datascience Documentation

Release 0.5.12

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Release 0.5.12

Date March 08, 2016

The datascience package was written for use in Berkeley's DS 8 course and contains useful functionality for investigating and graphically displaying data.

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Start Here: datascience Tutorial

This is a brief introduction to the functionality in datascience. For a complete reference guide, please see *Tables* (*datascience.tables*).

For other useful tutorials and examples, see:

- The textbook introduction to Tables
- Example notebooks

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1.1 Getting Started

The most important functionality in the package is is the Table class, which is the structure used to represent columns of data. First, load the class:

```
In [1]: from datascience import Table
```

In the IPython notebook, type Table. followed by the TAB-key to see a list of members.

Note that for the Data Science 8 class we also import additional packages and settings for all assignments and labs. This is so that plots and other available packages mirror the ones in the textbook more closely. The exact code we use is:

```
import matplotlib
matplotlib.use('Agg')
from datascience import Table
%matplotlib inline
import matplotlib.pyplot as plt
```

```
import numpy as np
plt.style.use('fivethirtyeight')
```

In particular, the lines involving matplotlib allow for plotting within the IPython notebook.

1.2 Creating a Table

A Table is a sequence of labeled columns of data.

A Table can be constructed from scratch by extending an empty table with columns.

```
In [2]: t = Table().with_columns([
             'letter', ['a', 'b', 'c', 'z'],
   . . . :
            'count', [ 9, 3, 3, 'points', [ 1, 2, 2,
   . . . :
                                     2, 10],
   . . . :
   ...: ])
   . . . :
In [3]: print(t)
letter | count | points
              | 1
       | 9
        1 3
               | 2
        | 3
               | 2
        | 1
               | 10
```

More often, a table is read from a CSV file (or an Excel spreadsheet). Here's the content of an example file:

```
In [4]: cat sample.csv
x,y,z
1,10,100
2,11,101
3,12,102
```

And this is how we load it in as a Table using read_table():

CSVs from URLs are also valid inputs to read_table():

```
In [6]: Table.read_table('http://data8.org/textbook/notebooks/sat2014.csv')
Out[6]:
          | Participation Rate | Critical Reading | Math | Writing | Combined
State
                              | 612
                                                | 620 | 584
                                                             | 1816
North Dakota | 2.3
                               | 599
                                                | 616 | 587
                                                               | 1802
Illinois | 4.6
                               | 605
                                                | 611 | 578
                                                               | 1794
          | 3.1
South Dakota | 2.9
                               1 604
                                                | 609 | 579
                                                               | 1792
Minnesota | 5.9
                               | 598
                                                | 610 | 578
                                                               | 1786
          | 3.8
                               1 593
Michigan
                                                | 610 | 581
                                                                | 1784
Wisconsin | 3.9
                               1 596
                                                | 608 | 578
                                                                | 1782
Missouri
           | 4.2
                               | 595
                                                | 597
                                                      | 579
                                                                | 1771
          | 3.3
                               | 590
                                                | 599 | 573
                                                                | 1762
Wyoming
```

```
Kansas | 5.3 | 591 | 596 | 566 | 1753 | ... (41 rows omitted)
```

It's also possible to add columns from a dictionary, but this option is discouraged because dictionaries do not preserve column order.

```
In [7]: t = Table().with_columns({
            'letter': ['a', 'b', 'c', 'z'],
   . . . :
            'count': [ 9, 3, 3,
                                        1],
   . . . :
            'points': [ 1, 2,
                                  2, 10],
   . . . :
   ...: })
   . . . :
In [8]: print(t)
count | letter | points
9
      | a
              | 1
3
      | b
               | 2
3
               | 2
      | C
               | 10
1
      | z
```

1.3 Accessing Values

To access values of columns in the table, use column (), which takes a column label or index and returns an array. Alternatively, columns () returns a list of columns (arrays).

```
In [9]: t
Out[9]:
count | letter | points
     | 1
  | a
   | b
       | 2
3
3
   | C
       | 2
   | z
       | 10
In [10]: t.column('letter')
array(['a', 'b', 'c', 'z'],
   dtype='<U1')
In [11]: t.column(1)
array(['a', 'b', 'c', 'z'],
   dtype='<U1')
```

You can use bracket notation as a shorthand for this method:

To access values by row, row() returns a row by index. Alternatively, rows() returns an list-like Rows object that contains tuple-like Row objects.

```
In [14]: t.rows
Out [14]:
Rows (count | letter | points
     | 1
  | a
  | b
      1 2
3
  l c
      1 2
      | 10)
  l z
In [15]: t.rows[0]
In [16]: t.row(0)
In [17]: second = t.rows[1]
In [18]: second
Out[18]: Row(count=3, letter='b', points=2)
In [19]: second[0]
In [20]: second[1]
```

To get the number of rows, use num_rows.

```
In [21]: t.num_rows
Out[21]: 4
```

1.4 Manipulating Data

Here are some of the most common operations on data. For the rest, see the reference (Tables (datascience.tables)).

Adding a column with with_column():

```
In [22]: t
Out [22]:
count | letter | points
   | a
         | 1
   | b
         1 2
   | C
         | 2
         | 10
In [23]: t.with_column('vowel?', ['yes', 'no', 'no', 'no'])
count | letter | points | vowel?
   | a
        | 1
              | yes
3
   l b
         1 2
              l no
3
         | 2
   | C
              | no
   | z
         | 10
              l no
In [24]: t # .with_column returns a new table without modifying the original
```

```
count | letter | points
       | 1
   | a
         | 2
   l b
         | 2
   | C
   Z
         | 10
In [25]: t.with_column('2 * count', t['count'] * 2) # A simple way to operate on columns
count | letter | points | 2 * count
   | a
        | 1
              | 18
   | b
         | 2
                | 6
3
   l c
         | 2
                | 6
         | 10
               | 2
   | z
```

Selecting columns with select ():

Renaming columns with relabeled():

```
In [28]: t
Out [28]:
count | letter | points
  | a
      | 1
   | b
        | 2
3
   l c
        | 2
   | z
        | 10
In [29]: t.relabeled('points', 'other name')
count | letter | other name
      | 1
   l a
        | 2
3
   l b
3
        | 2
   | C
        | 10
   | Z
In [30]: t
count | letter | points
   Ιa
       | 1
3
   l b
        | 2
3
        | 2
   | C
        | 10
In [31]: t.relabeled(['letter', 'count', 'points'], ['x', 'y', 'z'])
```

Selecting out rows by index with take() and conditionally with where():

```
In [32]: t
Out [32]:
count | letter | points
  | a
      | 1
  l b
       1 2
  | C
       | 2
3
       | 10
1
  | Z
In [33]: t.take(2) # the third row
count | letter | points
3 | c | 2
In [34]: t.take[0:2] # the first and second rows
count | letter | points
  | a | 1
   | b
       | 2
```

```
In [35]: t.where('points', 2) # rows where points == 2
count | letter | points
 l b
      | 2
  l c
        1 2
In [36]: t.where(t['count'] < 8) # rows where count < 8</pre>
count | letter | points
      | 2
  l b
      | 2
  | C
       | 10
  | Z
In [37]: t['count'] < 8 # .where actually takes in an array of booleans</pre>
In [38]: t.where([False, True, True, True]) # same as the last line
count | letter | points
3 | b | 2
   | C
        | 2
1
   Z
        | 10
```

