

3.1. Statistics in Python

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Requirements

- Standard scientific Python environment (numpy, scipy, matplotlib)
- [Pandas](#)
- [Statsmodels](#)
- [Seaborn](#)

To install Python and these dependencies, we recommend that you download [Anaconda Python](#) or [Enthought Canopy](#), or preferably use the package manager if you are under Ubuntu or other linux.

See also:

- **Bayesian statistics in Python:** This chapter does not cover tools for Bayesian statistics. Of particular interest for Bayesian modelling is [PyMC](#), which implements a probabilistic programming language in Python.
- **Read a statistics book:** The [Think stats](#) book is available as free PDF or in print and is a great introduction to statistics.

Why Python for statistics?

R is a language dedicated to statistics. Python is a general-purpose language with statistics modules. R has more statistical analysis features than Python, and specialized syntaxes. However, when it comes to building complex analysis pipelines that mix statistics with e.g. image analysis, text mining, or control of a physical experiment, the richness of Python is an invaluable asset.

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In this document, the Python inputs are represented with the sign “>>>”.

Disclaimer: Gender questions

Some of the examples of this tutorial are chosen around gender questions. The reason is that on such questions controlling the truth of a claim actually matters to many people.

3.1.1. Data representation and interaction

3.1.1.1. Data as a table

The setting that we consider for statistical analysis is that of multiple *observations* or *samples* described by a set of different *attributes* or *features*. The data can then be seen as a 2D table, or matrix, with columns giving the different attributes of the data, and rows the observations. For instance, the data contained in `examples/brain_size.csv`:

```
"";"Gender";"FSIQ";"VIQ";"PIQ";"Weight";"Height";"MRI_Count"
"1";"Female";133;132;124;"118";"64.5";816932
"2";"Male";140;150;124;".";"72.5";1001121
"3";"Male";139;123;150;"143";"73.3";1038437
"4";"Male";133;129;128;"172";"68.8";965353
"5";"Female";137;132;134;"147";"65.0";951545
```

3.1.1.2. The pandas data-frame

We will store and manipulate this data in a `pandas.DataFrame`, from the `pandas` module. It is the Python equivalent of the spreadsheet table. It is different from a 2D numpy array as it has named columns, can contain a mixture of different data types by column, and has elaborate selection and pivotal mechanisms.

Creating dataframes: reading data files or converting arrays

Separator

It is a CSV file, but the separator is “;”

Reading from a CSV file: Using the above CSV file that gives observations of brain size and weight and IQ (Willerman et al. 1991), the data are a mixture of numerical and categorical values:

```
>>> import pandas
>>> data = pandas.read_csv('examples/brain_size.csv', sep=';', na_values=
    ".")
>>> data
```

	Unnamed: 0	Gender	FSIQ	VIQ	PIQ	Weight	Height	MRI_Count
0	1	Female	133	132	124	118.0	64.5	816932
1	2	Male	140	150	124	NaN	72.5	1001121

2	3	Male	139	123	150	143.0	73.3	1038437
3	4	Male	133	129	128	172.0	68.8	965353
4	5	Female	137	132	134	147.0	65.0	951545
...								

⚠ Missing values

The weight of the second individual is missing in the CSV file. If we don't specify the missing value (NA = not available) marker, we will not be able to do statistical analysis.

Creating from arrays: A `pandas.DataFrame` can also be seen as a dictionary of 1D 'series', eg arrays or lists. If we have 3 numpy arrays:

```
>>> import numpy as np
>>> t = np.linspace(-6, 6, 20)
>>> sin_t = np.sin(t)
>>> cos_t = np.cos(t)
```

>>>

We can expose them as a `pandas.DataFrame`:

```
>>> pandas.DataFrame({'t': t, 'sin': sin_t, 'cos': cos_t})
      t      sin      cos
0 -6.000000  0.279415  0.960170
1 -5.368421  0.792419  0.609977
2 -4.736842  0.999701  0.024451
3 -4.105263  0.821291 -0.570509
4 -3.473684  0.326021 -0.945363
5 -2.842105 -0.295030 -0.955488
6 -2.210526 -0.802257 -0.596979
7 -1.578947 -0.999967 -0.008151
8 -0.947368 -0.811882  0.583822
...
```

>>>

Other inputs: `pandas` can input data from SQL, excel files, or other formats. See the [pandas documentation](#).

Manipulating data

`data` is a `pandas.DataFrame`, that resembles R's dataframe:

```
>>> data.shape      # 40 rows and 8 columns
(40, 8)

>>> data.columns    # It has columns
Index([u'Unnamed: 0', u'Gender', u'FSIQ', u'VIQ', u'PIQ', u'Weight',
      u'Height', u'MRI_Count'], dtype='object')

>>> print(data['Gender']) # Columns can be addressed by name
0      Female
1       Male
```

>>>

```
2      Male
3      Male
4      Female
...
```

```
>>> # Simpler selector
>>> data[data['Gender'] == 'Female']['VIQ'].mean()
109.45
```

Note: For a quick view on a large dataframe, use its *describe* method:
`pandas.DataFrame.describe()`.

