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The datascience package was written for use in Berkeley’s DS 8 course and contains useful functionality for investigating and graphically displaying data.
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 the `Table` class, which is the structure used to represent columns of data. First, load the class:

```python
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:

```python
# HIDDEN
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, 1],
    ...:     'points', [1, 2, 2, 10],
    ...: )

In [3]: print(t)
letter | count | points
a | 9 | 1
b | 3 | 2
c | 3 | 2
z | 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()`:

```
In [5]: Table.read_table('sample.csv')
Out[5]:
x | y | z
1 | 10 | 100
2 | 11 | 101
3 | 12 | 102
```

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

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

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

```python
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)
letter | points | count
a      | 1      | 9
b      | 2      | 3
c      | 2      | 3
z      | 10     | 1
```

### 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).

```python
In [9]: t
Out[9]:
letter | points | count
a      | 1      | 9
b      | 2      | 3
c      | 2      | 3
z      | 10     | 1

In [10]: t.column('letter')
array(['a', 'b', 'c', 'z'],
      dtype='<U1')

In [11]: t.column(1)
array([ 1, 2, 2, 10])
```

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

```python
In [12]: t['letter'] # This is a shorthand for t.column('letter')
Out[12]:
array(['a', 'b', 'c', 'z'],
      dtype='<U1')

In [13]: t[1]  # This is a shorthand for t.column(1)
Out[13]: array([ 1, 2, 2, 10])
```

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(letter | points | count
a   | 1   | 9
b   | 2   | 3
c   | 2   | 3
z   | 10  | 1)

In [15]: t.rows[0]
\nIn [16]: t.row(0)
\nIn [17]: second = t.rows[1]

In [18]: second
Out[18]: Row(letter='b', points=2, count=3)

In [19]: second[0]
\n\nIn [20]: second[1]
\nTo 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]:
letter | points | count
a   | 1   | 9
b   | 2   | 3
c   | 2   | 3
z   | 10  | 1

In [23]: t.with_column('vowel?', ['yes', 'no', 'no', 'no'])
letter | points | count | vowel?
|-------|-------|-------|-------
| a     | 1     | 9     | yes   |
| b     | 2     | 3     | no    |
| c     | 2     | 3     | no    |
| z     | 10    | 1     | no    |

In [24]: t # .with_column returns a new table without modifying the original
In [25]: t.with_column('2 * count', t['count'] * 2)  # A simple way to operate on columns

letter | points | count | 2 * count
a | 1 | 9 | 18
b | 2 | 3 | 6
c | 2 | 3 | 6
z | 10 | 1 | 2

Selecting columns with `select()`:

In [26]: t.select('letter')
Out[26]:
letter
a
b
c
z

In [27]: t.select(['letter', 'points'])

letter | points
a | 1
b | 2
c | 2
z | 10

Renaming columns with `relabeled()`:

In [28]: t
Out[28]:
letter | points | count
a | 1 | 9
b | 2 | 3
c | 2 | 3
z | 10 | 1

In [29]: t.relabeled('points', 'other name')

letter | other name | count
a | 1 | 9
b | 2 | 3
c | 2 | 3
z | 10 | 1

In [30]: t

letter | points | count
a | 1 | 9
b | 2 | 3
c | 2 | 3
z | 10 | 1

In [31]: t.relabeled(['letter', 'count', 'points'], ['x', 'y', 'z'])

x | z | y
a | 1 | 9
Selecting out rows by index with `take()` and conditionally with `where()`:

In [32]: t
Out[32]:
<table>
<thead>
<tr>
<th>letter</th>
<th>points</th>
<th>count</th>
</tr>
</thead>
<tbody>
<tr>
<td>a</td>
<td>1</td>
<td>9</td>
</tr>
<tr>
<td>b</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>c</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>z</td>
<td>10</td>
<td>1</td>
</tr>
</tbody>
</table>

In [33]: t.take(2) # the third row
Out[33]:
<table>
<thead>
<tr>
<th>letter</th>
<th>points</th>
<th>count</th>
</tr>
</thead>
<tbody>
<tr>
<td>c</td>
<td>2</td>
<td>3</td>
</tr>
</tbody>
</table>

In [34]: t.take[0:2] # the first and second rows
Out[34]:
<table>
<thead>
<tr>
<th>letter</th>
<th>points</th>
<th>count</th>
</tr>
</thead>
<tbody>
<tr>
<td>a</td>
<td>1</td>
<td>9</td>
</tr>
<tr>
<td>b</td>
<td>2</td>
<td>3</td>
</tr>
</tbody>
</table>

In [35]: t.where('points', 2) # rows where points == 2
Out[35]:
<table>
<thead>
<tr>
<th>letter</th>
<th>points</th>
<th>count</th>
</tr>
</thead>
<tbody>
<tr>
<td>b</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>c</td>
<td>2</td>
<td>3</td>
</tr>
</tbody>
</table>

In [36]: t.where(t['count'] < 8) # rows where count < 8
Out[36]:
<table>
<thead>
<tr>
<th>letter</th>
<th>points</th>
<th>count</th>
</tr>
</thead>
<tbody>
<tr>
<td>b</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>c</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>z</td>
<td>10</td>
<td>1</td>
</tr>
</tbody>
</table>

In [37]: t['count'] < 8 # .where actually takes in an array of booleans

In [38]: t.where([False, True, True, True]) # same as the last line
Out[38]:
<table>
<thead>
<tr>
<th>letter</th>
<th>points</th>
<th>count</th>
</tr>
</thead>
<tbody>
<tr>
<td>b</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>c</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>z</td>
<td>10</td>
<td>1</td>
</tr>
</tbody>
</table>

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

In [39]: t
Out[39]:
<table>
<thead>
<tr>
<th>letter</th>
<th>points</th>
<th>count</th>
</tr>
</thead>
<tbody>
<tr>
<td>a</td>
<td>1</td>
<td>9</td>
</tr>
<tr>
<td>b</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>c</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>z</td>
<td>10</td>
<td>1</td>
</tr>
</tbody>
</table>

In [40]: t.sort('count')
# You may pass a reducing function into the collect arg
# Note the renaming of the points column because of the collect arg
In [42]: t.select(['count', 'points']).group('count', collect=sum)
Out[42]:
  count | points sum
     1   |       10
     3   |       4
     9   |

1.5 Visualizing Data

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

In [46]: normal_data = Table().with_columns([....:   'data1', np.random.normal(loc = 1, scale = 2, size = 100),....:   'data2', np.random.normal(loc = 4, scale = 3, size = 100)])....:

In [47]: normal_data
Out[47]:
data1    | data2
---      | ---
0.166218 | -0.574452
2.93638  | 2.2193
-0.870999 | 1.13241
0.11712  | 0.345876
0.928033 | 1.16795
-1.66621 | 7.39356
-0.248231 | -1.86525
3.02329  | -0.527577
2.75636  | 6.53435
1.47817  | 0.311778
... (90 rows omitted)

Draw histograms with `hist()`:

In [48]: normal_data.hist()

_draw/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()`:
In [51]: normal_data.sort('data1').plot('data1')  # Sort first to make plot nicer

_build/latex/_images/plot.png

In [52]: normal_data.scatter('data1')

_build/latex/_images/scatter.png
In [53]: normal_data.scatter('data1', fit_line = True)

Use `barh()` to display categorical data.