Operate on table data with sort (), group (), and pivot ()

```
1 | z | 10
In [40]: t.sort('count')
count | letter | points
     | 10
  Z
      | 2
  | b
3
  | C
      | 2
9
  | a
      | 1
In [41]: t.sort('letter', descending = True)
count | letter | points
  | z
     | 10
       | 2
  | C
  | b
       | 2
9
       | 1
  | a
```

```
In [43]: other_table = Table().with_columns([
  ....: 'mar_status', ['married', 'married', 'partner', 'partner', 'married'],
           'empl_status', ['Working as paid', 'Working as paid', 'Not working',
                        'Not working', 'Not working'],
  . . . . :
                      [1, 1, 1, 1, 1]])
         'count',
  . . . . :
  . . . . :
In [44]: other_table
Out [44]:
mar_status | empl_status
                     | count
married | Working as paid | 1
married
         | Working as paid | 1
partner | Not working
                     | 1
partner | Not working
                        1 1
married | Not working
                       | 1
In [45]: other_table.pivot('mar_status', 'empl_status', 'count', collect=sum)
| married | partner
empl status
             | 1
                   | 2
Not working
                     | 0
Working as paid | 2
```

1.5 Visualizing Data

We'll start with some data drawn at random from two normal distributions:

Draw histograms with hist():

```
In [48]: normal_data.hist()

_build/latex/_images/hist.png
```

```
In [49]: normal_data.hist(bins = range(-5, 10))
```



If we treat the normal_data table as a set of x-y points, we can plot () and scatter():

1	[n [51]: no	rmal_da	ta.sort('	data1')	.plot('data1')	# S	ort	first	to	make	plot	nicer	
	_build/la	tex/_im	ages/plo	ot.png										
	[n [52]: no	rmal da	ta scatte	r(!data	1!)									
ᅼ	nc	ormar_da.	La.SCATTE	:r (. agta	.± ')									
-	_build/la	tex/_im	ages/sca	utter.p	ng									

```
In [53]: normal_data.scatter('datal', fit_line = True)
_build/latex/_images/scatter_line.png
```

Use barh () to display categorical data.

```
_build/latex/_images/barh.png
```

1.6 Exporting

Exporting to CSV is the most common operation and can be done by first converting to a pandas dataframe with $to_df()$:

```
In [56]: normal_data
Out [56]:
data1
        | data2
1.89785 | 4.30097
0.274448 | 7.70979
3.59141 | 2.11838
2.22658 | 4.83009
-4.00548 | 5.2149
6.84537 | 2.66324
3.23994 | -0.0205213
1.62204 | 5.3063
4.20877 | 6.93846
1.39406 | 3.24598
... (90 rows omitted)
# index = False prevents row numbers from appearing in the resulting CSV
In [57]: normal_data.to_df().to_csv('normal_data.csv', index = False)
```

1.7 An Example

We'll recreate the steps in Chapter 3 of the textbook to see if there is a significant difference in birth weights between smokers and non-smokers using a bootstrap test.

For more examples, check out the TableDemos repo.

From the text:

The table baby contains data on a random sample of 1,174 mothers and their newborn babies. The column birthwt contains the birth weight of the baby, in ounces; gest_days is the number of gestational days, that is, the number of days the baby was in the womb. There is also data on maternal age, maternal height, maternal pregnancy weight, and whether or not the mother was a smoker.

```
In [58]: baby = Table.read_table('http://data8.org/textbook/notebooks/baby.csv')
In [59]: baby # Let's take a peek at the table
Out [59]:
Birth Weight | Gestational Days | Maternal Age | Maternal Height | Maternal Pregnancy Weight | Maternal
          | 284
                          | 27
                                       | 62
                                                       | 100
                                                                                | False
                                                        | 135
113
          | 282
                          1 33
                                        1 64
                                                                                | False
128
          1 279
                          | 28
                                        | 64
                                                       | 115
                                                                                 | True
                                                        | 125
                                                                                 | True
108
          | 282
                          1 23
                                        1 67
          | 286
                                                        | 93
                          | 25
136
                                        | 62
                                                                                I False
                          | 33
                                                        | 178
138
          | 244
                                        | 62
                                                                                | False
                                                        | 140
132
           | 245
                           | 23
                                        | 65
                                                                                 | False
120
           1 289
                           1 25
                                        1 62
                                                        | 125
                                                                                 | False
           | 299
                           | 30
                                        | 66
143
                                                        | 136
                                                                                | True
140
          | 351
                          | 27
                                       | 68
                                                        | 120
                                                                                 | False
... (1164 rows omitted)
# Select out columns we want.
In [60]: smoker_and_wt = baby.select(['m_smoker', 'birthwt'])
KevError
                                    Traceback (most recent call last)
<ipython-input-60-98d2eeccb488> in <module>()
----> 1 smoker_and_wt = baby.select(['m_smoker', 'birthwt'])
../datascience/tables.py in select(self, column_label_or_labels)
   559
        table = Table()
   560
             for label in labels:
--> 561
                self._add_column_and_format(table, label, np.copy(self[label]))
   562
             return table
   563
../datascience/tables.py in __getitem__(self, index_or_label)
   def __getitem__(self, index_or_label):
             label = self._as_label(index_or_label)
   164
--> 165
             return self.column(label)
   166
   167
          def __setitem__(self, label, values):
../datascience/tables.py in column(self, index_or_label)
   261
                An instance of ``numpy.array`
   2.62
--> 263
             return self._columns[self._as_label(index_or_label)]
   264
   265
          # Deprecated
KeyError: 'm_smoker'
In [61]: smoker_and_wt
NameError
                                    Traceback (most recent call last)
```

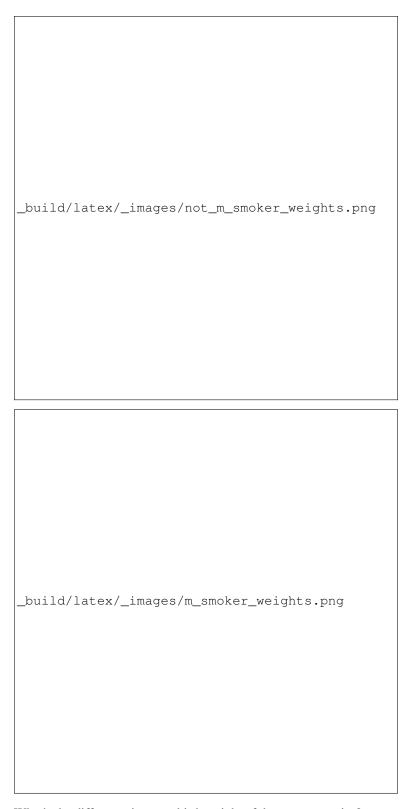
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```
<ipython-input-61-1c8046ed122a> in <module>()
----> 1 smoker_and_wt
NameError: name 'smoker_and_wt' is not defined
```

Let's compare the number of smokers to non-smokers.

```
In [62]: smoker_and_wt.select('m_smoker').hist(bins = [0, 1, 2]);
__build/latex/_images/m_smoker.png
```

We can also compare the distribution of birthweights between smokers and non-smokers.



What's the difference in mean birth weight of the two categories?

1.7. An Example

```
<ipython-input-65-f41eeafe6bd7> in <module>()
---> 1 nonsmoking_mean = smoker_and_wt.where('m_smoker', 0).column('birthwt').mean()
NameError: name 'smoker_and_wt' is not defined
In [66]: smoking_mean = smoker_and_wt.where('m_smoker', 1).column('birthwt').mean()
NameError
                               Traceback (most recent call last)
<ipython-input-66-025a08c0d2c6> in <module>()
----> 1 smoking_mean = smoker_and_wt.where('m_smoker', 1).column('birthwt').mean()
NameError: name 'smoker_and_wt' is not defined
In [67]: observed_diff = nonsmoking_mean - smoking_mean
NameError
                               Traceback (most recent call last)
<ipython-input-67-1dc0201ce02b> in <module>()
----> 1 observed_diff = nonsmoking_mean - smoking_mean
NameError: name 'nonsmoking_mean' is not defined
In [68]: observed_diff
Traceback (most recent call last)
NameError
<ipython-input-68-dbee69efdaa6> in <module>()
----> 1 observed_diff
NameError: name 'observed_diff' is not defined
```

Let's do the bootstrap test on the two categories.

