Inference

COGS 108 Spring 2025

Jason Chen

Week 6

xic007@ucsd.edu

OH: Thu 3-5 pm

Discussion slides and materials adapted from Ruby Ying & previous quarter

Due dates

• D5 is due Friday, 05/09 @ 11:59 PM

Project Proposal

- Feedback will be out this week
- Make revisions in next checkpoint

D5: Inference

Part I: Data & Wrangling

df['height'] = df['height'].apply(function)

The apply() method allows you to apply a function along one of the axis of the DataFrame, default 0, which is the index (row) axis.

Syntax

dataframe.apply(func, axis, raw, result_type, args, kwds)

Parameters

The axis, raw, result_type, and args parameters are keyword arguments.

Parameter	Value	Description
func		Required. A function to apply to the DataFrame.
axis	0 1 'index' 'columns'	Optional, Which axis to apply the function to. default 0.
raw	True False	Optional, default False. Set to true if the row/column should be passed as an ndarray object
result_type	'expand' 'reduce' 'broadcast' None	Optional, default None. Specifies how the result wil be returned
args	a tuple	Optional, arguments to send into the function
kwds	keyword arguments	Optional, keyword arguments to send into the function

Part II: EDA

```
fig =
pd.plotting.scatter_matrix(
df[['column_1',
column_2']])
```

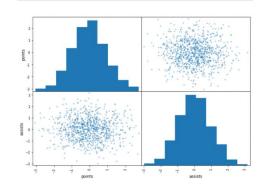
A scatter matrix is useful because it allows you to visualize the relationship between multiple variables in a dataset at once.

pandas.plotting.scatter_matrix

```
pandas.plotting.scatter_matrix(frame, alpha=0.5, figsize=None, ax=None,
grid=False, diagonal='hist', marker='.', density kwds=None, hist kwds=None,
range_padding=0.05, **kwargs)
    Draw a matrix of scatter plots.
    Parameters:
         frame : DataFrame
         alpha: float, optional
             Amount of transparency applied.
         figsize: (float,float), optional
             A tuple (width, height) in inches.
         ax : Matplotlib axis object, optional
         grid: bool, optional
             Setting this to True will show the grid.
         diagonal : ('hist', 'kde')
             Pick between 'kde' and 'hist' for either Kernel Density Estimation or Histogram plot in
             the diagonal.
         marker: str, optional
             Matplotlib marker type, default ::.
         density_kwds : keywords
             Keyword arguments to be passed to kernel density estimate plot.
         hist_kwds : keywords
             Keyword arguments to be passed to hist function.
         range padding: float, default 0.05
             Relative extension of axis range in x and y with respect to (x max - x min) or (y max -
         **kwargs
             Keyword arguments to be passed to scatter function.
     Returns:
         numpy.ndarray
             A matrix of scatter plots.
```

```
import pandas as pd
import numpy as np
#make this example reproducible
np.random.seed(0)
#create DataFrame
df = pd.DataFrame({'points': np.random.randn(1000),
                   'assists': np.random.randn(1000),
                   'rebounds': np.random.randn(1000)})
#view first five rows of DataFrame
df.head()
        points
                        assists
                                         rebounds
        1.764052
                        0.555963
                                         -1.532921
        0.400157
                        0.892474
                                         -1.711970
        0.978738
                         -0.422315
                                         0.046135
        2.240893
                        0.104714
                                         -0.958374
        1.867558
                        0.228053
                                         -0.080812
```

pd.plotting.scatter_matrix(df.iloc[:, 0:2])



Part III: ttest_ind

t_val, p_val = ttest_ind(df_1, df_2)

ttest_ind used to check whether the unknown population means of given pair of groups are equal.

tt allows one to test the null hypothesis that the means of two groups are equal

scipy.stats.ttest_ind

scipy.stats.ttest_ind(a, b, $axis=\theta$, $equal_var=True$, $nan_policy='propagate'$, permutations=None, $random_state=None$, alternative='two-sided', $trim=\theta$, *, keepdims=False) [source]

Calculate the T-test for the means of two independent samples of scores.

This is a test for the null hypothesis that 2 independent samples have identical average (expected) values. This test assumes that the populations have identical variances by default.

Parameters: a, b : array_like

The arrays must have the same shape, except in the dimension corresponding to axis (the first, by default).

axis: int or None, default: 0

If an int, the axis of the input along which to compute the statistic. The statistic of each axis-slice (e.g. row) of the input will appear in a corresponding element of the output. If None, the input will be raveled before computing the statistic.

equal var : bool, optional

If True (default), perform a standard independent 2 sample test that assumes equal population variances [1]. If False, perform Welch's t-test, which does not assume equal population variance [2].



nan_policy : {'propagate', 'omit', 'raise'}

Defines how to handle input NaNs.

- propagate: if a NaN is present in the axis slice (e.g. row) along which the statistic is computed, the corresponding entry of the output will be NaN.
- omit: NaNs will be omitted when performing the calculation. If insufficient data remains in the axis slice along which the statistic is computed, the corresponding entry of the output will be NaN.
- raise: if a NaN is present, a valueError will be raised.

Part III : statsmodels.formula.api.ols

mod_1 = statsmodels.formula.api.ols(formula,
data)

$$res_1 = mod_1.fit()$$

Introduction:

A linear regression model establishes the relation between a dependent variable (y) and at least one independent variable (x) as :

$$\hat{y} = b_1 x + b_0$$

In OLS method, we have to choose the values of b_1 and b_0 such that, the total sum of squares of the difference between the calculated and observed values of y, is minimised.

Formula for OLS:

$$S = \sum_{i=1}^{n} (y_i - \hat{y}_i)^2 = \sum_{i=1}^{n} (y_i - b_1 x_1 - b_0)^2 = \sum_{i=1}^{n} (\hat{\epsilon}_i)^2 = \min$$

Where.

 \hat{y}_i = predicted value for the ith observation

 y_i = actual value for the ith observation

&= error/residual for the ith observation

n = total number of observations

To get the values of b_0 and b_1 which minimise S, we can take a partial derivative for each coefficient and equate it to zero.

Section Materials

https://github.com/JasonC1217/COGS 108 B03-B04 Sp25/tree/master

or:

https://tinyurl.com/4d8wx3ne



THANKS!

Questions on EdStem or office hours

Office hours: Thu, 3-5 PM

UNFINISHED WORK