In [54]: t
Out[54]:
<table>
<thead>
<tr>
<th>letter</th>
<th>points</th>
<th>count</th>
</tr>
</thead>
<tbody>
<tr>
<td>a</td>
<td>1</td>
<td>9</td>
</tr>
<tr>
<td>b</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>c</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>z</td>
<td>10</td>
<td>1</td>
</tr>
</tbody>
</table>

In [55]: t.barh('letter')
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()`:

```python
In [56]: normal_data
```
```
Out[56]:
data1    data2
0  0.166218 -0.574452
1  2.93638  2.2193
2 -0.870999  1.13241
... (90 rows omitted)
```

# index = False prevents row numbers from appearing in the resulting CSV

```python
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.
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('https://github.com/data-8/textbook/raw/9aa0a167bc514749338cd7754f2b9338cd7754f2b/notebooks/baby.csv')

In [59]: baby
Out[59]:
<table>
<thead>
<tr>
<th>birthwt</th>
<th>gest_days</th>
<th>mat_age</th>
<th>mat_ht</th>
<th>mat_pw</th>
<th>m_smoker</th>
</tr>
</thead>
<tbody>
<tr>
<td>120</td>
<td>284</td>
<td>27</td>
<td>62</td>
<td>100</td>
<td>0</td>
</tr>
<tr>
<td>113</td>
<td>282</td>
<td>33</td>
<td>64</td>
<td>135</td>
<td>0</td>
</tr>
<tr>
<td>128</td>
<td>279</td>
<td>28</td>
<td>64</td>
<td>115</td>
<td>1</td>
</tr>
<tr>
<td>108</td>
<td>282</td>
<td>23</td>
<td>67</td>
<td>125</td>
<td>1</td>
</tr>
<tr>
<td>136</td>
<td>286</td>
<td>25</td>
<td>62</td>
<td>93</td>
<td>0</td>
</tr>
<tr>
<td>138</td>
<td>244</td>
<td>33</td>
<td>62</td>
<td>178</td>
<td>0</td>
</tr>
<tr>
<td>132</td>
<td>245</td>
<td>23</td>
<td>65</td>
<td>140</td>
<td>0</td>
</tr>
<tr>
<td>120</td>
<td>289</td>
<td>25</td>
<td>62</td>
<td>125</td>
<td>0</td>
</tr>
<tr>
<td>143</td>
<td>299</td>
<td>30</td>
<td>66</td>
<td>136</td>
<td>1</td>
</tr>
<tr>
<td>140</td>
<td>351</td>
<td>27</td>
<td>68</td>
<td>120</td>
<td>0</td>
</tr>
</tbody>
</table>
... (1164 rows omitted)
```

Let's take a peek at the table.

```
In [59]: baby # Let's take a peek at the table
Out[59]:
<table>
<thead>
<tr>
<th>birthwt</th>
<th>gest_days</th>
<th>mat_age</th>
<th>mat_ht</th>
<th>mat_pw</th>
<th>m_smoker</th>
</tr>
</thead>
<tbody>
<tr>
<td>120</td>
<td>284</td>
<td>27</td>
<td>62</td>
<td>100</td>
<td>0</td>
</tr>
<tr>
<td>113</td>
<td>282</td>
<td>33</td>
<td>64</td>
<td>135</td>
<td>0</td>
</tr>
<tr>
<td>128</td>
<td>279</td>
<td>28</td>
<td>64</td>
<td>115</td>
<td>1</td>
</tr>
<tr>
<td>108</td>
<td>282</td>
<td>23</td>
<td>67</td>
<td>125</td>
<td>1</td>
</tr>
<tr>
<td>136</td>
<td>286</td>
<td>25</td>
<td>62</td>
<td>93</td>
<td>0</td>
</tr>
<tr>
<td>138</td>
<td>244</td>
<td>33</td>
<td>62</td>
<td>178</td>
<td>0</td>
</tr>
<tr>
<td>132</td>
<td>245</td>
<td>23</td>
<td>65</td>
<td>140</td>
<td>0</td>
</tr>
<tr>
<td>120</td>
<td>289</td>
<td>25</td>
<td>62</td>
<td>125</td>
<td>0</td>
</tr>
<tr>
<td>143</td>
<td>299</td>
<td>30</td>
<td>66</td>
<td>136</td>
<td>1</td>
</tr>
<tr>
<td>140</td>
<td>351</td>
<td>27</td>
<td>68</td>
<td>120</td>
<td>0</td>
</tr>
</tbody>
</table>
... (1164 rows omitted)
```

Let's select out columns we want.

```
In [60]: smoker_and_wt = baby.select(['m_smoker', 'birthwt'])

In [61]: smoker_and_wt
Out[61]:
<table>
<thead>
<tr>
<th>m_smoker</th>
<th>birthwt</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>120</td>
</tr>
<tr>
<td>0</td>
<td>113</td>
</tr>
<tr>
<td>1</td>
<td>128</td>
</tr>
<tr>
<td>1</td>
<td>108</td>
</tr>
<tr>
<td>0</td>
<td>136</td>
</tr>
<tr>
<td>0</td>
<td>138</td>
</tr>
<tr>
<td>0</td>
<td>132</td>
</tr>
<tr>
<td>0</td>
<td>120</td>
</tr>
<tr>
<td>1</td>
<td>143</td>
</tr>
<tr>
<td>0</td>
<td>140</td>
</tr>
</tbody>
</table>
... (1164 rows omitted)
```

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

```
In [62]: smoker_and_wt.select('m_smoker').hist(bins = [0, 1, 2]);
```
We can also compare the distribution of birthweights between smokers and non-smokers.

```python
# Non smokers
# We do this by grabbing the rows that correspond to mothers that don't
# smoke, then plotting a histogram of just the birthweights.
In [63]: smoker_and_wt.where('m_smoker', 0).select('birthwt').hist()

# Smokers
In [64]: smoker_and_wt.where('m_smoker', 1).select('birthwt').hist()
```
What's the difference in mean birth weight of the two categories?

In [65]: nonsmoking_mean = smoker_and_wt.where('m_smoker', 0).column('birthwt').mean()

In [66]: smoking_mean = smoker_and_wt.where('m_smoker', 1).column('birthwt').mean()
Let’s do the bootstrap test on the two categories.

```python
In [69]: num_nonsmokers = smoker_and_wt.where('m_smoker', 0).num_rows

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) ])

In [73]: bootstrapped_diff_means[:10]
Out[73]: array([ 0.64817405, 0.31037068, -1.47215747, -1.31250362, -0.17084876,
                   -1.3440529 , 0.32526471, 1.20186785, 0.38423755, 0.09761872])

In [74]: num_diffs_greater = (abs(bootstrapped_diff_means) > abs(observed_diff)).sum()

In [75]: p_value = num_diffs_greater / len(bootstrapped_diff_means)

In [76]: p_value
Out[76]: 0.0
```

### 1.8 Drawing Maps

To come.
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

```python
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`:

```python
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:

```python
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

<table>
<thead>
<tr>
<th>Method</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><code>Table.__init__([labels, _deprecated, formatter])</code></td>
<td>Create an empty table with column labels.</td>
</tr>
<tr>
<td><code>Table.empty([labels])</code></td>
<td>Creates an empty table.</td>
</tr>
<tr>
<td><code>Table.from_records(records)</code></td>
<td>Create a table from a sequence of records (dicts with fixed keys).</td>
</tr>
<tr>
<td><code>Table.from_columns_dict(columns)</code></td>
<td>Create a table from a mapping of column labels to column values.</td>
</tr>
<tr>
<td><code>Table.read_table(filepath_or_buffer, *args, ...)</code></td>
<td>Read a table from a file or web address.</td>
</tr>
<tr>
<td><code>Table.from_df(df)</code></td>
<td>Convert a Pandas DataFrame into a Table.</td>
</tr>
<tr>
<td><code>Table.from_array(arr)</code></td>
<td>Convert a structured NumPy array into a Table.</td>
</tr>
</tbody>
</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.