```
In [69]: num_nonsmokers = smoker_and_wt.where('m_smoker', 0).num_rows
NameError
                                          Traceback (most recent call last)
<ipython-input-69-66887dbcfe42> in <module>()
---> 1 num_nonsmokers = smoker_and_wt.where('m_smoker', 0).num_rows
NameError: name 'smoker_and_wt' is not defined
In [70]: def bootstrap_once():
  . . . . :
           Computes one bootstrapped difference in means.
   . . . . :
           The table.sample method lets us take random samples.
           We then split according to the number of nonsmokers in the original sample
  . . . . :
           resample = smoker_and_wt.sample(with_replacement = True)
           bootstrap_diff = resample.column('birthwt')[:num_nonsmokers].mean() - \
  ...:
               resample.column('birthwt')[num_nonsmokers:].mean()
   . . . . :
           return bootstrap_diff
   . . . . :
In [71]: repetitions = 1000
In [72]: bootstrapped_diff_means = np.array(
           [ bootstrap_once() for _ in range(repetitions) ])
  . . . . :
   . . . . :
                                          Traceback (most recent call last)
```

```
<ipython-input-72-167c9ef03394> in <module>()
    1 bootstrapped_diff_means = np.array(
         [ bootstrap_once() for _ in range(repetitions) ])
<ipython-input-72-167c9ef03394> in <listcomp>(.0)
    1 bootstrapped_diff_means = np.array(
         [ bootstrap_once() for _ in range(repetitions) ])
<ipython-input-70-0af534563c70> in bootstrap_once()
         We then split according to the number of nonsmokers in the original sample.
    6
 --> 7
        resample = smoker_and_wt.sample(with_replacement = True)
        bootstrap_diff = resample.column('birthwt')[:num_nonsmokers].mean() -
                                                                        resample.co
        return bootstrap_diff
NameError: name 'smoker_and_wt' is not defined
In [73]: bootstrapped_diff_means[:10]
Traceback (most recent call last)
NameError
<ipython-input-73-7ce34a8f5ddf> in <module>()
----> 1 bootstrapped_diff_means[:10]
NameError: name 'bootstrapped_diff_means' is not defined
In [74]: num_diffs_greater = (abs(bootstrapped_diff_means) > abs(observed_diff)).sum()
NameError
                                 Traceback (most recent call last)
<ipython-input-74-fea00db6f1ce> in <module>()
---> 1 num_diffs_greater = (abs(bootstrapped_diff_means) > abs(observed_diff)).sum()
NameError: name 'bootstrapped_diff_means' is not defined
In [75]: p_value = num_diffs_greater / len(bootstrapped_diff_means)
Traceback (most recent call last)
NameError
<ipython-input-75-053c4de37c34> in <module>()
---> 1 p_value = num_diffs_greater / len(bootstrapped_diff_means)
NameError: name 'num_diffs_greater' is not defined
In [76]: p_value
Traceback (most recent call last)
<ipython-input-76-1336a4a7d880> in <module>()
----> 1 p_value
NameError: name 'p_value' is not defined
```

1.8 Drawing Maps

To come.

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Reference

2.1 Tables (datascience.tables)

Summary of methods for Table. Click a method to see its documentation.

One note about reading the method signatures for this page: each method is listed with its arguments. However, optional arguments are specified in brackets. That is, a method that's documented like

```
Table.foo (first_arg, second_arg[, some_other_arg, fourth_arg])
```

means that the Table.foo method must be called with first_arg and second_arg and optionally some_other_arg and fourth_arg. That means the following are valid ways to call Table.foo:

```
some_table.foo(1, 2)
some_table.foo(1, 2, 'hello')
some_table.foo(1, 2, 'hello', 'world')
some_table.foo(1, 2, some_other_arg='hello')
```

But these are not valid:

```
some_table.foo(1) # Missing arg
some_table.foo(1, 2[, 'hi']) # SyntaxError
some_table.foo(1, 2[, 'hello', 'world']) # SyntaxError
```

If that syntax is confusing, you can click the method name itself to get to the details page for that method. That page will have a more straightforward syntax.

At the time of this writing, most methods only have one or two sentences of documentation, so what you see here is all that you'll get for the time being. We are actively working on documentation, prioritizing the most complicated methods (mostly visualizations).

Creation

<pre>Tableinit([labels, _deprecated, formatter])</pre>	Create an empty table with column labels.		
Table.from_records(records)	Create a table from a sequence of records (dicts with fixed keys).		
Table.read_table(filepath_or_buffer, *args,)	Read a table from a file or web address.		
Table.from_df(df)	Convert a Pandas DataFrame into a Table.		
Table.from_array(arr)	Convert a structured NumPy array into a Table.		

2.1.1 datascience.tables.Table.__init__

Table.__init__ (labels=None, _deprecated=None, *, formatter=<datascience.formats.Formatter object>)

Create an empty table with column labels.

```
>>> tiles = Table(['letter', 'count', 'points'])
>>> tiles
letter | count | points
```

Args: labels (list of strings): The column labels.

formatter (Formatter): An instance of Formatter that formats the columns' values.

2.1.2 datascience.tables.Table.from_records

```
classmethod Table.from_records (records)
```

Create a table from a sequence of records (dicts with fixed keys).

2.1.3 datascience.tables.Table.read table

classmethod Table.read_table (filepath_or_buffer, *args, **vargs)

Read a table from a file or web address.

filepath_or_buffer – string or file handle / StringIO; The string could be a URL. Valid URL schemes include http, ftp, s3, and file.

2.1.4 datascience.tables.Table.from df

```
classmethod Table.from_df(df)
```

Convert a Pandas DataFrame into a Table.

2.1.5 datascience.tables.Table.from_array

classmethod Table.from_array(arr)

Convert a structured NumPy array into a Table.

Extension (does not modify original table)

Table.with_column(label, values)	Return a table with an additional or replaced column.			
Table.with_columns(labels_and_values)	Return a table with additional or replaced columns.			
Table.with_row(row)	Return a table with an additional row.			
Table.with_rows(rows)	Return a table with additional rows.			
Table.relabeled(label, new_label)	Returns a table with label changed to new_label.			

2.1.6 datascience.tables.Table.with_column

Table.with column(label, values)

Return a table with an additional or replaced column.

Args:

label (str): The column label. If an existing label is used, that column will be replaced in the returned table.

values (single value or sequence): If a single value, every value in the new column is values.

If a sequence, the new column contains the values in values. values must be the same length as the table.

Raises:

ValueError: If

- label is not a valid column name
- values is a list/array and does not have the same length as the number of rows in the table.

2.1.7 datascience.tables.Table.with columns

Table.with_columns(labels_and_values)

Return a table with additional or replaced columns.

Args:

labels_and_values: An alternating list of labels and values or a list of label-values pairs.