groupby: splitting a dataframe on values of categorical variables:

```
>>> groupby_gender = data.groupby('Gender')
>>> for gender, value in groupby_gender['VIQ']:
...     print((gender, value.mean()))
('Female', 109.45)
('Male', 115.25)
```

>>>

groupby_gender is a powerful object that exposes many operations on the resulting group of dataframes:

```
>>> groupby_gender.mean()
      Unnamed: 0    FSIQ    VIQ    PIQ    Weight    Height    MRI_Count
Gender
Female      19.65   111.9   109.45   110.45   137.200000    65.765000    862654.6
Male       21.35   115.0   115.25   111.60   166.444444    71.431579    954855.4
```

>>>

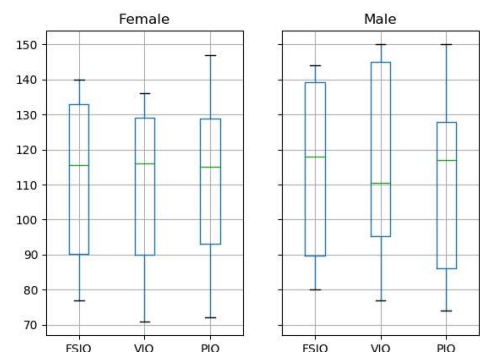
Use tab-completion on *groupby_gender* to find more. Other common grouping functions are median, count (useful for checking to see the amount of missing values in different subsets) or sum. Groupby evaluation is lazy, no work is done until an aggregation function is applied.

Exercise

- What is the mean value for VIQ for the full population?
- How many males/females were included in this study?

Hint use 'tab completion' to find out the methods that can be called, instead of 'mean' in the above example.

- What is the average value of MRI counts expressed in log units, for males and females?



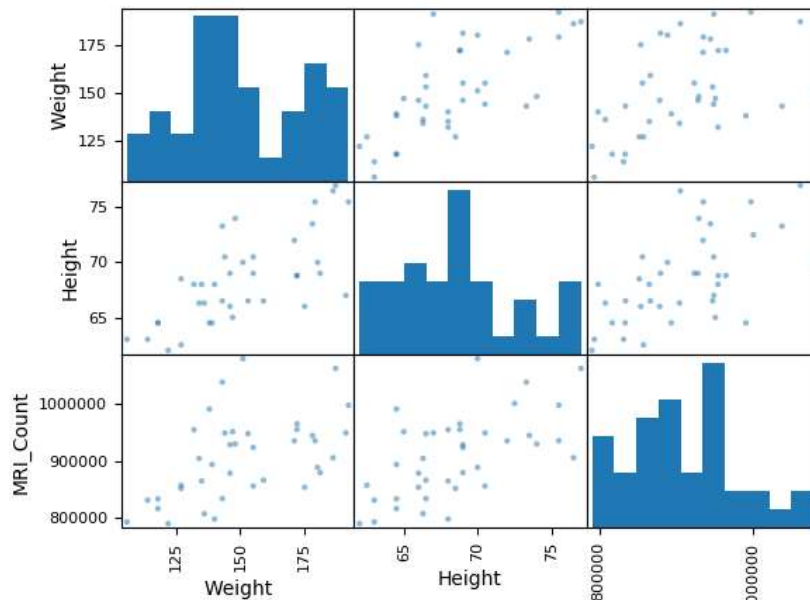
Note: *groupby_gender.boxplot* is used for the plots above (see [this example](#)).

Plotting data

Pandas comes with some plotting tools (**pandas.tools.plotting**, using matplotlib behind the scene) to display statistics of the data in dataframes:

Scatter matrices:

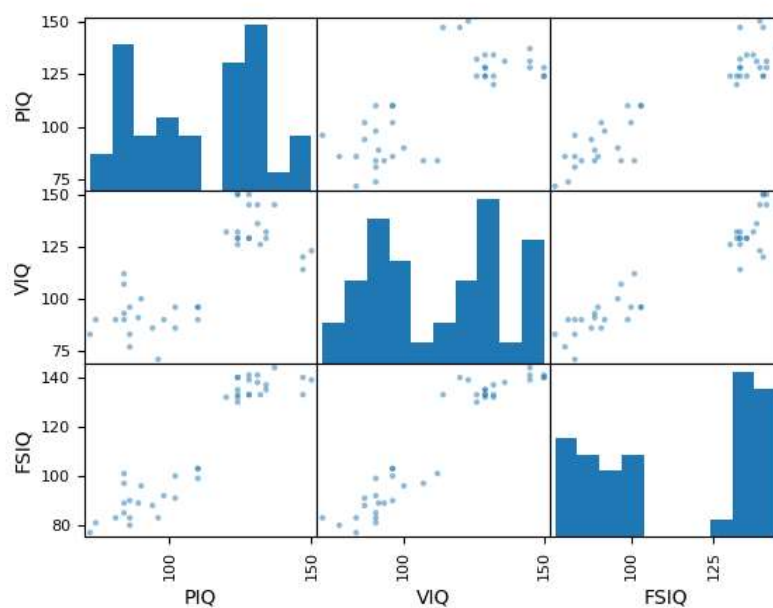
```
>>> from pandas.tools import plotting
>>> plotting.scatter_matrix(data[['Weight', 'Height', 'MRI_Count']])
```



```
>>> plotting.scatter_matrix(data[['PIQ', 'VIQ', 'FSIQ']])
```

Two populations

The IQ metrics are bimodal, as if there are 2 sub-populations.



