Spring 2023 Data C100/C200 Midterm Reference Sheet

Pandas

Suppose df is a DataFrame; s is a Series. import pandas as pd

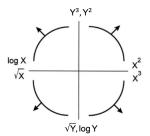
| Function | Description |
|--|---|
| df[col] | Returns the column labeled col from df as a Series. |
| df[[col1, col2]] | Returns a DataFrame containing the columns labeled col1 and col2. |
| s.loc[rows] / df.loc[rows, cols] | Returns a Series/DataFrame with rows (and columns) selected by their index values. |
| s.iloc[rows] / df.iloc[rows, cols] | Returns a Series/DataFrame with rows (and columns) selected by their positions. |
| s.isnull() / df.isnull() | Returns boolean Series/DataFrame identifying missing values |
| s.fillna(value) / df.fillna(value) | Returns a Series/DataFrame where missing values are replaced by value |
| s.isin(values) / df.isin(values) | Returns a Series/DataFrame of booleans indicating if each element is in values. |
| df.drop(labels, axis) | Returns a DataFrame without the rows or columns named labels along axis (either 0 or 1) |
| df.rename(index=None, columns=None) | Returns a DataFrame with renamed columns from a dictionary index and/or columns |
| df.sort_values(by, ascending=True) | Returns a DataFrame where rows are sorted by the values in columns by |
| s.sort_values(ascending=True) | Returns a sorted Series. |
| s.unique() | Returns a NumPy array of the unique values |
| s.value_counts() | Returns the number of times each unique value appears in a Series |
| <pre>pd.merge(left, right, how='inner', on='a')</pre> | Returns a DataFrame joining DataFrames left and right on the column labeled a; the join is of type inner |
| <pre>left.merge(right, left_on=col1, right_on=col2)</pre> | Returns a DataFrame joining DataFrames left and right on columns labeled col1 and col2. |
| <pre>df.pivot_table(index, columns, values=None, aggfunc='mean')</pre> | Returns a DataFrame pivot table where columns are unique values from columns (column name or list), and rows are unique values from index (column name or list); cells are collected values using aggfunc. If values is not provided, cells are collected for each remaining column with multi-level column indexing. |
| df.set_index(col) | Returns a DataFrame that uses the values in the column labeled col as the row index. |
| <pre>df.reset_index() Let grouped = df.groupby(by) where by can</pre> | Returns a DataFrame that has row index 0, 1, etc., and adds the current index as a column. be a column label or a list of labels. |
| Function | Description |
| | Description |
| grouped.count() | Return a Series containing the size of each group, excluding missing values |
| grouped.size() | Return a Series containing size of each group, including missing values |
| <pre>grouped.mean()/grouped.min()/grouped.max()</pre> | Return a Series/DataFrame containing mean/min/max of each group for each column, excluding missing values |
| <pre>grouped.filter(f) grouped.agg(f)</pre> | Filters or aggregates using the given function f |
| Function | Description |
| s.str.len() | Returns a Series containing length of each string |
| s.str[a:b] | Returns a Series where each element is a slice of the corresponding string indexed from a (inclusive, optional) to b (non-inclusive, optional) |
| s.str.lower()/s.str.upper() | Returns a Series of lowercase/uppercase versions of each string |
| s.str.replace(pat, repl) | Returns a Series that replaces occurences of substrings matching the regex pat with string repl |
| s.str.contains(pat) | Returns a boolean Series indicating if a substring matching the regex pat is contained in each string |
| s.str.extract(pat) | Returns a Series of the first subsequence of each string that matches the regex pat. If pat contains one group, then only the substring matching the group is extracted |

Visualization

Matplotlib: x and y are sequences of values. import matplotlib.pyplot as plt

| Function | Description |
|-----------------------------------|--|
| plt.plot(x, y) | Creates a line plot of x against y |
| <pre>plt.scatter(x, y)</pre> | Creates a scatter plot of x against y |
| <pre>plt.hist(x, bins=None)</pre> | Creates a histogram of x; bins can be an integer or a sequence |
| <pre>plt.bar(x, height)</pre> | Creates a bar plot of categories x and corresponding heights |

Tukey-Mosteller Bulge Diagram.



Seaborn: x and y are column names in a DataFrame data. import seaborn as sns

| Function | Description |
|--|--|
| <pre>sns.countplot(data, x)</pre> | Create a barplot of value counts of variable x from data |
| <pre>sns.histplot(data, x, stat='count', kde=False) sns.displot(x, data, stat='count', rug=True, kde=True)</pre> | Creates a histogram of x from data, where bin statistics stat is one of 'count', 'frequency', 'probability', 'percent', and 'density'; optionally overlay a kernel density estimator. displot is similar but can optionally overlay a rug plot |
| <pre>sns.boxplot(data, x=None, y) sns.violinplot(data, x=None, y)</pre> | Create a boxplot of y, optionally factoring by categorical x, from data. violinplot is similar but also draws a kernel density estimator of y |
| <pre>sns.rugplot(data, x)</pre> | Adds a rug plot on the x-axis of variable x from data |
| <pre>sns.scatterplot(data, x, y)</pre> | Create a scatterplot of x versus y from data |
| <pre>sns.lmplot(x, y, data, fit_reg=True)</pre> | Create a scatterplot of $\mathbf x$ versus $\mathbf y$ from data, and by default overlay a least-squares regression line |
| <pre>sns.jointplot(x, y, data, kind)</pre> | Combine a bivariate scatterplot of x versus y from data, with univariate density plots of each variable overlaid on the axes; kind determines the visualization type for the distribution plot, can be scatter, kde or hist |