```
>>> Table().with_columns([
       'letter', ['c', 'd'],
        'count', [2, 4],
...])
letter | count
  | 2
      | 4
>>> Table().with_columns([
... ['letter', ['c', 'd']],
       ['count', [2, 4]],
. . .
...])
letter | count
    | 2
      1 4
>>> Table().with_columns({'letter': ['c', 'd']})
letter
С
d
```

2.1.8 datascience.tables.Table.with row

Table.with row(row)

Return a table with an additional row.

Args: row (sequence): A value for each column.

Raises: ValueError: If the row length differs from the column count.

2.1.9 datascience.tables.Table.with_rows

Table.with rows (rows)

Return a table with additional rows.

Args: rows (sequence of sequences): Each row has a value per column.

If rows is a 2-d array, its shape must be (_, n) for n columns.

Raises: ValueError: If a row length differs from the column count.

2.1.10 datascience.tables.Table.relabeled

Table.relabeled(label, new_label)

Returns a table with label changed to new label.

label and new_label may be single values or lists specifying column labels to be changed and their new corresponding labels.

Args:

label (str or sequence of str): The label(s) of columns to be changed.

new_label (str or sequence of str): The new label(s) of columns to be changed. Same number of elements as label.

Accessing values

Table.num_columns Number of columns.

Table 2.3 – continued from previous page

Table.columns	
Table.column(index_or_label)	Return the values of a column as an array.
Table.num_rows	Number of rows.
Table.rows	Return a view of all rows.
Table.row(index)	Return a row.
Table.labels	Return a tuple of column labels.
Table.column_index(column_label)	Return the index of a column.
Table.apply(fn, column_label)	Returns an array where fn is applied to each set of elements by row from the specified c

2.1.11 datascience.tables.Table.num_columns

Table.num_columns

Number of columns.

2.1.12 datascience.tables.Table.columns

Table.columns

2.1.13 datascience.tables.Table.column

Table.column(index_or_label)

Return the values of a column as an array.

table.column(label) is equivalent to table[label].

Args: label (int or str): The index or label of a column

Returns: An instance of numpy.array.

2.1.14 datascience.tables.Table.num_rows

Table.num_rows

Number of rows.

2.1.15 datascience.tables.Table.rows

 ${\tt Table.rows}$

Return a view of all rows.

2.1.16 datascience.tables.Table.row

```
Table.row(index)
Return a row.
```

2.1.17 datascience.tables.Table.labels

Table.labels

Return a tuple of column labels.

2.1.18 datascience.tables.Table.column_index

```
Table.column_index (column_label)

Return the index of a column.
```

2.1.19 datascience.tables.Table.apply

```
Table.apply (fn, column_label)
```

Returns an array where fn is applied to each set of elements by row from the specified columns in column label.

Args:

fn (function): The function to be applied to elements specified by column_label.

column_label (single string or list of strings): Names of columns to be passed into function fn. Length must match number of elements fn takes.

Raises:

ValueError: column name in column_label is not an existing column in the table.

Returns: A numpy array consisting of results of applying fn to elements specified by column_label in each row.

```
>>> t = Table().with_columns([
       'letter', ['a', 'b', 'c', 'z'],
        'count', [9, 3, 3, 1],
        'points', [1, 2, 2, 10]])
>>> t
letter | count | points
    | 9
           | 1
       1 3
              | 2
       1 3
       | 1
              | 10
>>> t.apply(lambda x, y: x * y, ['count', 'points'])
array([ 9, 6, 6, 10])
>>> t.apply(lambda x: x - 1, 'points')
array([0, 1, 1, 9])
```

Mutation (modifies table in place)

	<pre>Table.set_format(column_label_or_labels,)</pre>	Set the format of a column.	
_	Table.move_to_start(column_label)	Move a column to the first in order.	
Ī	Table.move_to_end(column_label)	Move a column to the last in order.	
			Continued on ne

Table 2.4 – continued from previous page

Ī	Table.append(row_or_table)	Append a row or all rows of a table.				
Table.append_column(label, values)		Appends a column to the table or replaces a column.				
Ī	Table.relabel(column_label, new_label)	Change the labels of columns specified by column_label to labels in new_				

2.1.20 datascience.tables.Table.set_format

Table.set_format (column_label_or_labels, formatter)
Set the format of a column.

2.1.21 datascience.tables.Table.move_to_start

Table.move_to_start (column_label)

Move a column to the first in order.

2.1.22 datascience.tables.Table.move_to_end

Table.move_to_end(column_label)

Move a column to the last in order.

2.1.23 datascience.tables.Table.append

Table.append(row_or_table)

Append a row or all rows of a table. An appended table must have all columns of self.

2.1.24 datascience.tables.Table.append_column

Table.append_column (label, values)

Appends a column to the table or replaces a column.

```
__setitem__ is aliased to this method: table.append_column('new_col', [1, 2, 3]) is equivalent to table['new_col'] = [1, 2, 3].
```

Args: label (str): The label of the new column.

values (single value or list/array): If a single value, every value in the new column is values.

If a list or array, the new column contains the values in values, which must be the same length as the table.

Returns: Original table with new or replaced column

Raises:

ValueError: If

- label is not a string.
- values is a list/array and does not have the same length as the number of rows in the table.

```
>>> table = Table().with_columns([
... 'letter', ['a', 'b', 'c', 'z'],
... 'count', [9, 3, 3, 1],
... 'points', [1, 2, 2, 10]])
```

```
>>> table
letter | count | points
          | 1
    | 9
             | 2
      | 3
      | 3
             | 2
      | 1
             | 10
>>> table.append_column('new_col1', [10, 20, 30, 40])
>>> table
letter | count | points | new_col1
a | 9 | 1 | 10
     | 3
            | 2
                    | 20
     | 3
            | 2
                    | 30
     | 1 | 10 | 40
>>> table.append_column('new_col2', 'hello')
>>> table
letter | count | points | new_col1 | new_col2
a | 9 | 1 | 10
                           | hello
      1 3
             | 2
                     | 20
                                | hello
                  | 30
      | 3
                               | hello
             | 2
   | 1
           | 10
                               | hello
>>> table.append_column(123, [1, 2, 3, 4])
Traceback (most recent call last):
ValueError: The column label must be a string, but a int was given
>>> table.append_column('bad_col', [1, 2])
Traceback (most recent call last):
ValueError: Column length mismatch. New column does not have the same number of rows as table.
```

2.1.25 datascience.tables.Table.relabel

Table.relabel(column_label, new_label)

Change the labels of columns specified by column label to labels in new label.

Args:

column_label (single str or list/array of str): The label(s) of columns to be changed. Must be str.
new_label (single str or list/array of str): The new label(s) of columns to be changed. Must be str.

Number of elements must match number of elements in column_label.

Returns: Original table with modified labels

```
>>> table = Table().with_columns([
       'points', (1, 2, 3),
        'id',
                (12345, 123, 5123)])
>>> table.relabel('id', 'yolo')
points | yolo
1 | 12345
     | 123
     | 5123
>>> table.relabel(['points', 'yolo'], ['red', 'blue'])
red | blue
    | 12345
1
    | 123
    | 5123
>>> table.relabel(['red', 'green', 'blue'], ['cyan', 'magenta', 'yellow', 'key'])
Traceback (most recent call last):
```

Transformation (creates a new table)

Table.copy(*[, shallow])	Return a copy of a Table.
Table.select(column_label_or_labels)	Return a Table with selected column or columns by label or index.
Table.drop(column_label_or_labels)	Return a Table with only columns other than selected label or labels.
Table.take()	Return a new Table of a sequence of rows taken by number.
Table.exclude()	Return a new Table without a sequence of rows excluded by number.
Table.where(column_or_label[, value])	Return a Table of rows for which the column is value or a non-zero value.
Table.sort(column_or_label[, descending,])	Return a Table of rows sorted according to the values in a column.
Table.group(column_or_label[, collect])	Group rows by unique values in a column; count or aggregate others.
Table.groups(labels[, collect])	Group rows by multiple columns, count or aggregate others.
Table.pivot(columns, rows[, values,])	Generate a table with a column for rows (or a column for each row in rows list)
Table.stack(key[, labels])	Takes k original columns and returns two columns, with col.
Table.join(column_label, other[, other_label])	Generate a table with the columns of self and other, containing rows for all valu
Table.stats([ops])	Compute statistics for each column and place them in a table.
Table.percentile(p)	Returns a new table with one row containing the pth percentile for each column
Table.sample([k, with_replacement, weights])	Returns a new table where k rows are randomly sampled from the original table
Table.split(k)	Returns a tuple of two tables where the first table contains k rows randomly san
Table.bin([select])	Group values by bin and compute counts per bin by column.

2.1.26 datascience.tables.Table.copy

Table.copy (*, shallow=False)
Return a copy of a Table.

2.1.27 datascience.tables.Table.select

Table.select (column_label_or_labels)

Return a Table with selected column or columns by label or index.

Args: column_label_or_labels (string or list of strings): The header names or indices of the columns to be selected. column_label_or_labels must be an existing header name, or a valid column index.

Returns: An instance of Table containing only selected columns.

