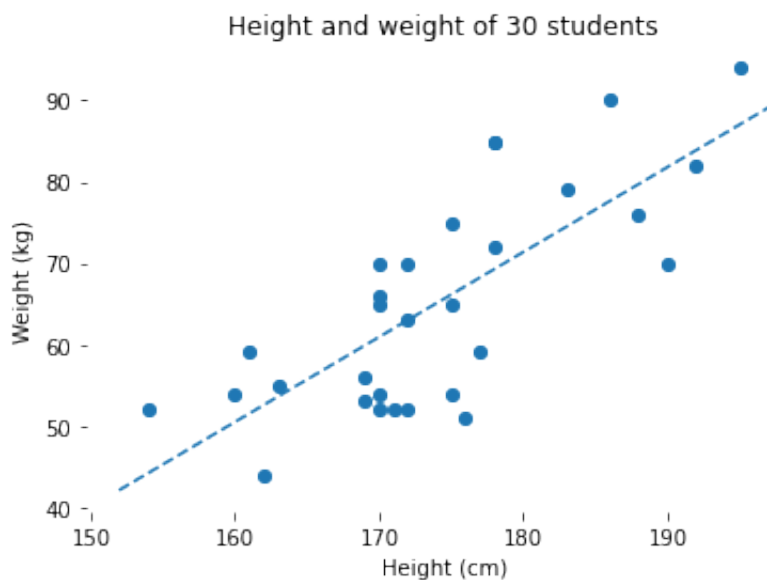
```
In [56]: results.pvalues
```

```
Out[56]: Intercept      0.000588  
df.height      0.000002  
dtype: float64
```

Add line of best fit to the scatter plot (see:

<https://stackoverflow.com/questions/7941226/how-to-add-line-based-on-slope-and-intercept-in-matplotlib>):

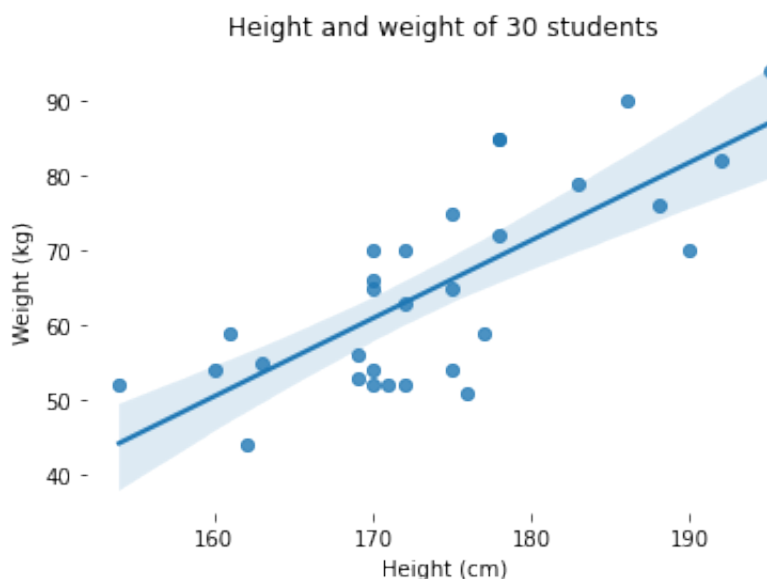
```
In [57]: import matplotlib.pyplot as plt  
import numpy as np  
import statsmodels.api as sms  
import statsmodels.formula.api as smf  
  
# Fit regression model:  
results = smf.ols('df.weight ~ df.height', data=df).fit()  
  
def abline(slope, intercept):  
    """Plot a line from slope and intercept"""  
    axes = plt.gca()  
    x_vals = np.array(axes.get_xlim())  
    y_vals = intercept + slope * x_vals  
    plt.plot(x_vals, y_vals, '--')  
  
plt.box(False) # get rid of the box  
plt.title("Height and weight of 30 students") # add title  
plt.xlabel("Height (cm)") # add label to x-axis  
plt.ylabel("Weight (kg)") # add label to y-axis  
plt.scatter(df.height, df.weight)  
abline(results.params[1], results.params[0]) # add line y=a*x+b (a = s
```



The **seaborn** library has more sophisticated ways to visualize data. For instance, to add a line of best fit with a 95 percent confidence interval to the scatter plot:

In [58]:

```
import seaborn as sns
sns.regplot(x=df.height, y=df.weight, data=df) # regplot: regression plot
plt.box(False) # get rid of the box
plt.title("Height and weight of 30 students") # add title
plt.xlabel("Height (cm)") # add label to x-axis
plt.ylabel("Weight (kg)") # add label to y-axis
plt.show()
```



Module 5: Randomness and Probability

(No Python code for this module.)

