

Introduction to Python and Machine Learning

31st July 2019 Koh Wyhow

Agenda



- Introduction to Python
- Data visualization using seaborn

Creating statistical plots easily with seaborn

Hypothesis testing: z-test

Getting started with statistical hypothesis testing – a simple z-test

■ Correlation: Contingency table & chi-square test

Estimating the correlation between two variables with a contingency table and a chisquare test

Predictive analytics: Logistic regression

Logistic regression using Python

Predictive analytics: Natural Language Processing

Learning from text from Naïve Bayes for NLP

Predictive analytics: GCP AutoML

Introduction to Google Cloud Platform's AutoML

Recommendations



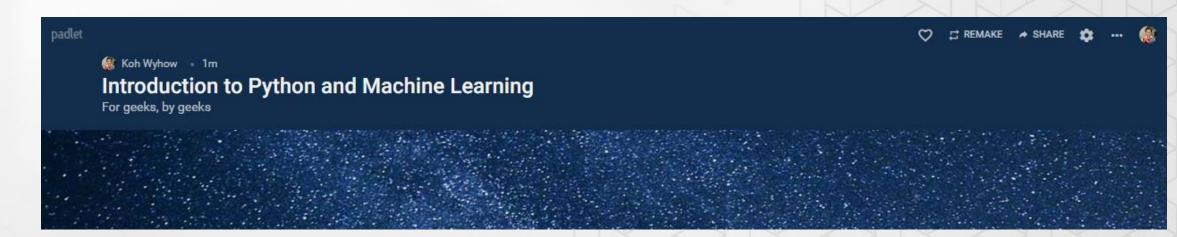
About me:

- Koh Wyhow (not SIR I have a name =P)
- Manager, Data Science @ Star Media Group
- Formerly:
 - Data Scientist @ INVOKE Malaysia
 - Consultant @ EY Data and Analytics
 - Further Mathematics lecturer @ Taylor's College
- Majored in Mathematics @ National University of Singapore, 2013



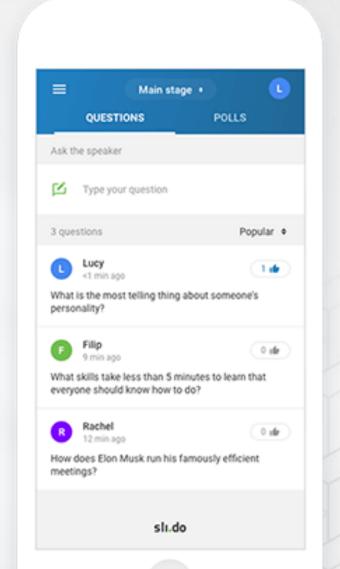
About you:

- https://padlet.com/kohwyhow/python20190731
 - What's your name?
 - What do you do?
 - What's your email?
 - How do you plan to use Python?



Questions:

■ www.sli.do code: #T458





Introduction to Python





Why Python?

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- Free!
- It's easy to learn (my first language was Java)
- Large community to support, and extensive libraries and documentation
- scikit-learn
- Giants are using it (e.g. NASA, Disney, Netflix, Electronic Arts, Google etc.)
- Reasons:
 - Machine learning and AI
 - Compatible with major platforms and systems
 - As a first language, easy to branch out to other languages
- Sources: PYPL, TIOBE

PYPL PopularitY of Programming Language

Worldwide, Jun 2019 compared to a year ago:

Rank	Change	Language	Share	Trend
1		Python	28.08 %	+4.7 %
2		Java	20.51 %	-1.8 %
3		Javascript	8.29 %	-0.2 %
4	1	C#	7.41 %	-0.5 %
5	Ψ.	PHP	6.96 %	-1.2 %
6		C/C++	5.76 %	-0.4 %

TIOBE Index for June 2019

June Headline: Python continues to soar in the TIOBE index

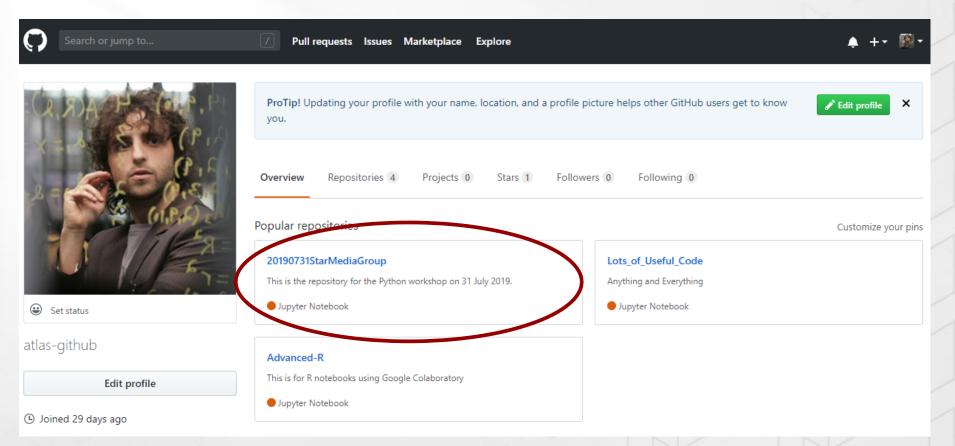
Jun 2019	Jun 2018	Change	Programming Language	Ratings
1	1		Java	15.004%
2	2		С	13.300%
3	4	^	Python	8.530%
4	3	v	C++	7.384%
5	6	^	Visual Basic .NET	4.624%
6	5	~	C#	4.483%
7	٩	^	lavaScrint	2 716%