```
cheeseburger | 743
hamburger | 651
veggie burger | 582
>>> t.select('prices')
prices
5
5
>>> t.select(1)
prices
>>> t.select([2, 0])
calories | burgers
743 | cheeseburger
651
        | hamburger
582
       | veggie burger
```

2.1.28 datascience.tables.Table.drop

Table.drop (column_label_or_labels)

Return a Table with only columns other than selected label or labels.

Args: column_label_or_labels (string or list of strings): The header names or indices of the columns to be dropped. column_label_or_labels must be an existing header name, or a valid column index.

Returns: An instance of Table with given columns removed.

```
>>> t = Table().with_columns([
       'burgers', ['cheeseburger', 'hamburger', 'veggie burger'],
       'prices', [6, 5, 5],
       'calories', [743, 651, 582]])
. . .
>>> t
burgers | prices | calories
cheeseburger | 6 | 743
hamburger | 5
veggie burger | 5
                    | 651
                    | 582
>>> t.drop('prices')
burgers | calories
cheeseburger | 743
hamburger | 651
veggie burger | 582
>>> t.drop(['burgers', 'calories'])
prices
6
5
>>> t.drop([0, 2])
prices
5
>>> t.drop(1)
burgers
        | calories
cheeseburger | 743
hamburger
            | 651
```

```
veggie burger | 582
```

2.1.29 datascience.tables.Table.take

Table.take()

Return a new Table of a sequence of rows taken by number.

Args: row_indices_or_slice (integer or list of integers or slice): The row index, list of row indices or a slice of row indices to be selected.

Returns: A new instance of Table.

```
>>> t = Table().with_columns([
       'letter grade', ['A+', 'A', 'A-', 'B+', 'B', 'B-'],
       'gpa', [4, 4, 3.7, 3.3, 3, 2.7]])
. . .
>>> t
letter grade | gpa
A+ | 4
Α
         | 3.7
A -
R+
          | 3.3
В
          | 3
   2.7
>>> t.take(0)
letter grade | gpa
A+ | 4
>>> t.take(5)
letter grade | gpa
>>> t.take(-1)
letter grade | gpa
      | 2.7
>>> t.take([2, 1, 0])
letter grade | gpa
A- | 3.7
          | 4
Α
>>> t.take([1, 5])
letter grade | gpa
      | 4
| 2.7
A
B-
>>> t.take(range(3))
letter grade | gpa
A+ | 4
Α
           | 4
     | 3.7
```

Note that take also supports NumPy-like indexing and slicing:

```
>>> t.take[2, 1, 0]
letter grade | gpa
```

```
A- | 3.7
A | 4
A+ | 4
```

2.1.30 datascience.tables.Table.exclude

Table.exclude()

Return a new Table without a sequence of rows excluded by number.

Args:

row_indices_or_slice (integer or list of integers or slice): The row index, list of row indices or a slice of row indices to be excluded.

Returns: A new instance of Table.

```
>>> t = Table().with_columns([
      'letter grade', ['A+', 'A', 'A-', 'B+', 'B', 'B-'],
      'gpa', [4, 4, 3.7, 3.3, 3, 2.7]])
>>> t
letter grade | gpa
A+ | 4
A
         | 4
A-
         | 3.7
          | 3.3
B+
В
          | 3
     | 2.7
>>> t.exclude(4)
letter grade | gpa
A+ | 4
         | 4
         | 3.7
A-
  | 3.3
B+
>>> t.exclude(-1)
letter grade | gpa
A+ | 4
          | 4
Α
         | 3.7
A-
B+
          | 3.3
>>> t.exclude([1, 3, 4])
letter grade | gpa
A+ | 4
         | 3.7
A-
       2.7
>>> t.exclude(range(3))
letter grade | gpa
B+ | 3.3
В
          | 3
         | 2.7
```

Note that exclude also supports NumPy-like indexing and slicing:

```
>>> t.exclude[:3]
letter grade | gpa
B+ | 3.3
```

```
B | 3
B- | 2.7
```

2.1.31 datascience.tables.Table.where

Table.where (column_or_label, value=None)

Return a Table of rows for which the column is value or a non-zero value.

If column_or_label contains Boolean values, returns rows corresponding to True.

Args: column_or_label: The header name of a column in the table or an array.

value: Value for comparison with items in column_or_label.

Returns: An instance of Table containing rows for which the column_or_label column or column_or_label itself is non-zero or True, or is equal to value, if provided.

```
>>> marbles = Table().with_columns([
      "Color", ["Red", "Green", "Blue", "Red", "Green", "Green"],
      "Shape", ["Round", "Rectangular", "Rectangular", "Round", "Rectangular", "Round"],
. . .
      "Amount", [4, 6, 12, 7, 9, 2],
      "Price", [1.30, 1.20, 2.00, 1.75, 1.40, 1.00]])
>>> marbles
Color | Shape
                  | Amount | Price
Red | Round | 4
                       | 1.3
Green | Rectangular | 6
                           | 1.2
                          | 2
Blue | Rectangular | 12
                          | 1.75
Red | Round | 7
                          | 1.4
Green | Rectangular | 9
Green | Round
             | 2
                          | 1
>>> marbles.where("Shape", "Round")
Color | Shape | Amount | Price
Red | Round | 4 | 1.3
Red | Round | 7
                    | 1.75
Green | Round | 2
                    | 1
>>> marbles.where(marbles.column("Shape") == "Round") # equivalent to the previous example
Color | Shape | Amount | Price
Red
    | Round | 4
                     | 1.3
    | Round | 7
                     | 1.75
Red
Green | Round | 2
                     | 1
>>> marbles.where(marbles.column("Price") > 1.5)
Color | Shape | Amount | Price
                        | 2
Blue | Rectangular | 12
    | Round
              | 7
Red
                           1 1.75
```

2.1.32 datascience.tables.Table.sort

Table.**sort** (*column_or_label*, *descending=False*, *distinct=False*)

Return a Table of rows sorted according to the values in a column.

Args: column_or_label: the column whose values are used for sorting.

descending: if True, sorting will be in descending, rather than ascending order.

distinct: if True, repeated values in column_or_label will be omitted.

Returns: An instance of Table containing rows sorted based on the values in column_or_label.

```
>>> marbles = Table().with columns([
... "Color", ["Red", "Green", "Blue", "Red", "Green", "Green"],
       "Shape", ["Round", "Rectangular", "Rectangular", "Round", "Rectangular", "Round"],
       "Amount", [4, 6, 12, 7, 9, 2],
       "Price", [1.30, 1.30, 2.00, 1.75, 1.40, 1.00]])
. . .
>>> marbles
Color | Shape | Amount | Price Red | Round | 4 | 1.3
Green | Rectangular | 6
                          | 2
Blue | Rectangular | 12
Red | Round | 7
                          | 1.75
Green | Rectangular | 9
Green | Round | 2
                          | 1
>>> marbles.sort("Amount")
Color | Shape | Amount | Price
Green | Round | 2
Red | Round | 4
                  | 2 | 1
                           | 1.3
>>> marbles.sort("Amount", descending = True)
Color | Shape | Amount | Price
Blue | Rectangular | 12 | 2
Green | Rectangular | 9
Red | Round | 7
                          | 1.75
                         | 1.3
Green | Rectangular | 6
Red | Round | 4
                          | 1.3
Green | Round | 2 | 1
>>> marbles.sort(3) # the Price column
Color | Shape | Amount | Price
Green | Round
                   | 2 | 1
Red | Round | 4
                           | 1.3
                          | 1.3
Green | Rectangular | 6
                          | 1.4
Green | Rectangular | 9

      kea | Round | 7
      | 1.75

      Blue | Rectangular | 12
      | 2

>>> marbles.sort(3, distinct = True)
Color | Shape | Amount | Price
Green | Round
                  | 2 | 1
Red | Round | 4
                          | 1.3
Green | Rectangular | 9 | 1.4
Red | Round | 7 | 1.7
                           | 1.75
Blue | Rectangular | 12
                           | 2
```

2.1.33 datascience.tables.Table.group

Table.group (column_or_label, collect=None)

Group rows by unique values in a column; count or aggregate others.

Args: column_or_label: values to group (column label or index, or array)

collect: a function applied to values in other columns for each group

Returns: A Table with each row corresponding to a unique value in <code>column_or_label</code>, where the first column contains the unique values from <code>column_or_label</code>, and the second contains counts for each of the unique values. If <code>collect</code> is provided, a Table is returned with all original columns, each containing values calculated by first grouping rows according to <code>column_or_label</code>, then applying <code>collect</code> to each set of grouped values in the other columns.

Note: The grouped column will appear first in the result table. If collect does not accept arguments with one of the column types, that column will be empty in the resulting table.