Module 6: Random variables and probability models

Calculate **binomial probability** (k = number of successes, n = number of trials, p = probability of success) (pmf stands for probability mass function—like pdf but for a discrete random variable):

```
In [59]: # calculate binomial probability (k= number of successes, n = number of trials)  
from scipy.stats import binom  
binom.pmf(k=10, n=12, p=0.6) # pmf: probability mass function (like pdf for continuous)
```

Out[59]: 0.063852281856

Calculate cumulative binomial probability:

```
In [60]: from scipy.stats import binom  
binom.cdf(k=10, n=12, p=0.6)
```

Out[60]: 0.980408958976

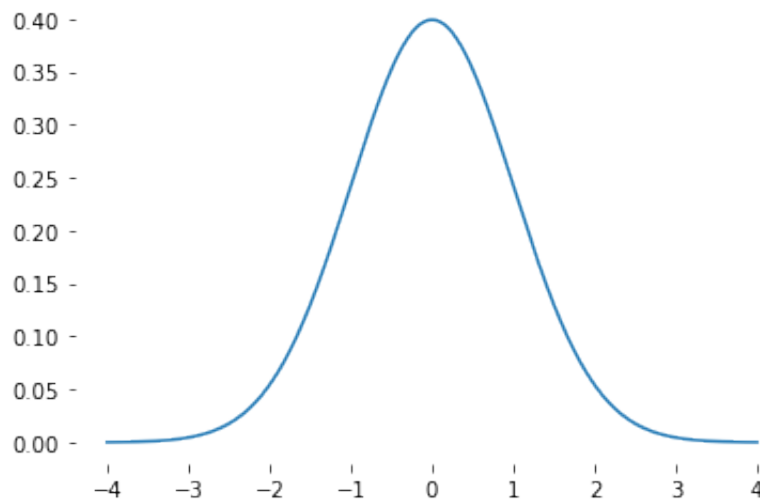
Module 7: The Normal distribution

Plot the **probability density function** (pdf) of the normal curve

(<https://www.statology.org/plot-normal-distribution-python/>):

```
In [61]: import numpy as np  
import matplotlib.pyplot as plt  
from scipy.stats import norm  
# x-axis ranges from -4 and 4 with .001 steps:  
x = np.arange(-4, 4, 0.001)  
  
# plot normal distribution with mean 0 and standard deviation 1  
plt.box(False) # get rid of the box around the plot  
plt.plot(x, norm.pdf(x, 0, 1))
```

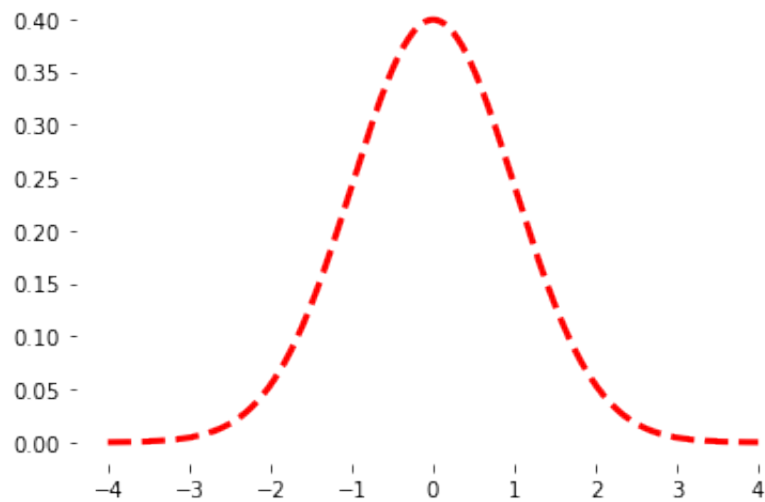
Out[61]: [



To change color, linewidth, linestyle:

```
In [62]: plt.box(False) # get rid of the box around the plot  
plt.plot(x, norm.pdf(x, 0, 1), color='red', linewidth=3, linestyle='dashed')
```