Exercise

Plot the scatter matrix for males only, and for females only. Do you think that the 2 sub-populations correspond to gender?

3.1.2. Hypothesis testing: comparing two groups

For simple statistical tests, we will use the `scipy.stats` sub-module of `scipy`:

```
>>> from scipy import stats
```

```
>>>
```

See also: Scipy is a vast library. For a quick summary to the whole library, see the [scipy](#) chapter.

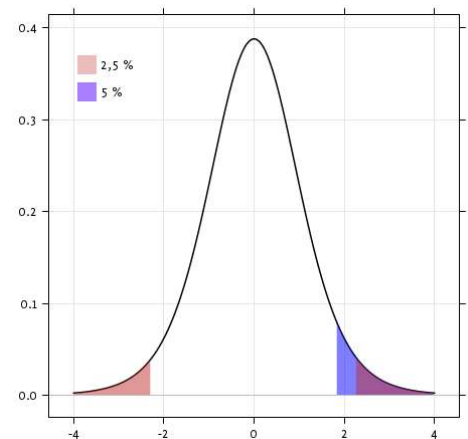
3.1.2.1. Student's t-test: the simplest statistical test

1-sample t-test: testing the value of a population mean

`scipy.stats.ttest_1samp()` tests if the population mean of data is likely to be equal to a given value (technically if observations are drawn from a Gaussian distributions of given population mean). It returns the **T statistic**, and the **p-value** (see the function's help):

```
>>> stats.ttest_1samp(data['VIQ'], 0)
Ttest_1sampResult(statistic=30.088099970...,
                  pvalue=1.32891964...e-28)
```

With a p-value of 10^{-28} we can claim that the population mean for the IQ (VIQ measure) is not 0.



2-sample t-test: testing for difference across populations

We have seen above that the mean VIQ in the male and female populations were different. To test if this is significant, we do a 2-sample t-test with `scipy.stats.ttest_ind()`:

```
>>> female_viq = data[data['Gender'] == 'Female']['VIQ']
>>> male_viq = data[data['Gender'] == 'Male']['VIQ']
>>> stats.ttest_ind(female_viq, male_viq)
Ttest_indResult(statistic=-0.77261617232..., pvalue=0.4445287677858...)
```

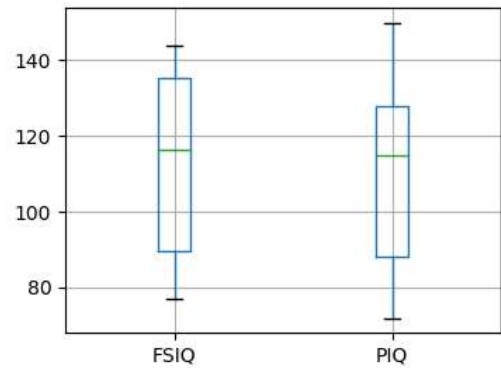
3.1.2.2. Paired tests: repeated measurements on the same individuals

PIQ, VIQ, and FSIQ give 3 measures of IQ. Let us test if FSIQ and PIQ are significantly different. We can use a 2 sample test:

```
>>> stats.ttest_ind(data['FSIQ'], data['PIQ'])
Ttest_indResult(statistic=0.46563759638..., pvalue=0.64277250...)
```

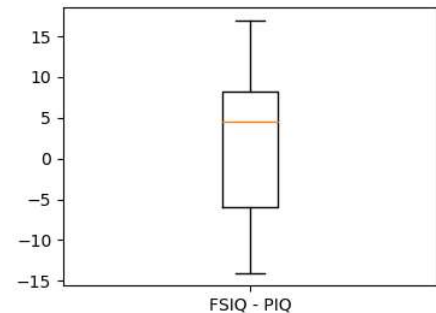
The problem with this approach is that it forgets that there are links between observations: FSIQ and PIQ are measured on the same individuals. Thus the variance due to inter-subject variability is confounding, and can be removed, using a “paired test”, or “repeated measures test”:

```
>>> stats.ttest_rel(data['FSIQ'], data['PIQ'])
Ttest_relResult(statistic=1.784201940...,
                 pvalue=0.082172638183...)
```



This is equivalent to a 1-sample test on the difference:

```
>>> stats.ttest_1samp(data['FSIQ'] - data['PIQ'], 0)
Ttest_1sampResult(statistic=1.784201940...,
                  pvalue=0.082172638183...)
```



T-tests assume Gaussian errors. We can use a [Wilcoxon signed-rank test](#), that relaxes this assumption:

```
>>> stats.wilcoxon(data['FSIQ'], data['PIQ'])
WilcoxonResult(statistic=274.5, pvalue=0.106594927...)
```

Note: The corresponding test in the non paired case is the [Mann–Whitney U test](#), `scipy.stats.mannwhitneyu()`.

Exercise

- Test the difference between weights in males and females.
- Use non parametric statistics to test the difference between VIQ in males and females.