```
>>> marbles = Table().with_columns([
      "Color", ["Red", "Green", "Blue", "Red", "Green", "Green"],
      "Shape", ["Round", "Rectangular", "Rectangular", "Round", "Rectangular", "Round"],
      "Amount", [4, 6, 12, 7, 9, 2],
      "Price", [1.30, 1.30, 2.00, 1.75, 1.40, 1.00]])
. . .
>>> marbles
                  | Amount | Price
Color | Shape
             | Amo
                       | 1.3
Red | Round
Green | Rectangular | 6
                           | 1.3
Blue | Rectangular | 12
                         | 2
Red | Round | 7
                         | 1.75
Green | Rectangular | 9 | 1.4
Green | Round | 2 | 1
>>> marbles.group("Color") # just gives counts
Color I count.
Blue | 1
Green | 3
>>> marbles.group("Color", max) # takes the max of each grouping, in each column
Color | Shape max | Amount max | Price max
Blue | Rectangular | 12
                       | 2
                             | 1.4
Green | Round | 9
Red | Round | 7
                              | 1.75
>>> marbles.group("Shape", sum) # sum doesn't make sense for strings
Shape | Color sum | Amount sum | Price sum
Rectangular | 27 | 4.7
Round |
                      | 13
                                 | 4.05
```

2.1.34 datascience.tables.Table.groups

Table.groups(labels, collect=None)

Group rows by multiple columns, count or aggregate others.

Args: labels: list of column names (or indices) to group on

collect: a function applied to values in other columns for each group

Returns: A Table with each row corresponding to a unique combination of values in the columns specified in labels, where the first columns are those specified in labels, followed by a column of counts for each of the unique values. If collect is provided, a Table is returned with all original columns, each containing values calculated by first grouping rows according to to values in the labels column, then applying collect to each set of grouped values in the other columns.

Note: The grouped columns will appear first in the result table. If collect does not accept arguments with one of the column types, that column will be empty in the resulting table.

```
>>> marbles = Table().with_columns([
      "Color", ["Red", "Green", "Blue", "Red", "Green", "Green"],
      "Shape", ["Round", "Rectangular", "Rectangular", "Round", "Rectangular", "Round"],
      "Amount", [4, 6, 12, 7, 9, 2],
      "Price", [1.30, 1.30, 2.00, 1.75, 1.40, 1.00]])
. . .
>>> marbles
Color | Shape
                  | Amount | Price
                       | 1.3
Red | Round
                  | 4
Green | Rectangular | 6
                          | 1.3
                          | 2
Blue | Rectangular | 12
Red | Round | 7
                          | 1.75
Green | Rectangular | 9
                          | 1.4
Green | Round | 2
                           | 1
>>> marbles.groups(["Color", "Shape"])
Color | Shape
              | count
Blue | Rectangular | 1
Green | Rectangular | 2
Green | Round
Red | Round
                  | 2
>>> marbles.groups(["Color", "Shape"], sum)
Color | Shape | Amount sum | Price sum
Blue | Rectangular | 12
                           | 2
Green | Rectangular | 15
                               | 2.7
Green | Round
                 | 2
                               | 1
Red
    Round
                  | 11
                               | 3.05
```

2.1.35 datascience.tables.Table.pivot

Table.pivot (columns, rows, values=None, collect=None, zero=None)

Generate a table with a column for rows (or a column for each row in rows list) and a column for each unique value in columns. Each row counts/aggregates the values that match both row and column.

columns – column label in self rows – column label or a list of column labels values – column label in self (or None to produce counts) collect – aggregation function over values zero – zero value for non-existent row-column combinations

2.1.36 datascience.tables.Table.stack

Table.stack (key, labels=None)

Takes k original columns and returns two columns, with col. 1 of all column names and col. 2 of all associated data.

2.1.37 datascience.tables.Table.join

Table.join (column_label, other, other_label=None)

Generate a table with the columns of self and other, containing rows for all values of a column that appear in both tables. If a join value appears more than once in self, each row will be used, but in the other table, only the first of each will be used.

If the result is empty, return None.

2.1.38 datascience.tables.Table.stats

Table.stats(ops=(<built-in function min>, <built-in function max>, <function median at 0x7f9aaa0f22f0>, <built-in function sum>))

Compute statistics for each column and place them in a table.

2.1.39 datascience.tables.Table.percentile

Table.percentile(p)

Returns a new table with one row containing the pth percentile for each column.

Assumes that each column only contains one type of value.

Returns a new table with one row and the same column labels. The row contains the pth percentile of the original column, where the pth percentile of a column is the smallest value that at at least as large as the p% of numbers in the column.

2.1.40 datascience.tables.Table.sample

Table.sample(*k*=*None*, *with_replacement*=*False*, *weights*=*None*)

Returns a new table where k rows are randomly sampled from the original table.

Kwargs:

k (int or None): If None (default), all the rows in the table are sampled. If an integer, k rows from the original table are sampled.

with_replacement (bool): If False (default), samples the rows without replacement. If True, samples the rows with replacement.

weights (list/array or None): If None (default), samples the rows using a uniform random distribution. If a list/array is passed in, it must be the same length as the number of rows in the table and the values must sum to 1. The rows will then be sampled according to the probability distribution in weights.

Returns: A new instance of Table.

```
>>> jobs.sample()
job | wage
    | 20
b
    | 15
    | 10
    | 8
>>> jobs.sample(k = 2)
job | wage
  | 20
    | 15
>>> jobs.sample(k = 2, with_replacement = True,
       weights = [0.5, 0.5, 0, 0])
job | wage
    | 10
а
    | 10
а
```

2.1.41 datascience.tables.Table.split

```
Table.split (k)
```

Returns a tuple of two tables where the first table contains k rows randomly sampled and the second contains the remaining rows.

Args:

k (int): The number of rows randomly sampled into the first table. k must be between 1 and num_rows - 1.

Raises: ValueError: k is not between 1 and num_rows - 1.

Returns: A tuple containing two instances of Table.