Out[62]: [



Area under normal curve:

To find an area under the normal curve, use the **cumulative density function** (cdf) of the normal distribution. (documentation: see:

<https://docs.scipy.org/doc/scipy/reference/stats.html>)

In [63]:

```
import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import norm
mu = 0          # mean (for the standard normal distribution, mu = 0)
sigma = 1       # standard deviation (for the standard normal distribution, sigma = 1)
x1 = -1.96      # lower boundary
x2 = 1.96       # lower boundary
# area under normal curve between x1 and x2:
area = norm.cdf(x2, loc=mu, scale=sigma)-norm.cdf(x1, loc=mu, scale=sigma)
print('The area under the normal curve is', area)
```

The area under the normal curve is 0.950004209703559

To **plot** the area under the normal curve (see:

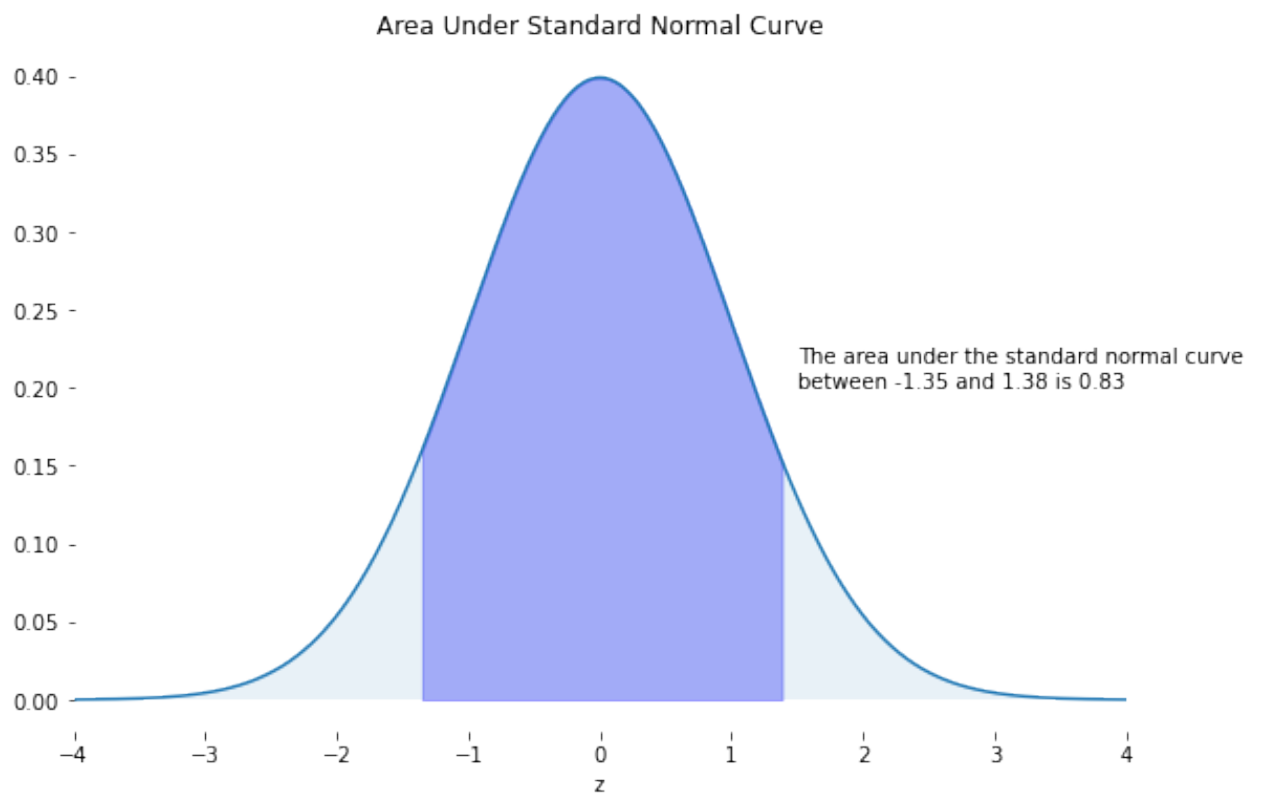
<https://pythonforundergradengineers.com/plotting-normal-curve-with-python.html>)

