Spring 2023 Data C100/C200 Midterm Reference Sheet

Pandas

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

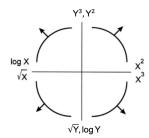
Function	Description
df[col]	Returns the column labeled col from df as a Series.
df[[col1, col2]]	Returns a DataFrame containing the columns labeled col1 and col2.
s.loc[rows] / df.loc[rows, cols]	Returns a Series/DataFrame with rows (and columns) selected by their index values.
s.iloc[rows] / df.iloc[rows, 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 height

Tukey-Mosteller Bulge Diagram.



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

Function	Description
sns.countplot(data, x)	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
sns.boxplot(data, x=None, y) sns.violinplot(data, x=None, y)	Create a boxplot of y, optionally factoring by categorical x, from data. violinplot is similar but also draws a kernel density estimator of y
sns.scatterplot(data, x, y)	Create a scatterplot of x versus y from data
<pre>sns.lmplot(x, y, data, fit_reg=True)</pre>	Create a scatterplot of ${\bf x}$ versus ${\bf y}$ from ${\tt data}$, and by default overlay a least-squares regression line
sns.jointplot(x, y, data, kind)	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
I	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)
re sub(nattern renl string)	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	$L_1(y,\hat{y}) = \mid y - \hat{y} \mid$		$\hat{y}=\theta_0+\theta_1 x$
L_2 loss	$L_2(y,\hat{y})=(y-\hat{y})^2$	Least squares linear regression	$\hat{ heta}_0 = ar{y} - \hat{ heta}_1 ar{x} \qquad \hat{ heta}_1 = r rac{\sigma_{ar{z}}}{\sigma_{ar{z}}}$
mpirical 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	$\mathbb{X}^T \mathbb{X} \hat{ heta} = \mathbb{X}^T \mathbb{Y}$
		Least squares estimate, if $\mathbb X$ 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)	$R^2 = rac{ ext{variance of fitted values}}{ ext{variance of } y}$