GitHub:

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www.github.com/atlas-github



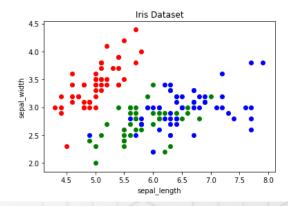
Data visualization using seaborn



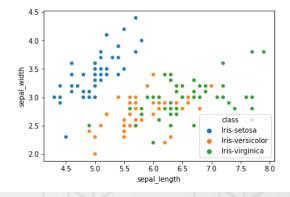
What is seaborn?

- Numerous other data visualization packages
 - Matplotlib, pandas visualization, ggplot, plotly
- Reasons to use seaborn
 - Can create graphs in 1 line of code which takes 10 in matplotlib
 - Awesome standard designs and colour palettes
 - Easy to learn <u>here</u>
- Source: Medium

```
# create color dictionary
colors = {'Iris-setosa':'r', 'Iris-versicolor':'g', 'Iris-virginica':'b'}
# create a figure and axis
fig, ax = plt.subplots()
# plot each data-point
for i in range(len(iris['sepal_length'])):
ax.scatter(iris['sepal_length'][i], iris['sepal_width'][i],color=colors[iris['class'][i]])
# set a title and labels
ax.set_title('Iris Dataset')
ax.set_title('Iris Dataset')
ax.set_vlabel('sepal_length')
ax.set_ylabel('sepal_width')
matplotlib_simple_scatterplot_with_colors.py hosted with \(\infty\) by GitHub
```



```
1 sns.scatterplot(x='sepal_length', y='sepal_width', hue='class', data=iris)
seaborn_simple_scatterplot_colored.py hosted with ♥ by GitHub view raw
```





How to use seaborn?

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seaborn.boxplot(x=None, y=None, hue=None, data=None, order=None, hue_order=None, orient=None, color=None, palette=None, saturation=0.75, width=0.8, dodge=True, fliersize=5, linewidth=None, whis=1.5, notch=False, ax=None, **kwargs)

Parameters:

x, y, hue: names of variables in data or vector data, optional Inputs for plotting long-form data. See examples for interpretation.

data: DataFrame, array, or list of arrays, optional Dataset for plotting. If x and y are absent, this is interpreted as wide-form. Otherwise it is expected to be long-form.

order, hue_order: lists of strings, optional
Order to plot the categorical levels in, otherwise the levels are inferred
from the data objects.

orient: "v" | "h", optional

Orientation of the plot (vertical or horizontal). This is usually inferred from the dtype of the input variables, but can be used to specify when the "categorical" variable is a numeric or when plotting wide-form data.

color: matplotlib color, optional Color for all of the elements, or seed for a gradient palette.

Thur Fri Sat Sun

Draw a vertical boxplot grouped by a categorical variable:

>>> ax = sns.boxplot(x="day", y="total_bill", data=tips)

Source: Seaborn

Hypothesis testing – z-test





Hypothesis-testing

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A few types:

Normality

Sample size > 30, population variance known

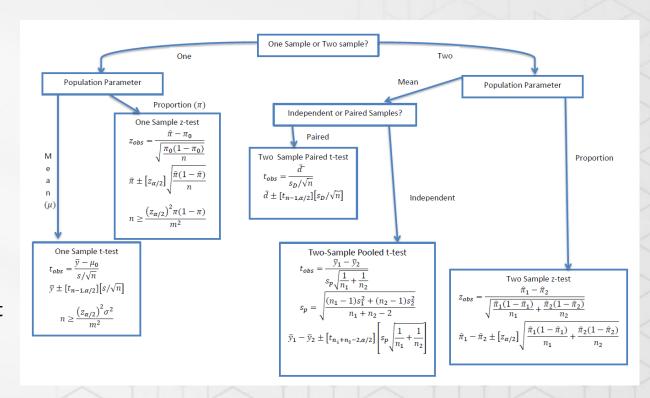
- T-test
 - Sample size < 30, population variance unknown
- Chi-square test

To test whether variables are independent

ANOVA

To analyse the differences among group means in a sample

■ Source: Towards Data Science





How to use statsmodels.stats?