In [64]:

```
import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import norm
# define constants
mu = 998.8 # mean
sigma = 73.10 # standard deviation
x1 = 900 # lower boundary
x2 = 1100 # lower boundary
# calculate the standardized values:
z1 = ( x1 - mu ) / sigma
z2 = ( x2 - mu ) / sigma
x = np.arange(z1, z2, 0.001) # range of x in spec
x_all = np.arange(-10, 10, 0.001) # entire range of x, both in and out of spec
# for standard normal distribution, mean = 0, stddev = 1:
y = norm.pdf(x,0,1)
y2 = norm.pdf(x_all,0,1)
# find area under normal curve between x1 and x2:
area = norm.cdf(x2, loc=mu, scale=sigma)-norm.cdf(x1, loc=mu, scale=sigma)

# build the plot
fig, ax = plt.subplots(figsize=(9,6))
ax.plot(x_all,y2)
ax.fill_between(x,y,0, alpha=0.3, color='b')
ax.fill_between(x_all,y2,0, alpha=0.1)
ax.set_xlim([-4,4])
ax.set_xlabel('z')
# ax.set_yticklabels([])
ax.set_title('Area Under Standard Normal Curve')
plt.box(False)
plt.text(1.5,0.2, f"The area under the standard normal curve \nbetween {np.round(z1,2)} and {np.round(z2,2)} is {area:.4f}")
plt.savefig('normal_curve.png', dpi=72, bbox_inches='tight')
plt.show()

print(f"The area under the standard normal curve between {np.round(z1,2)} and {np.round(z2,2)} is {area:.4f}")
```



The area under the standard normal curve between -1.35 and 1.38 is 0.83.

Other continuous distributions

Uniform distribution

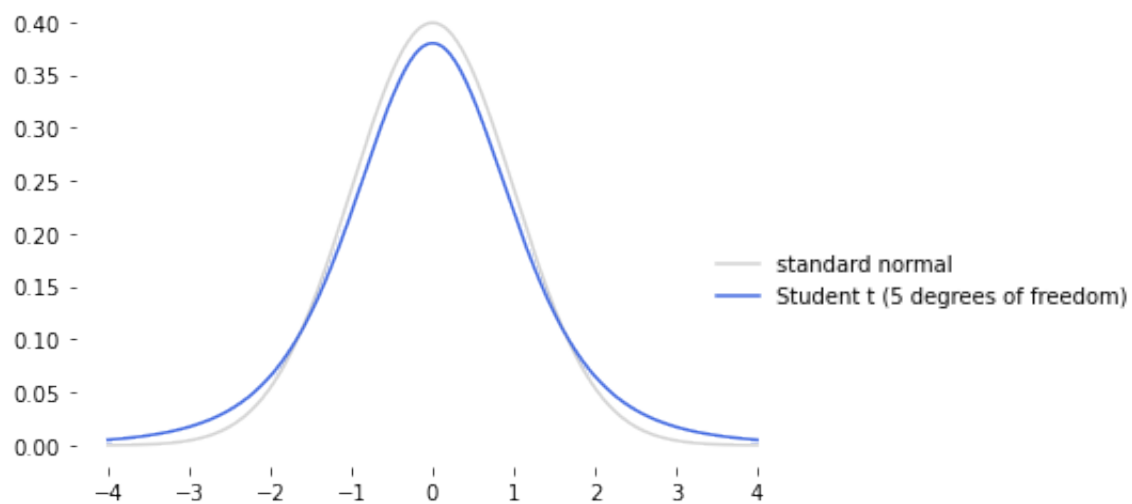
(<https://docs.scipy.org/doc/scipy/reference/generated/scipy.stats.uniform.html>):

```
In [65]: from scipy.stats import uniform
# In the standard form, the distribution is uniform on [0, 1].
# Using the parameters loc and scale, one obtains the uniform distribution
uniform.cdf(0.9)
```

Out[65]: 0.9

Student *t* distribution (covered in Statistics II)

```
In [66]: import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import t
dof = 5 # degrees of freedom of Student t distribution
# x-axis ranges from -4 and 4 with .001 steps:
x = np.arange(-4, 4, 0.001)
plt.figure()
plt.box(False)
plt.plot(x, norm.pdf(x,0,1 ), color="lightgrey") # plot the standard normal
plt.plot(x, t.pdf(x, dof) , color="royalblue") # the second argument
plt.legend(["standard normal" , f"Student t ({dof} degrees of freedom)"], 1)
# plt.text(3.5,0.35, f"({dof} degrees of freedom)")
plt.show()
```