```python
tiles = Table(make_array('letter', 'count', 'points'))
tiles
```

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.empty

classmethod Table.empty(labels=None)

Creates an empty table. Column labels are optional. [Deprecated]

Args:

labels (None or list): If None, a table with 0 columns is created. If a list, each element is a column label in a table with 0 rows.

Returns: A new instance of Table.

2.1.3 datascience.tables.Table.from_records

classmethod Table.from_records(records)

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

2.1.4 datascience.tables.Table.from_columns_dict

classmethod Table.from_columns_dict(columns)

Create a table from a mapping of column labels to column values. [Deprecated]

2.1.5 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.

2.1.6 datascience.tables.Table.from_df

classmethod Table.from_df(df)

Convert a Pandas DataFrame into a Table.
2.1.7 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 new 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 new table with an additional row.
- **Table.with_rows**(rows) Return a table with additional rows.
- **Table.relabeled**(label, new_label) Returns a new table with label specifying column label(s) replaced by corresponding.

2.1.8 datascience.tables.Table.with_column

Table.**with_column**(label, values)
Return a new table with an additional or replaced column.

**Args:**

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

- **values** (single value or sequence): If a single value, every value in the new column is values. If sequence of values, new column takes on values in values.

**Raises:**

- **ValueError**: If
  - label is not a valid column name
  - if label is not of type (str)
  - values is a list/array that does not have the same length as the number of rows in the table.

**Returns:** copy of original table with new or replaced column

```python
>>> alphabet = Table().with_column('letter', make_array('c','d'))
>>> alphabet = alphabet.with_column('count', make_array(2, 4))
>>> alphabet
letter | count
--- | ---
c | 2
d | 4

>>> alphabet.with_column('permutes', make_array('a', 'g'))
letter | count | permutes
--- | --- | ---
c | 2 | a
d | 4 | g

>>> alphabet
letter | count
--- | ---
c | 2
d | 4

>>> alphabet.with_column('count', 1)
Traceback (most recent call last):
...
ValueError: The column label must be a string, but a int was given
```
>>> alphabet.with_column('bad_col', make_array(1))
Traceback (most recent call last):
...
    ValueError: Column length mismatch. New column does not have the same number of rows as table.

2.1.9 data science.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-value pairs. If one of the labels is in existing table, then every value in the corresponding column is set to that value. If label has only a single value (int), every row of corresponding column takes on that value.

Raises:

ValueError: If

- any label in labels_and_values is not a valid column name, i.e if label is not of type (str).
- if any value in labels_and_values is a list/array and does not have the same length as the number of rows in the table.

AssertionError:

- ‘incorrect columns format’, if passed more than one sequence (iterables) for labels_and_values.
- ‘even length sequence required’ if missing a pair in label-value pairs.

Returns: Copy of original table with new or replaced columns. Columns added in order of labels. Equivalent to with_column(label, value) when passed only one label-value pair.

>>> players = Table().with_columns('player_id', ...
...    make_array(110234, 110235), 'wOBA', make_array(.354, .236))
>>> players
player_id | wOBA
110234   | 0.354
110235   | 0.236

>>> players = players.with_columns('salaries', 'N/A', 'season', 2016)
>>> players
player_id | wOBA | salaries | season
110234   | 0.354 | N/A      | 2016
110235   | 0.236 | N/A      | 2016

>>> salaries = Table().with_column('salary', ...
...    make_array('$500,000', '$15,500,000'))
>>> players.with_columns('salaries', salaries.column('salary'), ...
...    'years', make_array(6, 1))

>>> players.with_columns(2, make_array('$600,000', '$20,000,000'))
Traceback (most recent call last):
...
    ValueError: The column label must be a string, but a int was given
>>> players.with_columns('salaries', make_array('$600,000'))
Traceback (most recent call last):
2.1.10 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.

```python
>>> tiles = Table(make_array('letter', 'count', 'points'))
>>> tiles.with_row(['c', 2, 3]).with_row(['d', 4, 2])
letter | count | points
--- | --- | ---
c | 2 | 3
d | 4 | 2
```

2.1.11 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.

```python
>>> tiles = Table(make_array('letter', 'count', 'points'))
>>> tiles.with_rows(make_array(make_array('c', 2, 3),
... make_array('d', 4, 2)))
letter | count | points
--- | --- | ---
c | 2 | 3
d | 4 | 2
```

2.1.12 datascience.tables.Table.relabeled

Table.relabeled(label, new_label)
Returns a new table with label specifying column label(s) replaced by corresponding new_label.

Args:

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

Raises:

- ValueError – if label does not exist in table, or if the label and new_label are not of equal length. Also, raised if label and/or new_label are not str.

Returns: New table with new_label in place of label.
```python
>>> tiles = Table().with_columns('letter', make_array('c', 'd'),
... 'count', make_array(2, 4))

>>> tiles
letter | count
c    | 2
+-----+-----
d    | 4

>>> tiles.relabeled('count', 'number')
letter | number
letter c    | 2
|        |    |
letter d    | 4

>>> tiles.relabeled('# original table unmodified')
letter | count
letter c    | 2
|        |
letter d    | 4

>>> tiles.relabeled(make_array('letter', 'count'),
... ... make_array('column1', 'column2'))
column1 | column2
...    | 2
...    |
...    |
...    |

>>> tiles.relabeled(make_array('letter', 'number'),
... ... make_array('column1', 'column2'))
Traceback (most recent call last):
...
ValueError: Invalid labels. Column labels must already exist in table in order to be replaced.
```

Accessing values

<table>
<thead>
<tr>
<th>Method</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><code>Table.num_columns</code></td>
<td>Number of columns.</td>
</tr>
<tr>
<td><code>Table.columns</code></td>
<td>Return the values of a column as an array.</td>
</tr>
<tr>
<td><code>Table.num_rows</code></td>
<td>Number of rows.</td>
</tr>
<tr>
<td><code>Table.rows</code></td>
<td>Return a view of all rows.</td>
</tr>
<tr>
<td><code>Table.row(index)</code></td>
<td>Return a row.</td>
</tr>
<tr>
<td><code>Table.labels</code></td>
<td>Return a tuple of column labels.</td>
</tr>
<tr>
<td><code>Table.column_index(column_label)</code></td>
<td>Return the index of a column.</td>
</tr>
<tr>
<td><code>Table.apply(fn[, column_label])</code></td>
<td>Apply fn to each element of column_label.</td>
</tr>
</tbody>
</table>

**2.1.13 datascience.tables.Table.num_columns**

`Table.num_columns`

Number of columns.

**2.1.14 datascience.tables.Table.columns**

`Table.columns`

Return the values of a column as an array.

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

**2.1.15 datascience.tables.Table.column**

`Table.column(index_or_label)`

Return the values of a column as an array.
>>> tiles = Table().with_columns(
...    'letter', make_array('c', 'd'),
...    'count', make_array(2, 4),
...)

>>> tiles.column('letter')
array(['c', 'd'],
      dtype='<U1')

>>> tiles.column(1)
array([2, 4])

Args: label (int or str): The index or label of a column
Returns: An instance of numpy.array.
Raises: ValueError: When the index_or_label is not in the table.

2.1.16 datascience.tables.Table.num_rows

Table.num_rows
Number of rows.

2.1.17 datascience.tables.Table.rows

Table.rows
Return a view of all rows.

2.1.18 datascience.tables.Table.row

Table.row(index)
Return a row.

2.1.19 datascience.tables.Table.labels

Table.labels
Return a tuple of column labels.

2.1.20 datascience.tables.Table.column_index

Table.column_index(column_label)
Return the index of a column.

2.1.21 datascience.tables.Table.apply

Table.apply(fn, column_label=None)
Apply fn to each element of column_label. If no column_label provided, fn is applied to each row of table.

Args:

fn (function) – The function to be applied to elements of column_label.
column_label (single str or array of str) – Names of columns to be passed into fn. Length must match number of arguments in fn signature.

Raises:

* ValueError – if column_label is not an existing column in the table.
* TypeError – if insufficient number of column_label passed to fn.

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