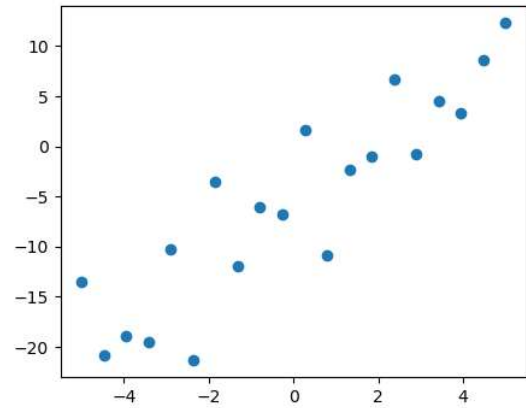
Conclusion: we find that the data does not support the hypothesis that males and females have different VIQ.

3.1.3. Linear models, multiple factors, and analysis of variance

3.1.3.1. “formulas” to specify statistical models in Python

A simple linear regression

Given two set of observations, x and y , we want to test the hypothesis that y is a linear function of x . In other terms:



$$y = x * coef + intercept + e$$

where e is observation noise. We will use the `statsmodels` module to:

1. Fit a linear model. We will use the simplest strategy, `ordinary least squares` (OLS).
2. Test that `coef` is non zero.

First, we generate simulated data according to the model:

```
>>> import numpy as np
>>> x = np.linspace(-5, 5, 20)
>>> np.random.seed(1)
>>> # normal distributed noise
>>> y = -5 + 3*x + 4 * np.random.normal(size=x.shape)
>>> # Create a data frame containing all the relevant variables
>>> data = pandas.DataFrame({'x': x, 'y': y})
```

“formulas” for statistics in Python

[See the statsmodels documentation](#)

Then we specify an OLS model and fit it:

```
>>> from statsmodels.formula.api import ols
>>> model = ols("y ~ x", data).fit()
```

We can inspect the various statistics derived from the fit:

```
>>> print(model.summary())
```

OLS Regression Results

```
=====...
Dep. Variable:          y    R-squared:
    0.804
Model:                OLS    Adj. R-squared:
    0.794
Method:             Least Squares    F-statistic:
    74.03
Date:                ...    Prob (F-statistic):
    08                                8.56e-
Time:                ...    Log-Likelihood:
    -57.988
```



```

No. Observations:          20    AIC:
    120.0
Df Residuals:              18    BIC:
    122.0
Df Model:                  1
Covariance Type:          nonrobust
=====...
              coef      std err          t      P>|t|      [0.025
              0.975]
-----...
Intercept    -5.5335      1.036     -5.342      0.000     -7.710
              -3.357
x             2.9369      0.341      8.604      0.000      2.220
              3.654
=====...
Omnibus:              0.100    Durbin-Watson:
    2.956
Prob(Omnibus):        0.951    Jarque-Bera (JB):
    0.322
Skew:                -0.058    Prob(JB):
    0.851
Kurtosis:            2.390    Cond. No.
    3.03
=====...

Warnings:
[1] Standard Errors assume that the covariance matrix of the errors is cor-
rectly specified.

```

Terminology:

Statsmodels uses a statistical terminology: the y variable in statsmodels is called 'endogenous' while the x variable is called exogenous. This is discussed in more detail [here](#).

To simplify, y (endogenous) is the value you are trying to predict, while x (exogenous) represents the features you are using to make the prediction.

Exercise

Retrieve the estimated parameters from the model above. **Hint:** use tab-completion to find the relevant attribute.

Categorical variables: comparing groups or multiple categories

Let us go back the data on brain size:

```
>>> data = pandas.read_csv('examples/brain_size.csv', sep=';', na_values=
    ".")
```

We can write a comparison between IQ of male and female using a linear model:

```
>>> model = ols("VIQ ~ Gender + 1", data).fit()
>>> print(model.summary())
```

OLS Regression Results

```
=====...
```

```

Dep. Variable:          VIQ    R-squared:
      0.015
Model:                  OLS    Adj. R-squared:
      -0.010
Method:                 Least Squares    F-statistic:
      0.5969
Date:                  ...          Prob (F-statistic):
      0.445
Time:                  ...          Log-Likelihood:
      -182.42
No. Observations:      40    AIC:
      368.8
Df Residuals:          38    BIC:
      372.2
Df Model:              1
Covariance Type:       nonrobust
=====...
               coef      std err          t      P>|t|      [0.025      0.975]
-----
Intercept      109.4500      5.308      20.619      0.000      98.704
      120.196
Gender[T.Male]    5.8000      7.507       0.773      0.445      -9.397
      20.997
=====...
Omnibus:              26.188    Durbin-Watson:
      1.709
Prob(Omnibus):       0.000    Jarque-Bera (JB):
      3.703
Skew:               0.010    Prob(JB):
      0.157
Kurtosis:           1.510    Cond. No.
      2.62
=====...

Warnings:
[1] Standard Errors assume that the covariance matrix of the errors is cor-
rectly specified.

```

Tips on specifying model

Forcing categorical: the 'Gender' is automatically detected as a categorical variable, and thus each of its different values are treated as different entities.