```
>>> jobs = Table().with_columns([
        'job', ['a', 'b', 'c', 'd'],
        'wage', [10, 20, 15, 8]])
>>> jobs
job | wage
    | 10
    | 20
b
    | 15
С
d
    | 8
>>> sample, rest = jobs.split(3)
>>> sample
job | wage
    | 15
    | 10
а
b
    | 20
>>> rest
job | wage
     | 8
```

2.1.42 datascience.tables.Table.bin

```
Table.bin (select=None, **vargs)
```

Group values by bin and compute counts per bin by column.

By default, bins are chosen to contain all values in all columns. The following named arguments from numpy.histogram can be applied to specialize bin widths:

If the original table has n columns, the resulting binned table has n+1 columns, where column 0 contains the lower bound of each bin.

Args:

select (columns): Columns to be binned. If None, all columns are binned.

bins (int or sequence of scalars): If bins is an int, it defines the number of equal-width bins in the given range (10, by default). If bins is a sequence, it defines the bin edges, including the rightmost edge, allowing for non-uniform bin widths.

range ((float, float)): The lower and upper range of the bins. If not provided, range contains all values in the table. Values outside the range are ignored.

density (bool): If False, the result will contain the number of samples in each bin. If True, the result is the value of the probability density function at the bin, normalized such that the integral over the range is 1. Note that the sum of the histogram values will not be equal to 1 unless bins of unity width are chosen; it is not a probability mass function.

Exporting / Displaying

Table.show([max_rows])	Display the table.
<pre>Table.as_text([max_rows, sep])</pre>	Format table as text.
Table.as_html([max_rows])	Format table as HTML.
Table.index_by(column_or_label)	Return a dict keyed by values in a column that contains lists of rows corresponding to each
Table.to_array()	Convert the table to a NumPy array.
Table.to_df()	Convert the table to a Pandas DataFrame.
Table.to_csv(filename)	Creates a CSV file with the provided filename.

2.1.43 datascience.tables.Table.show

Table.**show** (*max_rows=0*) Display the table.

2.1.44 datascience.tables.Table.as text

Table.as_text (max_rows=0, sep='|')
Format table as text.

2.1.45 datascience.tables.Table.as html

Table.as_html (max_rows=0)
Format table as HTML.

2.1.46 datascience.tables.Table.index_by

Table.index_by (column_or_label)

Return a dict keyed by values in a column that contains lists of rows corresponding to each value.

2.1.47 datascience.tables.Table.to array

```
Table.to_array()
Convert the table to a NumPy array.
```

2.1.48 datascience.tables.Table.to_df

```
Table.to_df()
```

Convert the table to a Pandas DataFrame.

2.1.49 datascience.tables.Table.to_csv

```
Table.to_csv(filename)
```

Creates a CSV file with the provided filename.

The CSV is created in such a way that if we run table.to_csv('my_table.csv') we can recreate the same table with Table.read_table('my_table.csv').

Args: filename (str): The filename of the output CSV file.

Returns: None, outputs a file with name filename.

Visualizations

<pre>Table.plot([column_for_xticks, select, overlay])</pre>	Plot line charts for the table.
Table.bar([column_for_categories, select,])	Plot bar charts for the table.
Table.barh([column_for_categories, select,])	Plot horizontal bar charts for the table.
Table.pivot_hist(pivot_column_label,[,])	Draw histograms of each category in a column.
<pre>Table.hist([select, overlay, bins, counts, unit])</pre>	Plots one histogram for each column in the table.
Table.points(columnlat, columnlong[,])	Draw points from latitude and longitude columns.
Table.scatter(column_for_x[, select,])	Creates scatterplots, optionally adding a line of best fit.
Table.boxplot(**vargs)	Plots a boxplot for the table.

2.1.50 datascience.tables.Table.plot

```
Table.plot (column_for_xticks=None, select=None, overlay=True, **vargs)

Plot line charts for the table.
```

Each plot is labeled using the values in *column_for_xticks* and one plot is produced for every other column (or for the columns designated by *select*).

Every selected column except for *column_for_xticks* must be numerical.

Args: column_for_xticks (str/array): A column containing x-axis labels

Kwargs:

overlay (bool): create a chart with one color per data column; if False, each will be displayed separately.

vargs: Additional arguments that get passed into *plt.plot*. See http://matplotlib.org/api/pyplot_api.html#matplotlib.pyplot for additional arguments that can be passed into vargs.

2.1.51 datascience.tables.Table.bar

Table.bar (column_for_categories=None, select=None, overlay=True, **vargs)
Plot bar charts for the table.

Each plot is labeled using the values in *column_for_categories* and one plot is produced for every other column (or for the columns designated by *select*).

Every selected except column for *column_for_categories* must be numerical.

Args: column_for_categories (str): A column containing x-axis categories

Kwargs:

overlay (bool): create a chart with one color per data column; if False, each will be displayed separately.

vargs: Additional arguments that get passed into *plt.bar*. See http://matplotlib.org/api/pyplot_api.html#matplotlib.pyplo for additional arguments that can be passed into vargs.

2.1.52 datascience.tables.Table.barh

Table .barh (column_for_categories=None, select=None, overlay=True, **vargs)
Plot horizontal bar charts for the table.

Each plot is labeled using the values in *column_for_categories* and one plot is produced for every other column (or for the columns designated by *select*).

Every selected except column for *column_for_categories* must be numerical.

Args: column_for_categories (str): A column containing y-axis categories

Kwargs:

overlay (bool): create a chart with one color per data column; if False, each will be displayed separately.

vargs: Additional arguments that get passed into *plt.barh*. See http://matplotlib.org/api/pyplot_api.html#matplotlib.pypl for additional arguments that can be passed into vargs.

2.1.53 datascience.tables.Table.pivot_hist

Table.pivot_hist (pivot_column_label, value_column_label, overlay=True, **vargs)

Draw histograms of each category in a column.

2.1.54 datascience.tables.Table.hist

Table.hist (select=None, overlay=True, bins=None, counts=None, unit=None, **vargs)
Plots one histogram for each column in the table.

Every column must be numerical.

Kwargs:

overlay (bool): If True, plots 1 chart with all the histograms overlaid on top of each other (instead of the default behavior of one histogram for each column in the table). Also adds a legend that matches each bar color to its column.

bins (column name or list): Lower bound for each bin in the histogram. If None, bins will be chosen automatically.

counts (column name or column): A column of counted values. All other columns are treated as counts of these values. If None, each value in each row is assigned a count of 1.

vargs: Additional arguments that get passed into:func:plt.hist. See http://matplotlib.org/api/pyplot_api.html#matplotlib for additional arguments that can be passed into vargs. These include: range, normed, cumulative, and orientation, to name a few.

```
>>> t = Table().with_columns([
        'count', [9, 3, 3, 1],
. . .
        'points', [1, 2, 2, 10]])
. . .
count | points
      1 1
3
      1 2
3
      | 2
1
      | 10
>>> t.hist()
<histogram of values in count>
<histogram of values in points>
```

2.1.55 datascience.tables.Table.points

Table.points (column_lat, column_long, labels=None, colors=None, **kwargs)

Draw points from latitude and longitude columns. [Deprecated]

2.1.56 datascience.tables.Table.scatter

Table.scatter (column_for_x, select=None, overlay=True, fit_line=False, **vargs)
Creates scatterplots, optionally adding a line of best fit.

Each plot uses the values in *column_for_x* for horizontal positions. One plot is produced for every other column as y (or for the columns designated by *select*).

Every selected except column for *column_for_categories* must be numerical.

Args:

column_for_x (str): The name to use for the x-axis values of the scatter plots.

Kwargs:

overlay (bool): create a chart with one color per data column; if False, each will be displayed separately.

fit_line (bool): draw a line of best fit for each set of points

vargs: Additional arguments that get passed into *plt.scatter*. See http://matplotlib.org/api/pyplot_api.html#matplotlib.pr for additional arguments that can be passed into vargs. These include: *marker* and *norm*, to name a couple.

```
>>> table = Table().with_columns([
       'x', [9, 3, 3, 1],
        'y', [1, 2, 2, 10],
        'z', [3, 4, 5, 6]])
. . .
>>> table
            Z
    ΙУ
9
    | 1
           | 3
3
    | 2
           | 4
3
    | 2
           | 5
    | 10
          | 6
>>> table.scatter('x')
<scatterplot of values in y and z on x>
```

```
>>> table.scatter('x', overlay=False)
<scatterplot of values in y on x>
<scatterplot of values in z on x>
```

```
>>> table.scatter('x', fit_line=True)
<scatterplot of values in y and z on x with lines of best fit>
```

2.1.57 datascience.tables.Table.boxplot

Table.boxplot(**vargs)

Plots a boxplot for the table.

Every column must be numerical.

Kwargs:

vargs: Additional arguments that get passed into *plt.boxplot*. See http://matplotlib.org/api/pyplot_api.html#matplotlib.py for additional arguments that can be passed into vargs. These include *vert* and *showmeans*.

Returns: None

Raises: ValueError: The Table contains columns with non-numerical values.