```python
>>> t = Table().with_columns(
...    'letter', make_array('a', 'b', 'c', 'z'),
...    'count', make_array(9, 3, 3, 1),
...    'points', make_array(1, 2, 2, 10))
```

<table>
<thead>
<tr>
<th>letter</th>
<th>count</th>
<th>points</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>9</td>
<td>1</td>
</tr>
<tr>
<td></td>
<td>3</td>
<td>2</td>
</tr>
<tr>
<td></td>
<td>3</td>
<td>2</td>
</tr>
<tr>
<td></td>
<td>1</td>
<td>10</td>
</tr>
</tbody>
</table>

```python
>>> t.apply(lambda x: x - 1, 'points')
array([0, 1, 1, 9])
```

```python
>>> t.apply(lambda x, y: x * y, make_array('count', 'points'))
array([ 9, 6, 6, 10])
```

```python
>>> t.apply(lambda x: x - 1, make_array('count', 'points'))
Traceback (most recent call last):
...
TypeError: <lambda>() takes 1 positional argument but 2 were given
```

```python
>>> t.apply(lambda x: x - 1, 'counts')
Traceback (most recent call last):
...
ValueError: The column "counts" is not in the table. The table contains these columns: letter, count, points
```

Whole rows are passed to the function if no columns are specified.

```python
>>> t.apply(lambda row: row[1] * 2)
array([18, 6, 6, 2])
```

Mutation (modifies table in place)

<table>
<thead>
<tr>
<th>Method</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><code>table.set_format</code></td>
<td>Set the format of a column.</td>
</tr>
<tr>
<td><code>table.move_to_start</code></td>
<td>Move a column to the first in order.</td>
</tr>
<tr>
<td><code>table.move_to_end</code></td>
<td>Move a column to the last in order.</td>
</tr>
<tr>
<td><code>table.append</code></td>
<td>Append a row or all rows of a table.</td>
</tr>
<tr>
<td><code>table.append_column</code></td>
<td>Appends a column to the table or replaces a column.</td>
</tr>
<tr>
<td><code>table.relabel</code></td>
<td>Changes the label(s) of column(s) specified by column_label to labels in new_label.</td>
</tr>
</tbody>
</table>

2.1.22 | **datascience.tables.Table.set_format**

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

2.1.23 | **datascience.tables.Table.move_to_start**

Table.move_to_start (column_label) Move a column to the first in order.
2.1.24 datascience.tables.Table.move_to_end

Table.move_to_end(column_label)
Move a column to the last in order.

2.1.25 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.

2.1.26 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', make_array(1, 2, 3)) is equivalent to table['new_col'] = make_array(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.

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

>>> table
letter | count | points
a | 9 | 1
b | 3 | 2
c | 3 | 2
z | 1 | 10

>>> table.append_column('new_col1', make_array(10, 20, 30, 40))

>>> table
letter | count | points | new_col1
a | 9 | 1 | 10
b | 3 | 2 | 20
c | 3 | 2 | 30
z | 1 | 10 | 40

>>> table.append_column('new_col2', 'hello')

>>> table
letter | count | points | new_col1 | new_col2
a | 9 | 1 | 10 | hello
b | 3 | 2 | 20 | hello
c | 3 | 2 | 30 | hello
z | 1 | 10 | 40 | hello
```
>>> table.append_column(123, make_array(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.

2.1.27 `datascience.tables.Table.relabel`

Table.relabel(column_label, new_label)

Changes the label(s) of column(s) specified by column_label to labels in new_label.

Args:

- column_label – (single str or array of str) The label(s) of columns to be changed to new_label.
- new_label – (single str or array of str): The label name(s) of columns to replace column_label.

Raises:

- ValueError – if column_label is not in table, or if column_label and new_label are not of equal length.
- TypeError – if column_label and/or new_label is not str.

Returns: Original table with new_label in place of column_label.

```python
>>> table = Table().with_columns(
...   'points', make_array(1, 2, 3),
...   'id', make_array(12345, 123, 5123))

>>> table.relabel('id', 'yolo')
points | yolo
1     | 12345
2     | 123
3     | 5123

>>> table.relabel(make_array('points', 'yolo'),
...   make_array('red', 'blue'))
red    | blue
1     | 12345
2     | 123
3     | 5123

>>> table.relabel(make_array('red', 'green', 'blue'),
...   make_array('cyan', 'magenta', 'yellow', 'key'))
Traceback (most recent call last):
...
ValueError: Invalid arguments. column_label and new_label must be of equal length.
```
Table 2.5 – continued from previous page

<table>
<thead>
<tr>
<th>Method</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><code>Table.where(column_or_label[, ...])</code></td>
<td>Return a new <code>Table</code> containing rows where <code>value_or_predicate</code> returns True</td>
</tr>
<tr>
<td><code>Table.sort(column_or_label[, descending, ...])</code></td>
<td>Return a <code>Table</code> of rows sorted according to the values in a column.</td>
</tr>
<tr>
<td><code>Table.group(column_or_label[, collect])</code></td>
<td>Group rows by unique values in a column; count or aggregate others.</td>
</tr>
<tr>
<td><code>Table.pivot(columns, rows[, values, ...])</code></td>
<td>Generate a table with a column for each unique value in <code>columns</code>, with rows for others.</td>
</tr>
<tr>
<td><code>Table.stack(key[, labels])</code></td>
<td>Takes k original columns and returns two columns, with col.</td>
</tr>
<tr>
<td><code>Table.join(column_label, other[, other_label])</code></td>
<td>Creates a new <code>Table</code> with the columns of self and other, containing rows for all values of column_label.</td>
</tr>
<tr>
<td><code>Table.stats([ops])</code></td>
<td>Compute statistics for each column and place them in a table.</td>
</tr>
<tr>
<td><code>Table.percentile(p)</code></td>
<td>Returns a new table with one row containing the pth percentile for each column.</td>
</tr>
<tr>
<td><code>Table.sample([k, with replacement, weights])</code></td>
<td>Returns a new <code>Table</code> where k rows are randomly sampled from the original table.</td>
</tr>
<tr>
<td><code>Table.split(k)</code></td>
<td>Returns a tuple of two tables where the first table contains k rows randomly sampled.</td>
</tr>
<tr>
<td><code>Table.bin([select])</code></td>
<td>Group values by bin and compute counts per bin by column.</td>
</tr>
</tbody>
</table>

2.1.28 `datascience.tables.Table.copy`

`Table.copy(*, shallow=False)`

Return a copy of a `Table`.