An integer column can be forced to be treated as categorical using:

```
>>> model = ols('VIQ ~ C(Gender)', data).fit()
```

```
>>>
```

Intercept: We can remove the intercept using `- 1` in the formula, or force the use of an intercept using `+ 1`.

By default, statsmodels treats a categorical variable with K possible values as K-1 'dummy' boolean variables (the last level being absorbed into the intercept term). This is almost always a good default choice - however, it is possible to specify different encodings for categorical variables (<http://statsmodels.sourceforge.net/devel/contrasts.html>).

Link to t-tests between different FSIQ and PIQ

To compare different types of IQ, we need to create a “long-form” table, listing IQs, where the type of IQ is indicated by a categorical variable:

```
>>> data_fisq = pandas.DataFrame({'iq': data['FSIQ'], 'type': 'fsiq'})
>>> data_piq = pandas.DataFrame({'iq': data['PIQ'], 'type': 'piq'})
>>> data_long = pandas.concat((data_fisq, data_piq))
>>> print(data_long)
      iq  type
0   133  fsiq
1   140  fsiq
2   139  fsiq
...
31  137  piq
32  110  piq
33   86  piq
...
```

```
>>> model = ols("iq ~ type", data_long).fit()
>>> print(model.summary())
```

OLS Regression Results

```
...
=====...
              coef      std err          t      P>|t|      [0.025
-----
Intercept    113.4500      3.683      30.807      0.000     106.119
type[T.piq]   -2.4250      5.208     -0.466      0.643     -12.793
...
0.975]
```

We can see that we retrieve the same values for t-test and corresponding p-values for the effect of the type of iq than the previous t-test:

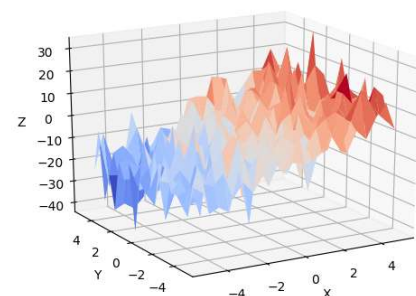
```
>>> stats.ttest_ind(data['FSIQ'], data['PIQ'])
Ttest_indResult(statistic=0.46563759638..., pvalue=0.64277250...)
```

3.1.3.2. Multiple Regression: including multiple factors

Consider a linear model explaining a variable z (the dependent variable) with 2 variables x and y :

$$z = x c_1 + y c_2 + i + e$$

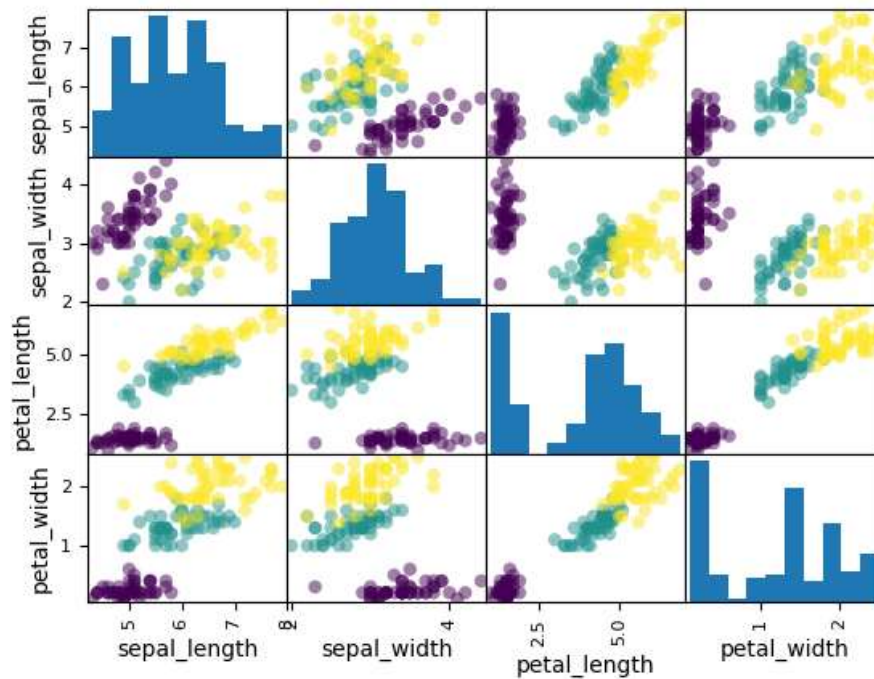
Such a model can be seen in 3D as fitting a plane to a cloud of (x, y, z) points.