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statsmodels.stats.proportion.proportions_ztest(count, nobs, value=None, alternative='two-sided', prop_var=False)

Parameters:

count: integer or array_like

the number of successes in nobs trials. If this is array_like, then the assumption is that this represents the number of successes for each independent sample

nobs: integer or array-like

the number of trials or observations, with the same length as count.

value: float, array_like or None, optional

This is the value of the null hypothesis equal to the proportion in the case of a one sample test. In the case of a two-sample test, the null hypothesis is that prop[0] - prop[1] = value, where prop is the proportion in the two samples. If not provided value = 0 and the null is prop[0] = prop[1]

alternative : string in ['two-sided', 'smaller', 'larger']

The alternative hypothesis can be either two-sided or one of the one-sided tests, smaller means that the alternative hypothesis is prop <value and larger means prop > value. In the two sample test, smaller means that the alternative hypothesis is p1 < p2 and larger means p1 > p2 where p1 is the proportion of the first sample and p2 of the second one.

Examples

```
>>> count = 5
>>> nobs = 83
>>> value = .05
>>> stat, pval = proportions_ztest(count, nobs, value)
>>> print('{0:0.3f}'.format(pval))
0.695

>>> import numpy as np
>>> from statsmodels.stats.proportion import proportions_ztest
>>> count = np.array([5, 12])
>>> nobs = np.array([83, 99])
>>> stat, pval = proportions_ztest(count, nobs)
>>> print('{0:0.3f}'.format(pval))
0.159
```

■ Source: <u>statsmodels.stats</u>

Contingency table & chi-square test





Contingency table and chi-square test

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Contingency table

- Aka. cross tabulation
- A table in a matrix format to display frequency distribution of variables
- Chi-square test
 - Aka. Pearson's chi-squared test
 - Used to determine whether there is significant difference between expected and observed frequencies in one or more categories
- Applications:
 - Data exploration
- Source: <u>Statistics How To</u>

AIDS * SEXPREF Crosstabulation

Count

	510	SEXPREF		
	Males	Females	Both	Total
AIDS Yes	4	2	3	9
No	3	16	2	21
Total	7	18	5	30

Chi-Square Tests

	Value	df	Asymp. Sig. (2-tailed)
Pearson Chi-Square	7.657 ^a	2	.022
Likelihood Ratio	7.803	2	.020
Linear-by-Linear Association	.062	1	.803
N of Valid Cases	30		

- a. 4 cells (66.7%) have expected count less than
 - 5. The minimum expected count is 1.50.



How to use scipy.stats?

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scipy.stats.chi2_contingency(observed, correction=True, lambda_=None)

Parameters:

bserved: array_like

The contingency table. The table contains the observed frequencies (i.e. number of occurrences) in each category. In the two-dimensional case, the table is often described as an "R x C table".

correction: bool, optional

If True, *and* the degrees of freedom is 1, apply Yates' correction for continuity. The effect of the correction is to adjust each observed value by 0.5 towards the corresponding expected value.

lambda:_float or str, optional.

By default, the statistic computed in this test is Pearson's chi-squared statistic. *lambda_* allows a statistic from the Cressie-Read power divergence family to be used instead. See power_divergence for details.

	Pizza Rolls	Chips & Dip	Cookies	Total
Poker	10	3	12	25
Trivial Pursuit	8	14	7	29
Monopoly	14	17	7	38
Wii Bowling	12	7	4	23
Total	44	41	30	115

Source: <u>scipy.stats</u>

Logistic regression

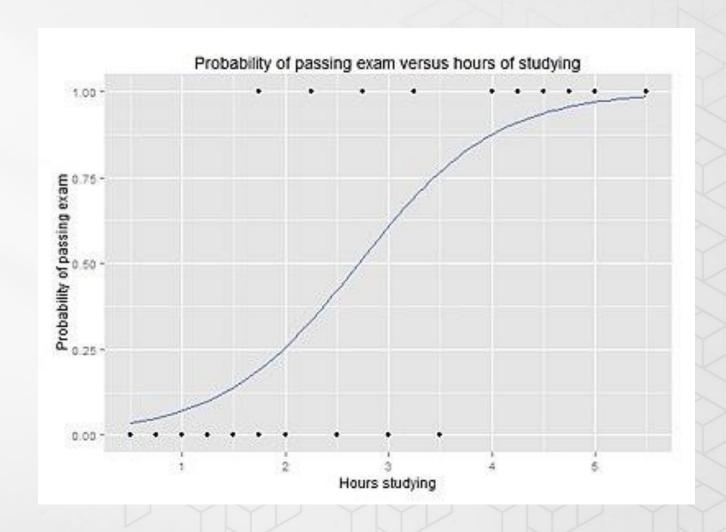




Logistic regression

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- Output is a probability a given input belongs to a certain class
- A type of classification algorithm
 - Binary labels: logistic regression
 - More than 2 labels: multinomial logistic regression
- Applications:
 - Profile behaviour
 - Which factors contribute to a certain health condition (yes vs. no)?