2.1.29 `datascience.tables.Table.select`

`Table.select(*column_label_or_labels)`

Returns a new `Table` with only the columns in `column_label_or_labels`.

**Args:**
- `column_label_or_labels`: Columns to select from the `Table` as either column labels (`str`) or column indices (`int`).

**Returns:**
An new instance of `Table` containing only selected columns. The columns of the new `Table` are in the order given in `column_label_or_labels`.

**Raises:**
- `KeyError` if any of `column_label_or_labels` are not in the table.

```python
>>> flowers = Table().with_columns(
...     'Number of petals', make_array(8, 34, 5),
...     'Name', make_array('lotus', 'sunflower', 'rose'),
...     'Weight', make_array(10, 5, 6)
... )

>>> flowers
Number of petals | Name     | Weight |
8                | lotus    | 10     |
34               | sunflower| 5      |
5                | rose     | 6      |

>>> flowers.select('Number of petals', 'Weight')
Number of petals | Weight |
8                | 10     |
34               | 5      |
5                | 6      |

>>> flowers # original table unchanged
Number of petals | Name     | Weight |
8                | lotus    | 10     |
34               | sunflower| 5      |
5                | rose     | 6      |
```
```
>>> flowers.select(0, 2)
Number of petals | Weight
8 | 10
34 | 5
5 | 6
```

### 2.1.30 `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', make_array('cheeseburger', 'hamburger', 'veggie burger'),
...   'prices', make_array(6, 5, 5),
...   'calories', make_array(743, 651, 582))

>>> t
burgers | prices | calories
cheeseburger | 6 | 743
hamburger | 5 | 651
veggie burger | 5 | 582

>>> t.drop('prices')
burgers | calories
cheeseburger | 743
hamburger | 651
veggie burger | 582

>>> t.drop(['burgers', 'calories'])
prices
6
5
5

>>> t.drop('burgers', 'calories')
prices
6
5
5

>>> t.drop([0, 2])
prices
6
5
5

>>> t.drop(0, 2)
prices
6
5
5

>>> t.drop(1)
burgers | calories
cheeseburger | 743
hamburger | 651
veggie burger | 582
```
2.1.31 data science.tables.Table.take

Table.take()
Return a new Table with selected rows taken by index.

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

Returns: A new instance of Table with selected rows in order corresponding to row_indices_or_slice.

Raises: IndexError, if any row_indices_or_slice is out of bounds with respect to column length.

```python
grades = Table().with_columns('letter grade', make_array('A+', 'A', 'A-', 'B+', 'B', 'B-'), 'gpa', make_array(4, 4, 3.7, 3.3, 3, 2.7))

grades.take(0)
letter grade | gpa
A+ | 4

grades.take(-1)
letter grade | gpa
B- | 2.7

grades.take(make_array(2, 1, 0))
letter grade | gpa
A- | 3.7
A | 4
A+ | 4

grades.take[:3]
letter grade | gpa
A+ | 4
A | 4
A- | 3.7

grades.take(np.arange(0,3))
letter grade | gpa
A+ | 4
A | 4
A- | 3.7

grades.take(10)
Traceback (most recent call last):
...
  IndexError: index 10 is out of bounds for axis 0 with size 6
```

2.1.32 data science.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.

```python
>>> t = Table().with_columns(
...     'letter grade', make_array('A+', 'A', 'A-', 'B+', 'B', 'B-'),
...     'gpa', make_array(4, 4, 3.7, 3.3, 3, 2.7))
>>> t
letter grade | gpa
A+   | 4
A    | 4
A-   | 3.7
B+   | 3.3
B    | 3
B-   | 2.7

>>> t.exclude(4)
letter grade | gpa
A+   | 4
A    | 4
A-   | 3.7
B+   | 3.3
B-   | 2.7

>>> t.exclude(-1)
letter grade | gpa
A+   | 4
A    | 4
A-   | 3.7
B+   | 3.3
B    | 3

>>> t.exclude(make_array(1, 3, 4))
letter grade | gpa
A+   | 4
A-   | 3.7
B-   | 2.7

>>> t.exclude(range(3))
letter grade | gpa
B+   | 3.3
B    | 3
B-   | 2.7
```

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

```python
>>> t.exclude[:3]
letter grade | gpa
B+   | 3.3
B    | 3
B-   | 2.7

>>> t.exclude[1, 3, 4]
letter grade | gpa
A+   | 4
A-   | 3.7
B-   | 2.7
```

2.1.33 `datascience.tables.Table.where`

Table.

```
Table.where(column_or_label, value_or_predicate=None, other=None)
```

Return a new Table containing rows where `value_or_predicate` returns True for values in `column_or_label`.
Args: column_or_label: A column of the Table either as a label (str) or an index (int). Can also be an array of booleans; only the rows where the array value is True are kept.

value_or_predicate: If a function, it is applied to every value in column_or_label. Only the rows where value_or_predicate returns True are kept. If a single value, only the rows where the values in column_or_label are equal to value_or_predicate are kept.

other: Optional additional column label for value_or_predicate to make pairwise comparisons. See the examples below for usage. When other is supplied, value_or_predicate must be a callable function.

Returns: If value_or_predicate is a function, returns a new Table containing only the rows where value_or_predicate(val) is True for the val’s in column_or_label.

If value_or_predicate is a value, returns a new Table containing only the rows where the values in column_or_label are equal to value_or_predicate.

If column_or_label is an array of booleans, returns a new Table containing only the rows where column_or_label is True.

```python
>>> marbles = Table().with_columns(  
...   "Color", make_array("Red", "Green", "Blue",  
...     "Red", "Green", "Green"),  
...   "Shape", make_array("Round", "Rectangular", "Rectangular",  
...     "Round", "Rectangular", "Round"),  
...   "Amount", make_array(4, 6, 12, 7, 9, 2),  
...   "Price", make_array(1.30, 1.20, 2.00, 1.75, 0, 3.00))

>>> marbles
Color | Shape | Amount | Price
Red   | Round | 4     | 1.3
Green | Rectangular | 6 | 1.2
Blue  | Rectangular | 12 | 2
Red   | Round | 7     | 1.75
Green | Rectangular | 9 | 0
Green | Round | 2     | 3

Use a value to select matching rows

```python
>>> marbles.where("Price", 1.3)
Color | Shape | Amount | Price
Red   | Round | 4     | 1.3

In general, a higher order predicate function such as the functions in datascience.predicates.are can be used.