Example: the iris data ([examples/iris.csv](#))

Sepal and petal size tend to be related: bigger flowers are bigger! But is there in addition a systematic effect of species?

blue: setosa, green: versicolor, red: virginica



```
>>> data = pandas.read_csv('examples/iris.csv')
>>> model = ols('sepal_width ~ name + petal_length', data).fit()
>>> print(model.summary())
```

>>>

OLS Regression Results

```
=====...
Dep. Variable:          sepal_width    R-squared:
    0.478
Model:                  OLS    Adj. R-squared:
    0.468
Method:                 Least Squares    F-statistic:
    44.63
Date:                  ...    Prob (F-statistic):
    20    1.58e-
Time:                  ...    Log-Likelihood:
    -38.185
No. Observations:      150    AIC:
    84.37
Df Residuals:          146    BIC:
    96.41
Df Model:               3
Covariance Type:        nonrobust
=====...
              coef    std err          t      P>|t|  [0.025
-----
Intercept      2.9813    0.099    29.989    0.000    2.785
name[T.versicolor] -1.4821    0.181   -8.190    0.000   -1.840
name[T.virginica] -1.6635    0.256   -6.502    0.000   -2.169
petal_length    0.2983    0.061    4.920    0.000    0.178
=====...
Omnibus:          2.868    Durbin-Watson:
    1.753
Prob(Omnibus):    0.238    Jarque-Bera (JB):
    2.885
```

```
Skew:          -0.082    Prob(JB) :
          0.236
Kurtosis:      3.659    Cond. No.
          54.0
=====...
```

Warnings:

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

3.1.3.3. Post-hoc hypothesis testing: analysis of variance (ANOVA)

In the above iris example, we wish to test if the petal length is different between versicolor and virginica, after removing the effect of sepal width. This can be formulated as testing the difference between the coefficient associated to versicolor and virginica in the linear model estimated above (it is an Analysis of Variance, [ANOVA](#)). For this, we write a **vector of 'contrast'** on the parameters estimated: we want to test "name[T.versicolor] - name[T.virginica]", with an **F-test**:

```
>>> print(model.f_test([0, 1, -1, 0]))
<F test: F=array([[3.24533535]]), p=0.07369..., df_denom=146, df_num=1>
```

Is this difference significant?

Exercise

Going back to the brain size + IQ data, test if the VIQ of male and female are different after removing the effect of brain size, height and weight.

3.1.4. More visualization: seaborn for statistical exploration

[Seaborn](#) combines simple statistical fits with plotting on pandas dataframes.

Let us consider a data giving wages and many other personal information on 500 individuals ([Berndt, ER. The Practice of Econometrics. 1991. NY: Addison-Wesley](#)).

The full code loading and plotting of the wages data is found in [corresponding example](#).

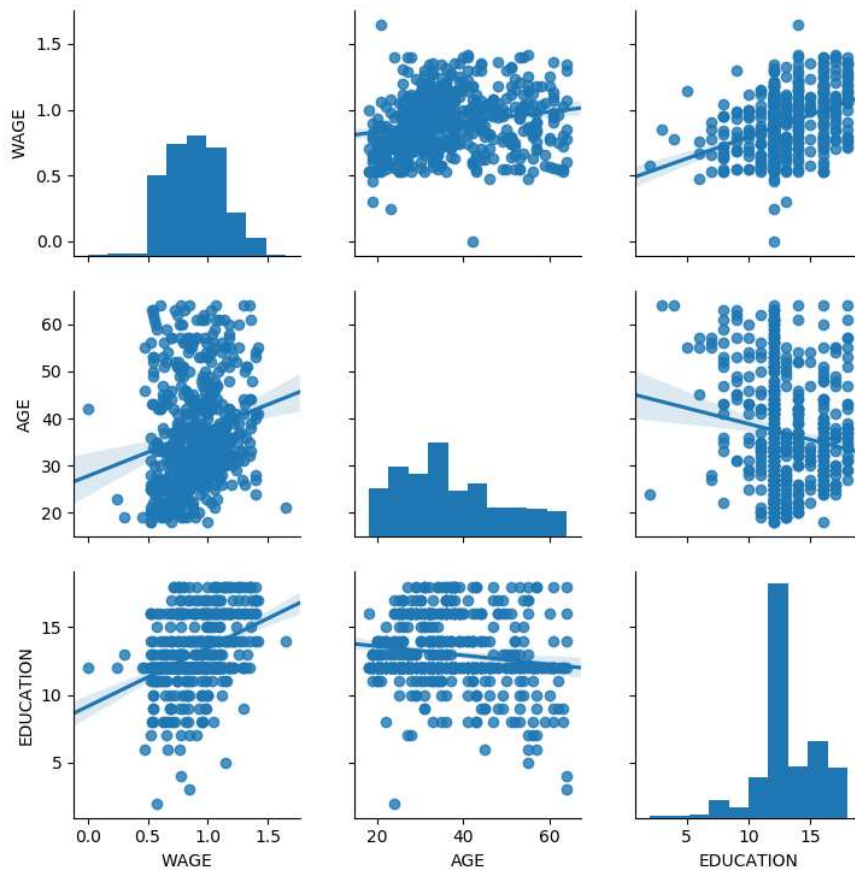
```
>>> print(data)
      EDUCATION  SOUTH  SEX  EXPERIENCE  UNION      WAGE  AGE  RACE  \
0             8       0    1          21       0  0.707570  35    2
1             9       0    1          42       0  0.694605  57    3
2            12       0    0           1       0  0.824126  19    3
3            12       0    0           4       0  0.602060  22    3
...
```

3.1.4.1. Pairplot: scatter matrices

We can easily have an intuition on the interactions between continuous variables using `seaborn.pairplot()` to display a scatter matrix:

```
>>> import seaborn
>>> seaborn.pairplot(data, vars=['WAGE', 'AGE', 'EDUCATION'],
...                  kind='reg')
```