Regular Expressions

| Operator | Description | Operator | Description |
|------------|--|----------|--|
| | Matches any character except \n | * | Matches preceding character/group zero or more times |
| \ | Escapes metacharacters | ? | Matches preceding character/group zero or one times |
| 1 | Matches expression on either side of expression; has lowest priority of any operator | + | Matches preceding character/group one or more times |
| \d, \w, \s | Predefined character group of digits (0-9), alphanumerics (a-z, A-Z, 0-9, and underscore), or whitespace, respectively | ^, \$ | Matches the beginning and end of the line, respectively |
| \D, \W, \S | Inverse sets of \d, \w, \s, respectively | () | Capturing group used to create a sub-expression |
| {m} | Matches preceding character/group exactly m times | [] | Character class used to match any of the specified characters or range (e.g. [abcde] is equivalent to [a-e]) |
| {m, n} | Matches preceding character/group at least m times and at most n times if either m or n are omitted, set lower/upper bounds to 0 and ∞ , respectively | [^] | Invert character class; e.g. [^a-c] matches all characters except a, b, c |

Modified lecture example for capture groups:

```
import re
lines = '169.237.46.168 - - [26/Jan/2014:10:47:58 -0800] "GET ... HTTP/1.1"'
re.findall(r'\[\d+\/(\w+)\/\d+:\d+:\d+ .+\]', line) # returns ['Jan']
```

| Function | Description |
|--|--|
| re.match(pattern, string) | Returns a match if zero or more characters at beginning of string matches pattern, else None |
| re.search(pattern, string) | Returns a match if zero or more characters anywhere in string matches pattern, else None |
| re.findall(pattern, string) | Returns a list of all non-overlapping matches of pattern in string (if none, returns empty list) |
| <pre>re.sub(pattern, repl, string)</pre> | Returns string that replaces all occurrences of pattern with repl |

Modeling

| Concept | Formula | Concept | Formula |
|---------------------------------|--|--|---|
| Variance, σ_x^2 | $\frac{1}{n}\sum_{i=1}^n (x_i - \bar{x})^2$ | Correlation r | $r = rac{1}{n} \sum_{i=1}^{n} rac{x_i - ar{x}}{\sigma_x} rac{y_i - ar{y}}{\sigma_y}$ |
| L_1 loss | $L_1(y,\hat{y}) = \mid y - \hat{y} \mid$ | Linear regression estimate of \boldsymbol{y} | $\hat{y}=	heta_0+	heta_1 x$ |
| L_2 loss | $L_2(y,\hat{y})=(y-\hat{y})^2$ | Least squares linear regression | $\hat{	heta}_0 = ar{y} - \hat{	heta}_1 ar{x} \qquad \hat{	heta}_1 = r rac{\sigma}{\sigma}$ |
| npirical risk with loss $\it L$ | $R(heta) = rac{1}{n} \sum_{i=1}^n L(y_i, \hat{y_i})$ | | |

Ordinary Least Squares

Multiple Linear Regression Model: $\hat{\mathbb{Y}} = \mathbb{X}\theta$ with design matrix \mathbb{X} , response vector \mathbb{Y} , and predicted vector $\hat{\mathbb{Y}}$. If there are p features plus a bias/intercept, then the vector of parameters $\theta = [\theta_0, \theta_1, \dots, \theta_p]^T \in \mathbb{R}^{p+1}$. The vector of estimates $\hat{\theta}$ is obtained from fitting the model to the sample (\mathbb{X}, \mathbb{Y}) .

| Concept | Formula | Concept | Formula |
|----------------------|--|---|---|
| Mean squared error | $R(heta) = rac{1}{n} \ \mathbb{Y} - \mathbb{X}	heta\ _2^2$ | Normal equation | $\mathbb{X}^T\mathbb{X}\hat{\theta}=\mathbb{X}^T\mathbb{Y}$ |
| | | Least squares estimate, if $\ensuremath{\mathbb{X}}$ is full rank | $\hat{\theta} = (\mathbb{X}^T \mathbb{X})^{-1} \mathbb{X}^T \mathbb{Y}$ |
| Residual vector, e | $e=\mathbb{Y}-\hat{\mathbb{Y}}$ | Multiple ${\cal R}^2$ (coefficient of determination) | $R^2 = rac{	ext{variance of fitted values}}{	ext{variance of } y}$ |