Plot Student t distributions with different degrees of freedom and compare with standard normal distribution (see Haslwanter (2016), p. 110) in an interactive diagram where you can change the degrees of freedom with a slider:

(Normally the `ipywidgets` library is installed in Anaconda. If that is not the case, remove the hashtag at the beginning of the following line and run the line: `pip install ipywidgets`.)

```
In [67]: import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import norm, t
from ipywidgets import interact # import ipywidgets.interact class. This class
# x-axis ranges from -4 to 4 with steps of .001 :
x = np.arange(-4, 4, 0.001)
# use interact decorator to decorate the function,
# so the function can receive the slide bar's value with the parameter (in
@interact(dof=(1,60,1))
def plt_t(dof):
    plt.figure()
    plt.box(False)
    plt.plot(x, norm.pdf(x,0,1), color="lightgrey") # plot the standard normal
    plt.plot(x, t.pdf(x, dof), color="royalblue") # plot Student t distribution
    plt.legend(["standard normal", "Student  $t$ "], loc="upper right")
    plt.show
```

To find an area under the Student t curve, use the **cumulative density function** (cdf) of the Student t distribution (documentation: see:

<https://docs.scipy.org/doc/scipy/reference/stats.html>):

```
In [68]: import numpy as np
from scipy.stats import t
dof = 30 # degrees of freedom
t1 = -1.00 # lower boundary
t2 = 1.00 # lower boundary
# area under Student t curve between t1 and t2:
area = t.cdf(t2,dof)-t.cdf(t1,dof)
print(f'The area under the Student t curve with {dof} degrees of freedom be
```

The area under the Student t curve with 30 degrees of freedom between -1.0 and 1.0 is 0.6746913845739702

(To **plot** the area, adapt the code above to plot the area under the normal curve.)

Module 9: Sampling Distributions and Confidence Intervals for Proportions

To find the **confidence interval for a proportion**:

(https://www.statsmodels.org/dev/generated/statsmodels.stats.proportion.proportion_confint)

```
In [69]: 310 / 1126      # sample proportion = p-hat = number of successes / number of trials

import statsmodels.api as sm
from statsmodels.stats.proportion import proportion_confint # Function for confidence interval

proportion_confint(count=310,          # count= number of successes
                   nobs=1126,          # nobs = number of trials
                   alpha=(1 - 0.95)) # alpha = 1 - confidence level
```

```
Out[69]: (0.24922129423231776, 0.30140037539468045)
```

Module 10: Sampling Distributions and Confidence Intervals for Means

To find the **confidence interval for a mean** (in this case: the mean height of all students in the dataframe df):

```
In [70]: import numpy as np
import scipy.stats as stats

degrees_of_freedom = len(df)-1          # degrees of freedom = sample size - 1
sample_mean = np.mean(df.height)        # sample mean
sample_standard_error = stats.sem(df.height) # sample standard error of the mean

# create confidence interval for the population mean:
stats.t.interval(alpha=0.05, df=degrees_of_freedom, loc=sample_mean, scale=sample_standard_error)
```

```
Out[70]: (173.9221744669485, 174.14449219971817)
```

Hypothesis tests (Statistics II)

For hypothesis tests (covered in Statistics II), see:

<https://www.statsmodels.org/stable/stats.html>

Interacting with the operating system (changing current working directory etc.)

The `os` library allows you to interact with the operating system using Python code.

The **current working directory** (cwd) is where Python will look for input (such as data files) and where it will store output (such as .png or .pdf figures and tables with results). To find out what the current working directory of Python is:

In [71]:

```
import os
print(os.getcwd())
```

```
/Users/luchens/Documents/jupyter-notebooks
```

If you want to change the current working directory, use the `chdir` command of the `os` library to do so. Here is how to change the current working directory (the expression in quotes is the path to the new working directory — it will be a different path for you, of course):

In [72]:

```
os.chdir('/Users/luchens/Documents/Data/')
print(os.getcwd())
```

```
/Users/luchens/Documents/Data
```

To display the files and directories in the working directory, use `os.listdir()`