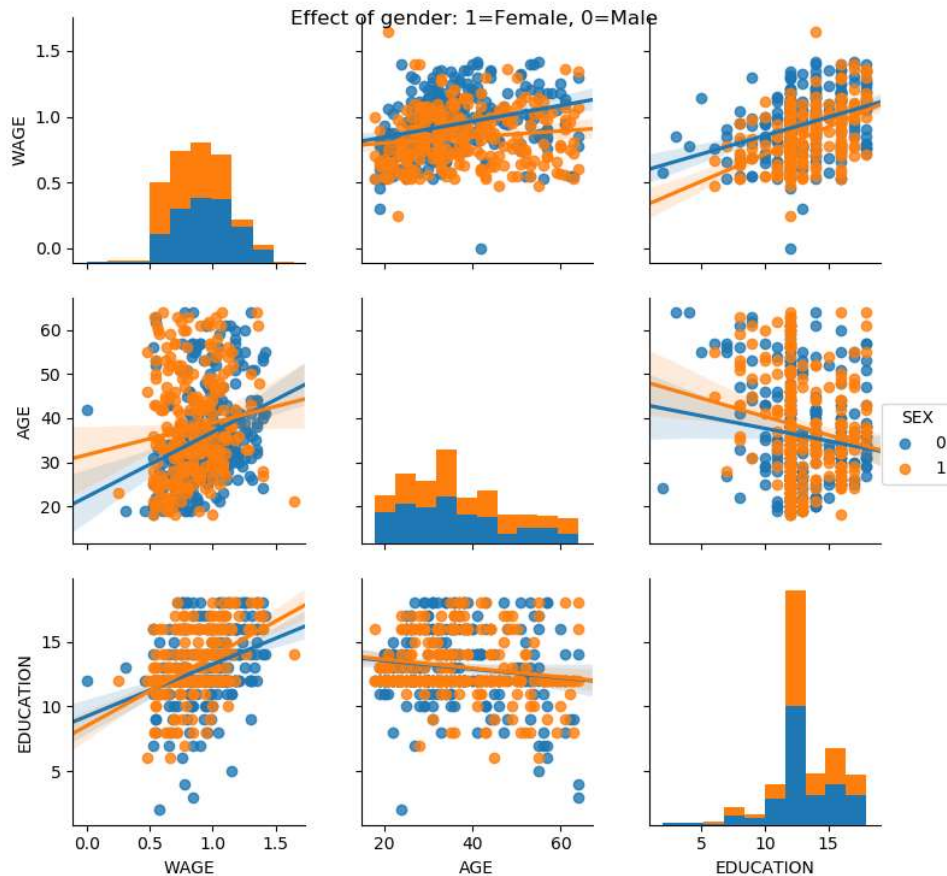
>>>



Categorical variables can be plotted as the hue:

```
>>> seaborn.pairplot(data, vars=['WAGE', 'AGE', 'EDUCATION'],
...                  kind='reg', hue='SEX')
```

>>>



Look and feel and matplotlib settings

Seaborn changes the default of matplotlib figures to achieve a more “modern”, “excel-like” look. It does that upon import. You can reset the default using:

```
>>> from matplotlib import pyplot as plt
>>> plt.rcParams()
```

```
>>>
```

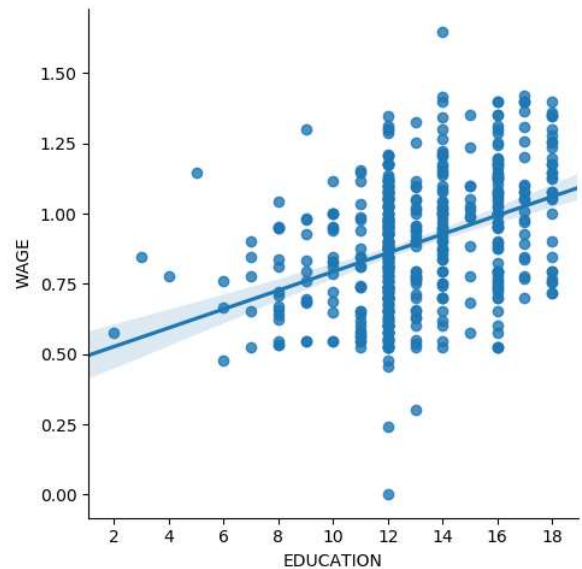
To switch back to seaborn settings, or understand better styling in seaborn, see the [relevant section of the seaborn documentation](#).