```python
>>> from datascience.predicates import are
>>> # equivalent to previous example
>>> marbles.where("Price", are.equal_to(1.3))
Color | Shape | Amount | Price
Red   | Round | 4     | 1.3

>>> marbles.where("Price", are.above(1.5))
Color | Shape | Amount | Price
Blue  | Rectangular | 12 | 2
Red   | Round | 7     | 1.75
Green | Round | 2     | 3

Use the optional argument other to apply predicates to compare columns.
>>> marbles.where("Price", are.above, "Amount")
Color | Shape | Amount | Price
Green | Round | 2 | 3

>>> marbles.where("Price", are.equal_to, "Amount") # empty table
Color | Shape | Amount | Price

2.1.34 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", make_array("Red", "Green", "Blue", "Red", "Green", "Green"), ...  "Shape", make_array("Round", "Rectangular", "Rectangular", "Round", "Rectangular", "Round"), ...  "Amount", make_array(4, 6, 12, 7, 9, 2), ...  "Price", make_array(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 | 1.3
Blue | Rectangular | 12 | 2
Red | Round | 7 | 1.75
Green | Rectangular | 9 | 1.4
Green | Round | 2 | 1

>>> marbles.sort("Amount")
Color | Shape | Amount | Price
Green | Round | 2 | 1
Red | Round | 4 | 1.3
Green | Rectangular | 6 | 1.3
Red | Round | 7 | 1.75
Green | Rectangular | 9 | 1.4
Blue | Rectangular | 12 | 2

>>> marbles.sort("Amount", descending = True)
Color | Shape | Amount | Price
Blue | Rectangular | 12 | 2
Green | Rectangular | 9 | 1.4
Red | Round | 7 | 1.75
Green | Rectangular | 6 | 1.3
Red | Round | 4 | 1.3
Green | Round | 2 | 1

>>> marbles.sort(3) # the Price column
Color | Shape | Amount | Price
Color | Shape | Amount | Price
Green | Round | 2 | 1
Red | Round | 4 | 1.3
Green | Rectangular | 6 | 1.3
Green | Rectangular | 9 | 1.4
Red | Round | 7 | 1.75
Blue | Rectangular | 12 | 2

>>> marbles.sort(3, distinct = True)
<table>
<thead>
<tr>
<th>Color</th>
<th>Shape</th>
<th>Amount</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td>Green</td>
<td>Round</td>
<td>2</td>
<td>1</td>
</tr>
<tr>
<td>Red</td>
<td>Round</td>
<td>4</td>
<td>1.3</td>
</tr>
<tr>
<td>Green</td>
<td>Rectangular</td>
<td>9</td>
<td>1.4</td>
</tr>
<tr>
<td>Red</td>
<td>Round</td>
<td>7</td>
<td>1.75</td>
</tr>
<tr>
<td>Blue</td>
<td>Rectangular</td>
<td>12</td>
<td>2</td>
</tr>
</tbody>
</table>

### 2.1.35 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 column_or_label, where the first column contains the unique values from column_or_label, and the second contains 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 column_or_label, then applying collect 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.

```python
def marbles = Table().with_columns(
    ... "Color", make_array("Red", "Green", "Blue", "Red", "Green", "Green"),
    ... "Shape", make_array("Round", "Rectangular", "Rectangular", "Round", "Rectangular", "Round"),
    ... "Amount", make_array(4, 6, 12, 7, 9, 2),
    ... "Price", make_array(1.30, 1.30, 2.00, 1.75, 1.40, 1.00))
```

```python
>>> marbles
<table>
<thead>
<tr>
<th>Color</th>
<th>Shape</th>
<th>Amount</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td>Red</td>
<td>Round</td>
<td>4</td>
<td>1.3</td>
</tr>
<tr>
<td>Green</td>
<td>Rectangular</td>
<td>6</td>
<td>1.3</td>
</tr>
<tr>
<td>Blue</td>
<td>Rectangular</td>
<td>12</td>
<td>2</td>
</tr>
<tr>
<td>Red</td>
<td>Round</td>
<td>7</td>
<td>1.75</td>
</tr>
<tr>
<td>Green</td>
<td>Rectangular</td>
<td>9</td>
<td>1.4</td>
</tr>
<tr>
<td>Green</td>
<td>Round</td>
<td>2</td>
<td>1</td>
</tr>
</tbody>
</table>
```

```python
>>> marbles.group("Color")  # just gives counts
<table>
<thead>
<tr>
<th>Color</th>
<th>count</th>
</tr>
</thead>
<tbody>
<tr>
<td>Blue</td>
<td>1</td>
</tr>
<tr>
<td>Green</td>
<td>3</td>
</tr>
<tr>
<td>Red</td>
<td>2</td>
</tr>
</tbody>
</table>
```

```python
>>> marbles.group("Color", max)  # takes the max of each grouping, in each column
<table>
<thead>
<tr>
<th>Color</th>
<th>Shape max</th>
<th>Amount max</th>
<th>Price max</th>
</tr>
</thead>
<tbody>
<tr>
<td>Blue</td>
<td>Rectangular</td>
<td>12</td>
<td>2</td>
</tr>
</tbody>
</table>
```

```python
>>> marbles.group("Shape", sum)  # sum doesn’t make sense for strings
<table>
<thead>
<tr>
<th>Shape</th>
<th>Color sum</th>
<th>Amount sum</th>
<th>Price sum</th>
</tr>
</thead>
<tbody>
<tr>
<td>Rectangular</td>
<td>27</td>
<td>4.7</td>
<td></td>
</tr>
<tr>
<td>Round</td>
<td>13</td>
<td>4.05</td>
<td></td>
</tr>
</tbody>
</table>
```
2.1.36 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.

```python
>>> marbles = Table().with_columns(
    ...   "Color", make_array("Red", "Green", "Blue", "Red", "Green", "Green"),
    ...   "Shape", make_array("Round", "Rectangular", "Rectangular", "Round", "Rectangular", "Round"),
    ...   "Amount", make_array(4, 6, 12, 7, 9, 2),
    ...   "Price", make_array(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       | 1.3
Blue  | Rectangular | 12     | 2
Red   | Round   | 7      | 1.75
Green | Rectangular | 9      | 1.4
Green | Round   | 2      | 1

>>> marbles.groups(["Color", "Shape"])

<table>
<thead>
<tr>
<th>Color</th>
<th>Shape</th>
<th>count</th>
</tr>
</thead>
<tbody>
<tr>
<td>Blue</td>
<td>Rectangular</td>
<td>1</td>
</tr>
<tr>
<td>Green</td>
<td>Rectangular</td>
<td>2</td>
</tr>
<tr>
<td>Green</td>
<td>Round</td>
<td>1</td>
</tr>
<tr>
<td>Red</td>
<td>Round</td>
<td>2</td>
</tr>
</tbody>
</table>

>>> marbles.groups(["Color", "Shape"], sum)

<table>
<thead>
<tr>
<th>Color</th>
<th>Shape</th>
<th>Amount sum</th>
<th>Price sum</th>
</tr>
</thead>
<tbody>
<tr>
<td>Blue</td>
<td>Rectangular</td>
<td>12</td>
<td>2</td>
</tr>
<tr>
<td>Green</td>
<td>Rectangular</td>
<td>15</td>
<td>2.7</td>
</tr>
<tr>
<td>Green</td>
<td>Round</td>
<td>2</td>
<td>1</td>
</tr>
<tr>
<td>Red</td>
<td>Round</td>
<td>11</td>
<td>3.05</td>
</tr>
</tbody>
</table>
```

2.1.37 datascience.tables.Table.pivot

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

Generate a table with a column for each unique value in `columns`, with rows for each unique value in `rows`. Each row counts/aggregates the values that match both row and column based on `collect`.

**Args:**
- **columns** – a single column label, (str), in table, used to create new columns, based on its unique values.
- **rows** – row labels, as (str) or array of strings, used to create new rows based on it’s unique values.
- **values** – column label in table for use in aggregation. Default None.
collect – aggregation function, used to group values over row-column combinations. Default None.

zero – zero value for non-existent row-column combinations.

Raises:

TypeError – if collect is passed in and values is not, vice versa.