How to use sklearn.linear_model?

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sklearn.linear_model.LogisticRegression(penalty='12', dual=False, tol=0.0001, C=1.0, fit_intercept=True, intercept_scaling=1, class_weight=None, random_state=None, solver='warn', max_iter=100, multi_class='warn', verbose=0, warm_start=False, n_jobs=None, l1_ratio=None)

Parameters:

penalty: str, '11', '12', 'elasticnet' or 'none', optional (default='12') Used to specify the norm used in the penalization. The 'newton-cg', 'sag' and 'lbfgs' solvers support only I2 penalties. 'elasticnet' is only supported by the 'saga' solver. If 'none' (not supported by the liblinear solver), no regularization is applied.

New in version 0.19: I1 penalty with SAGA solver (allowing 'multinomial' + L1)

dual: bool, optional (default=False)

Dual or primal formulation. Dual formulation is only implemented for I2 penalty with liblinear solver. Prefer dual=False when n_samples > n_f eatures.

tol: float, optional (default=1e-4) Tolerance for stopping criteria.

C: float, optional (default=1.0)

Inverse of regularization strength; must be a positive float. Like in support vector machines, smaller values specify stronger regularization.

Examples

■ Source: sklearn.linear model.LogisticRegression

Natural Language Processing

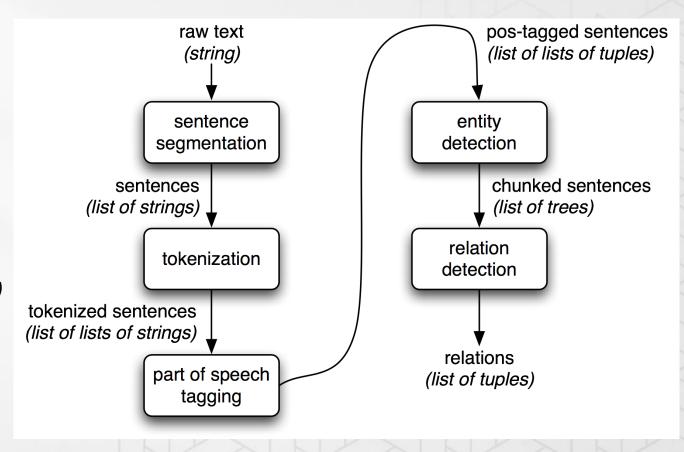




Natural Language Processing (NLP)

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- How to programme computers to process and analyse large amounts of unstructured text
- Applications:
 - Text classification
 - Behaviour profile of users
 - Taxonomy constructionNamed Entity Recognition
 - Natural Language Generation (WIP)
- Advanced applications:
 - <u>TalkToTransformer</u>
 - <u>BusUncle</u>
 - OpenAl GPT-2





How to use sklearn.naive_bayes?

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sklearn.naive_bayes. MultinomialNB(alpha=1.0, fit_prior=True, class_prior=None)

Parameters:

alpha: float, optional (default=1.0)

Additive (Laplace/Lidstone) smoothing parameter (0 for no smoothing).

fit_prior : boolean, optional (default=True)

Whether to learn class prior probabilities or not. If false, a uniform prior will be used.

class_prior: array-like, size (n_classes,), optional (default=None)
Prior probabilities of the classes. If specified the priors are not adjusted according to the data.

Examples

```
>>> import numpy as np
>>> X = np.random.randint(5, size=(6, 100))
>>> y = np.array([1, 2, 3, 4, 5, 6])
>>> from sklearn.naive_bayes import MultinomialNB
>>> clf = MultinomialNB()
>>> clf.fit(X, y)
MultinomialNB(alpha=1.0, class_prior=None, fit_prior=True)
>>> print(clf.predict(X[2:3]))
[3]
```

■ Source: sklearn.naive bayes.MultinomialNB

Google Cloud Platform's AutoML





Recommendations and Feedback

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- Need to continuously learn independently
- Meet up with others to look for ideas and inspiration
- Facebook:
 - TensorFlow & Deep Learning Malaysia
 - Malaysia R User Group (MYRUG)
- Meetup:
 - Kuala Lumpur Artificial Intelligence Meetup
- Workshop feedback:
 - https://www.surveymonkey.com/r/X6PT6GT