```
>>> table = Table().with_columns([
        'test1', [92.5, 88, 72, 71, 99, 100, 95, 83, 94, 93],
        'test2', [89, 84, 74, 66, 92, 99, 88, 81, 95, 94]])
>>> table
test1 | test2
92.5 | 89
88
      | 84
        74
71
      | 66
99
      1 92
100
      1 99
95
      | 88
8.3
      | 81
94
      | 95
93
      | 94
>>> table.boxplot()
<boxplot of test1 and boxplot of test2 side-by-side on the same figure>
```

2.2 Maps (datascience.maps)

Draw maps using folium.

```
class datascience.maps.Map (features=(), ids=(), width=960, height=500, **kwargs)
A map from IDs to features. Keyword args are forwarded to folium.
```

```
color (values, ids=(), key_on='feature.id', palette='YlOrBr', **kwargs)
Color map features by binning values.
```

values – a sequence of values or a table of keys and values ids – an ID for each value; if none are provided, indices are used key_on – attribute of each feature to match to ids palette – one of the following color brewer palettes:

```
'BuGn', 'BuPu', 'GnBu', 'OrRd', 'PuBu', 'PuBuGn', 'PuRd', 'RdPu', 'YlGn', 'YlGnBu', 'YlOrBr', and 'YlOrRd'.
```

Defaults from Folium:

threshold_scale: list, default None Data range for D3 threshold scale. Defaults to the following range of quantiles: [0, 0.5, 0.75, 0.85, 0.9], rounded to the nearest order-of-magnitude integer. Ex: 270 rounds to 200, 5600 to 6000.

fill opacity: float, default 0.6 Area fill opacity, range 0-1.

line_color: string, default 'black' GeoJSON geopath line color.

line_weight: int, default 1 GeoJSON geopath line weight.

line_opacity: float, default 1 GeoJSON geopath line opacity, range 0-1.

legend_name: string, default None Title for data legend. If not passed, defaults to columns[1].

features

```
format (**kwargs)
           Apply formatting.
     geojson()
          Render features as a FeatureCollection.
     classmethod read geojson (path or json or string)
           Read a geoJSON string, object, or file. Return a dict of features keyed by ID.
class datascience.maps.Marker(lat, lon, popup='', color='blue', **kwargs)
     A marker displayed with Folium's simple_marker method.
     popup – text that pops up when marker is clicked color – fill color
     Defaults from Folium:
     marker_icon: string, default 'info-sign' icon from (http://getbootstrap.com/components/) you want on the
     clustered_marker: boolean, default False boolean of whether or not you want the marker clustered with other
          markers
     icon angle: int, default 0 angle of icon
     popup_width: int, default 300 width of popup
     copy()
          Return a deep copy
     format (**kwargs)
           Apply formatting.
     geojson (feature_id)
           GeoJSON representation of the marker as a point.
     lat lons
     classmethod map (latitudes, longitudes, labels=None, colors=None, **kwargs)
           Return markers from columns of coordinates, labels, & colors.
     classmethod map_table (table, **kwargs)
           Return markers from the colums of a table.
class datascience.maps.Circle (lat, lon, popup='', color='blue', radius=10, **kwargs)
     A marker displayed with Folium's circle marker method.
     popup – text that pops up when marker is clicked color – fill color radius – pixel radius of the circle
     Defaults from Folium:
     fill opacity: float, default 0.6 Circle fill opacity
class datascience.maps.Region (geojson, **kwargs)
     A GeoJSON feature displayed with Folium's geo_json method.
     copy()
          Return a deep copy
     format (**kwargs)
           Apply formatting.
     geojson (feature_id)
          Return GeoJSON with ID substituted.
     lat lons
           A flat list of (lat, lon) pairs.
```

polygons

Return a list of polygons describing the region.

- •Each polygon is a list of linear rings, where the first describes the exterior and the rest describe interior holes.
- •Each linear ring is a list of positions where the last is a repeat of the first.
- •Each position is a (lat, lon) pair.

properties

type

The GEOJSON type of the regions: Polygon or MultiPolygon.

2.3 Formats (datascience.formats)

```
String formatting for table entries.
```

```
class datascience.formats.Formatter(min_width=None, max_width=None, etc=None)
     String formatter that truncates long values.
     static convert (value)
          Identity conversion (override to convert values).
     converts_values
          Whether this Formatter also converts values.
     etc = ' ...'
     format_column (label, column)
          Return a formatting function that pads & truncates values.
     static format_value(value)
          Pretty-print an arbitrary value.
     max width = 60
     min width = 4
class datascience.formats.NumberFormatter (decimals=2, decimal_point='.', separator=', ')
     Format numbers that may have delimiters.
     convert (value)
          Convert string 93,000.00 to float 93000.0.
     converts_values = True
     format\_value(value)
class datascience.formats.CurrencyFormatter(symbol='$', *args, **vargs)
     Format currency and convert to float.
     convert (value)
          Convert value to float. If value is a string, ensure that the first character is the same as symbol ie. the value
          is in the currency this formatter is representing.
     converts_values = True
     format_value(value)
```

Format currency.

```
class datascience.formats.DateFormatter(format='%Y-%m-%d
                                                                        %H:%M:%S.%f',
                                                                                             *args,
                                                  **vargs)
     Format date & time and convert to UNIX timestamp.
     convert (value)
          Convert 2015-08-03 to a Unix timestamp int.
     converts_values = True
     format value(value)
          Format timestamp as a string.
class datascience.formats.PercentFormatter(decimals=2, *args, **vargs)
     Format a number as a percentage.
     converts_values = False
     format_value(value)
          Format number as percentage.
2.4 Utility Functions (datascience.util)
Utility functions
datascience.util.percentile(p, arr=None)
     Returns the pth percentile of the input array (the value that is at least as great as p\% of the values in the array)
     If arr is not provided, percentile returns itself curried with p
     >>> percentile(67, [1, 3, 5, 9])
     >>> percentile(66, [1, 3, 5, 9])
     >>> f = percentile(66)
     >>> f([1, 3, 5, 9])
datascience.util.plot_cdf_area(rbound=None, lbound=None, mean=0, sd=1)
     Plots a normal curve with specified parameters and area below curve shaded between lbound and rbound.
     Args: rbound (numeric): right boundary of shaded region
          1bound (numeric): left boundary of shaded region; by default is negative infinity
          mean (numeric): mean/expectation of normal distribution
          sd (numeric): standard deviation of normal distribution
datascience.util.plot_normal_cdf(rbound=None, lbound=None, mean=0, sd=1)
     Plots a normal curve with specified parameters and area below curve shaded between lbound and rbound.
     Args: rbound (numeric): right boundary of shaded region
          lbound (numeric): left boundary of shaded region; by default is negative infinity
          mean (numeric): mean/expectation of normal distribution
          sd (numeric): standard deviation of normal distribution
datascience.util.table_apply (table, func, subset=None)
     Applies a function to each column and returns a Table.
```

Uses pandas apply under the hood, then converts back to a Table

table [instance of Table] The table to apply your function to

func [function] Any function that will work with DataFrame.apply

subset [list | None] A list of columns to apply the function to. If None, function will be applied to all columns in table

tab [instance of Table] A table with the given function applied. It will either be the shape == shape(table), or shape (1, table.shape[1])

datascience.util.minimize(f, start=None, **vargs)

Minimize a function f of one or more arguments.

Returns either:

- 1. the minimizing argument of a one-argument function
- 2. an array of minimizing arguments of a multi-argument function

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