3.1.4.2. Implot: plotting a univariate regression

A regression capturing the relation between one variable and another, eg wage and education, can be plotted using `seaborn.lmplot()`:

```
>>> seaborn.lmplot(y='WAGE', x='EDUCATION', data=data)
```

```
>>>
```

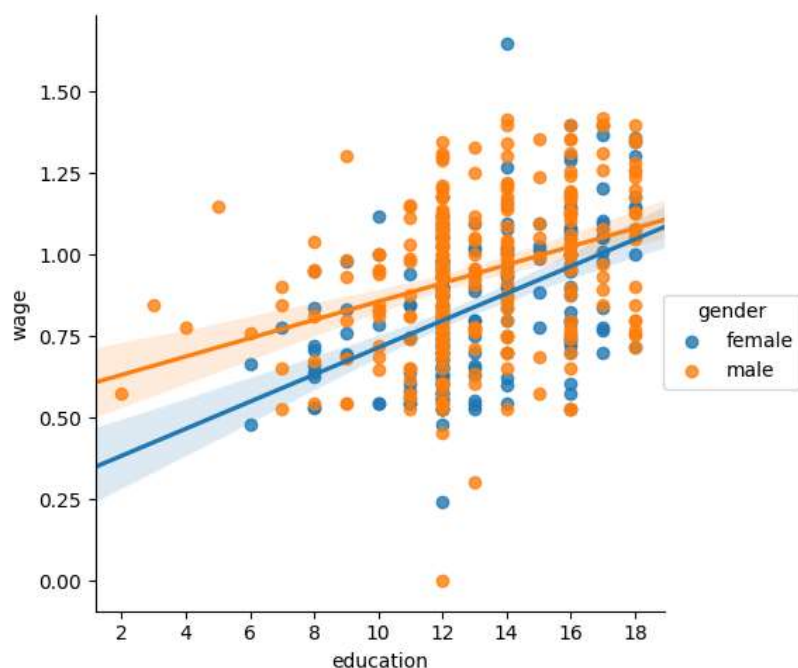



Robust regression

Given that, in the above plot, there seems to be a couple of data points that are outside of the main cloud to the right, they might be outliers, not representative of the population, but driving the regression.

To compute a regression that is less sensitive to outliers, one must use a [robust model](#). This is done in seaborn using `robust=True` in the plotting functions, or in statsmodels by replacing the use of the OLS by a “Robust Linear Model”, `statsmodels.formula.api.robustlm()`.

3.1.5. Testing for interactions



Do wages increase more with education for males than females?

The plot above is made of two different fits. We need to formulate a single model that tests for a variance of slope across the two populations. This is done via an “[interaction](#)”.


```
>>> result = sm.ols(formula='wage ~ education + gender + education * gender',
...                  data=data).fit()
>>> print(result.summary())
...
```

	coef	std err	t	P> t	[0.025	0.975]

Intercept	0.2998	0.072	4.173	0.000	0.159	0.441
gender[T.male]	0.2750	0.093	2.972	0.003	0.093	0.457
education	0.0415	0.005	7.647	0.000	0.031	0.052
education:gender[T.male]	-0.0134	0.007	-1.919	0.056	-0.027	0.000
=====	...					
...						

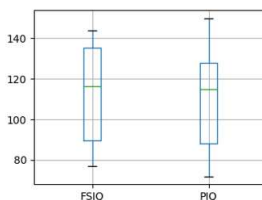
Can we conclude that education benefits males more than females?

Take home messages

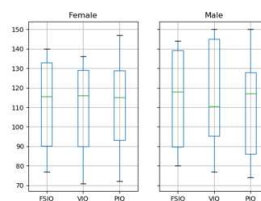
- Hypothesis testing and p-values give you the **significance** of an effect / difference.
- **Formulas** (with categorical variables) enable you to express rich links in your data.
- **Visualizing** your data and fitting simple models give insight into the data.
- **Conditioning** (adding factors that can explain all or part of the variation) is an important modeling aspect that changes the interpretation.

3.1.6. Full code for the figures

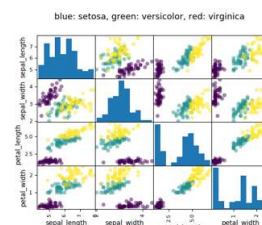
Code examples for the statistics chapter.



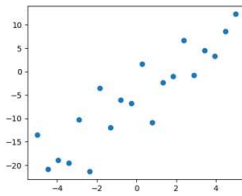
Boxplots and paired differences



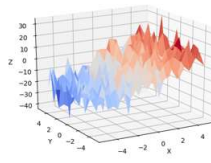
Plotting simple quantities of a pandas dataframe



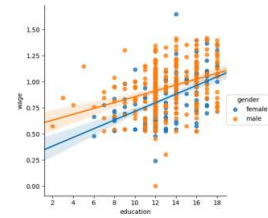
Analysis of Iris petal and sepal sizes



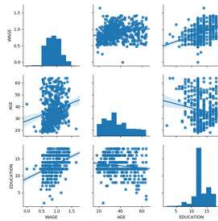
Simple Regression



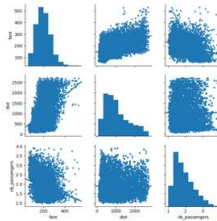
Multiple Regression



Test for an education/gender interaction in wages

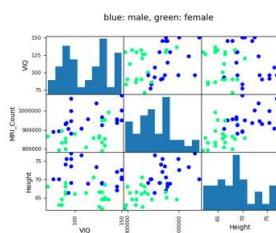


Visualizing factors influencing wages



Air fares before and after 9/11

3.1.7. Solutions to this chapter's exercises



Relating Gender and IQ

Download all examples in Python source code: `auto_examples_python.zip`

Download all examples in Jupyter notebooks: `auto_examples_jupyter.zip`