Returns: New pivot table, with row-column combinations, as specified, with aggregated values by collect across the intersection of columns and rows. Simple counts provided if values and collect are None, as default.

```python
>>> titanic = Table().with_columns('age', make_array(21, 44, 56, 89, 95,..., 40, 80, 45), 'survival', make_array(0,0,0,1, 1, 1, 0, 1),... 'gender', make_array('M', 'M', 'M', 'M', 'F', 'F', 'F', 'F'),... 'prediction', make_array(0, 0, 1, 1, 0, 1, 0, 1))
>>> titanic
age | survival | gender | prediction
21 | 0 | M | 0
44 | 0 | M | 0
56 | 0 | M | 1
89 | 1 | M | 1
95 | 1 | F | 0
40 | 1 | F | 1
80 | 0 | F | 0
45 | 1 | F | 1
```

```python
>>> titanic.pivot('survival', 'gender')
gender | 0 | 1
F | 1 | 3
M | 3 | 1
>>> titanic.pivot('prediction', 'gender')
gender | 0 | 1
F | 2 | 2
M | 2 | 2
>>> titanic.pivot('survival', 'gender', values='age', collect = np.mean)
gender | 0 | 1
F | 80 | 60
M | 40.3333 | 89
>>> titanic.pivot('survival', make_array('prediction', 'gender'))
prediction | gender | 0 | 1
0 | F | 1 | 1
0 | M | 2 | 0
1 | F | 0 | 2
1 | M | 1 | 1
>>> titanic.pivot('survival', 'gender', values = 'age')
Traceback (most recent call last):
... TypeError: values requires collect to be specified
>>> titanic.pivot('survival', 'gender', collect = np.mean)
Traceback (most recent call last):
... TypeError: collect requires values to be specified
```

2.1.38 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.

## 2.1.39 datascience.tables.Table.join

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

Creates a new table with the columns of self and other, containing rows for all values of a column that appear in both tables.

**Args:**

- column_label (str): label of column in self that is used to join rows of other.
- other: Table object to join with self on matching values of column_label.

**Kwargs:**

- other_label (str): default None, assumes column_label. Otherwise in other used to join rows.

**Returns:** New table self joined with other by matching values in column_label and other_label. If the resulting join is empty, returns None. If a join value appears more than once in self, each row with that value will appear in resulting join, but in other, only the first row with that value will be used.

```python
>>> table = Table().with_columns('a', make_array(9, 3, 3, 1),
...   'b', make_array(1, 2, 2, 10),
...   'c', make_array(3, 4, 5, 6))
>>> table
a | b | c
9 | 1 | 3
3 | 2 | 4
3 | 2 | 5
1 | 10| 6
>>> table2 = Table().with_columns('a', make_array(9, 1, 1, 1),
...   'd', make_array(1, 2, 2, 10),
...   'e', make_array(3, 4, 5, 6))
>>> table2
a | d | e
9 | 1 | 3
1 | 2 | 4
1 | 2 | 5
1 | 10| 6
>>> table.join('a', table2)
a | b | c | d | e
9 | 1 | 3 | 1 | 3
1 | 10| 6 | 2 | 4
>>> table.join('a', table2, 'a')  # Equivalent to previous join
a | b | c | d | e
1 | 10| 6 | 2 | 4
9 | 1 | 3 | 1 | 3
>>> table.join('a', table2, 'd')  # Repeat column labels relabeled
a | b | c | a_2 | e
1 | 10| 6 | 9 | 3
>>> table2  #table2 has three rows with a = 1
a | d | e
9 | 1 | 3
1 | 2 | 4
1 | 2 | 5
1 | 10| 6
>>> table  #table has only one row with a = 1
a | b | c
9 | 1 | 3
1 | 2 | 4
1 | 2 | 5
1 | 10| 6
```
2.1.40 datascience.tables.Table.stats

Table.stats(ops=(<built-in function min>, <built-in function max>, <function median at 0x7f9e4f88f840>, <built-in function sum>))

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

2.1.41 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 least as large as the p% of numbers in the column.

>>> table = Table().with_columns(
... 'count', make_array(9, 3, 3, 1),
... 'points', make_array(1, 2, 2, 10))
>>> table
count | points
9 | 1
3 | 2
3 | 2
1 | 10

>>> table.percentile(80)
count | points
9 | 10

2.1.42 datascience.tables.Table.sample

Table.sample(k=None, with_replacement=True, weights=None)

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

Args:

k – specifies the number of rows (int) to be sampled from the table. Default is k equal to number of rows in the table.
with_replacement – (bool) By default True; Samples k rows with replacement from table, else samples k rows without replacement.

weights – Array specifying probability the ith row of the table is sampled. Defaults to None, which samples each row with equal probability. weights must be a valid probability distribution – i.e. an array the length of the number of rows, summing to 1.

Raises:

ValueError – if weights is not length equal to number of rows in the table; or, if weights does not sum to 1.

Returns: A new instance of Table with k rows resampled.

```python
>>> jobs = Table().with_columns(
...   'job', make_array('a', 'b', 'c', 'd'),
...   'wage', make_array(10, 20, 15, 8))
>>> jobs
job  | wage
a    | 10
b    | 20
с    | 15
d    | 8

>>> jobs.sample()  # jobs without replacement
job  | wage
b    | 20
b    | 20
a    | 10
d    | 8

>>> jobs.sample(with_replacement=True)  # jobs with replacement
job  | wage
d    | 8
b    | 20
c    | 15
a    | 10

>>> jobs.sample(k = 2)  # sample 2 rows
job  | wage
b    | 20
c    | 15

>>> jobs.sample(k = 2, with_replacement = True,
...   weights = make_array(0.5, 0.5, 0, 0))  # with replacement
job  | wage
a    | 10
c    | 10

>>> jobs.sample(k = 2, weights = make_array(1, 0, 1, 0))
Traceback (most recent call last):
...
ValueError: probabilities do not sum to 1

# Weights must be length of table.
```
### 2.1.43 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 \(\text{num\_rows} - 1\).

**Raises:** ValueError: \(k\) is not between 1 and \(\text{num\_rows} - 1\).

**Returns:** A tuple containing two instances of Table.

```python
>>> jobs = Table().with_columns('job', make_array('a', 'b', 'c', 'd'),
                              'wage', make_array(10, 20, 15, 8))
>>> jobs
job | wage
-- | ----
a  | 10
b  | 20
c  | 15
d  | 8

>>> sample, rest = jobs.split(3)
>>> sample
job | wage
-- | ----
c  | 15
a  | 10
b  | 20

>>> rest
job | wage
-- | ----
d  | 8
```

### 2.1.44 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>
<thead>
<tr>
<th>Method</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><code>Table.show([max_rows])</code></td>
<td>Display the table.</td>
</tr>
<tr>
<td><code>Table.as_text([max_rows, sep])</code></td>
<td>Format table as text.</td>
</tr>
<tr>
<td><code>Table.as_html([max_rows])</code></td>
<td>Format table as HTML.</td>
</tr>
<tr>
<td><code>Table.index_by(column_or_label)</code></td>
<td>Return a dict keyed by values in a column that contains lists of rows corresponding to each value.</td>
</tr>
<tr>
<td><code>Table.to_array()</code></td>
<td>Convert the table to a structured NumPy array.</td>
</tr>
<tr>
<td><code>Table.to_df()</code></td>
<td>Convert the table to a Pandas DataFrame.</td>
</tr>
<tr>
<td><code>Table.to_csv(filename)</code></td>
<td>Creates a CSV file with the provided filename.</td>
</tr>
</tbody>
</table>

### 2.1.45 datascience.tables.Table.show

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

### 2.1.46 datascience.tables.Table.as_text

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

### 2.1.47 datascience.tables.Table.as_html

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

### 2.1.48 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.

### 2.1.49 datascience.tables.Table.to_array

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

### 2.1.50 datascience.tables.Table.to_df

`Table.to_df()`  
Convert the table to a Pandas DataFrame.

