# Spring 2023 Data C100/C200 Midterm Reference Sheet

## Pandas

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

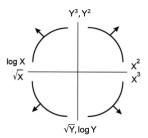
Function	Description
df[col]	Returns the column labeled col from df as a Series.
df[[col1, col2]]	Returns a DataFrame containing the columns labeled col1 and col2.
s.loc[rows] / df.loc[rows, cols]	Returns a Series/DataFrame with rows (and columns) selected by their index values.
s.iloc[rows] / df.iloc[rows, 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
plt.hist(x, bins=None)	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 ${\tt data}$ , 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 $\infty$ , 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(pattern, repl, string)	Returns string that replaces all occurrences of pattern with repl

## Modeling

Concept	Formula	Concept	Formula	
Variance, $\sigma_x^2$	Variance, $\sigma_x^2$ $\frac{1}{n} \sum_{i=1}^n (x_i - \bar{x})^2$		relation $r$ $r = rac{1}{n} \sum_{i=1}^n rac{x_i - ar{x}}{\sigma_x} rac{y_i - ar{y}}{\sigma_y}$	
$L_1$ 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)	$R^2 = rac{ ext{variance of fitted values}}{ ext{variance of } y}$