### 2.1.51 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.

```python
>>> jobs = Table().with_columns(
...   'job', make_array('a', 'b', 'c', 'd'),
...   'wage', make_array(10, 20, 15, 8))
>>> jobs
job | wage
a | 10
b | 20
c | 15
d | 8
>>> jobs.to_csv('my_table.csv')
<outputs a file called my_table.csv in the current directory>
```

Visualizations

<table>
<thead>
<tr>
<th>Method</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><code>Table.plot()</code></td>
<td>Plot line charts for the table.</td>
</tr>
<tr>
<td><code>Table.bar()</code></td>
<td>Plot bar charts for the table.</td>
</tr>
<tr>
<td><code>Table.barh()</code></td>
<td>Plot horizontal bar charts for the table.</td>
</tr>
<tr>
<td><code>Table.pivot_hist()</code></td>
<td>Draw histograms of each category in a column.</td>
</tr>
<tr>
<td><code>Table.hist()</code></td>
<td>Plots one histogram for each column in the table.</td>
</tr>
<tr>
<td><code>Table.scatter()</code></td>
<td>Creates scatterplots, optionally adding a line of best fit.</td>
</tr>
<tr>
<td><code>Table.boxplot()</code></td>
<td>Plots a boxplot for the table.</td>
</tr>
</tbody>
</table>

# 2.1.52 datascience.tables.Table.plot

`Table.plot(column_for_xticks=None, select=None, overlay=True, width=6, height=4, **vargs)`

Plot line charts for the table.

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

**Raises:** ValueError – Every selected column must be numerical.

**Returns:** Returns a line plot (connected scatter). Each plot is labeled using the values in `column_for_xticks` and one plot is produced for all other columns in self (or for the columns designated by `select`).

```python
>>> table = Table().with_columns(
...   'days', make_array(0, 1, 2, 3, 4, 5),
...   'price', make_array(90.5, 90.00, 83.00, 95.50, 82.00, 82.00),
...   'projection', make_array(90.75, 82.00, 82.50, 82.50, 83.00, 82.50))
>>> table
days | price | projection
0 | 90.5 | 90.75
1 | 90 | 82
2 | 83 | 82.5
3 | 95.5 | 82.5
4 | 82 | 83
5 | 82 | 82.5
>>> table.plot('days')
```

# 2.1. Tables (datascience.tables)
2.1.53 datascience.tables.Table.bar

Table.bar(column_for_categories=None, select=None, overlay=True, width=6, height=4, **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.pyplot.bar for additional arguments that can be passed into vargs.

2.1.54 datascience.tables.Table.barh

Table.barh(column_for_categories=None, select=None, overlay=True, width=6, **vargs)

Plot horizontal bar charts for the table.

Args:

column_for_categories (str): A column containing y-axis categories used to create buckets for bar chart.

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.pyplot.barh for additional arguments that can be passed into vargs.

Raises:

ValueError – Every selected except column for column_for_categories must be numerical.

Returns: Horizontal bar graph with buckets specified by column_for_categories. 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).

```python
>>> t = Table().with_columns(
...     'Furniture', make_array('chairs', 'tables', 'desks'),
...     'Count', make_array(6, 1, 2),
...     'Price', make_array(10, 20, 30)
... )
>>> t
Furniture | Count | Price
---|---|---
chairs | 6 | 10
tables | 1 | 20
desks | 2 | 30
```
chairs | 6 | 10
tables | 1 | 20
desks | 2 | 30

```python
>>> furniture_table.barh('Furniture')
<bar graph with furniture as categories and bars for count and price>
>>> furniture_table.barh('Furniture', 'Price')
<bar graph with furniture as categories and bars for price>
>>> furniture_table.barh('Furniture', make_array(1, 2))
<bar graph with furniture as categories and bars for count and price>
```

### 2.1.55 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.

### 2.1.56 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.pyplot.hist for additional arguments that can be passed into vargs. These include: range, normed, cumulative, and orientation, to name a few.

```python
>>> t = Table().with_columns(... 'count', make_array(9, 3, 3, 1),
... 'points', make_array(1, 2, 2, 10))

>>> t
count | points
9 | 1
3 | 2
3 | 2
1 | 10

>>> t.hist()
<histogram of values in count>
<histogram of values in points>
```

```python
>>> t = Table().with_columns(... 'value', make_array(101, 102, 103),
... 'proportion', make_array(0.25, 0.5, 0.25))

>>> t.hist(counts='value')
<histogram of values in prop weighted by corresponding values in value>
```
2.1.57 datascience.tables.Table.scatter

Table.scatter(column_for_x, select=None, overlay=True, fit_line=False, colors=None, labels=None, width=5, height=5, **vargs)

Creates scatterplots, optionally adding a line of best fit.

**Args:**

- **column_for_x (str):** The column to use for the x-axis values and label of the scatter plots.

**Kwargs:**

- **overlay (bool):** If true, creates a chart with one color per data column; if False, each plot 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.pyplot.scatter](http://matplotlib.org/api/pyplot_api.html#matplotlib.pyplot.scatter) for additional arguments that can be passed into vargs. These include: marker and norm, to name a couple.
- **colors:** A column of colors (labels or numeric values).
- **labels:** A column of text labels to annotate dots.

**Raises:** ValueError – Every column, column_for_x or select, must be numerical

**Returns:** Scatter plot of values of column_for_x plotted against values for all other columns in self. Each plot uses the values in column_for_x for horizontal positions. One plot is produced for all other columns in self as y (or for the columns designated by select).

```python
>>> table = Table().with_columns(
...   'x', make_array(9, 3, 3, 1),
...   'y', make_array(1, 2, 2, 10),
...   'z', make_array(3, 4, 5, 6))

>>> table
x | y | z  
9 | 1 | 3  
3 | 2 | 4  
3 | 2 | 5  
1 | 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>
```

2.1.58 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.pyplot.boxplot 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.

```python
>>> table = Table().with_columns(
...    'test1', make_array(92.5, 88, 72, 71, 99, 100, 95, 83, 94, 93),
...    'test2', make_array(89, 84, 74, 66, 92, 99, 88, 81, 95, 94))
```

```python
>>> 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.

```python
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:

    ‘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].

```copy ()
Copies the current Map into a new one and returns it.
features

format(**kwargs)
    Apply formatting.

geojson()
    Render features as a FeatureCollection.

overlay(feature, color='Blue', opacity=0.6)
    Overlays feature on the map. Returns a new Map.

    Args:
        feature: a Table of map features, a list of map features, a Map, a Region, or a circle marker
        map table. The features will be overlayed on the Map with specified color.
        color (str): Color of feature. Defaults to ‘Blue’
        opacity (float): Opacity of overlain feature. Defaults to 0.6.

    Returns: A new Map with the overlain feature.

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
    marker
    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(latitude, longitude, labels=None, colors=None, radii=None, **kwargs)
    Return markers from columns of coordinates, labels, & colors.
    The radii column is not applicable to markers, but sets circle radius.

classmethod map_table(table, **kwargs)
    Return markers from the columns 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

For example, to draw three circles:

t = Table().with_columns([  
    'lat', [37.8, 38, 37.9], 'lon', [-122, -122.1, -121.9], 'label', ['one', 'two', 'three'], 'color', ['red', 'green', 'blue'], 'radius', [3000, 4000, 5000],  
])  
Circle.map_table(t)

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.make_array (*elements)   
    Returns an array containing all the arguments passed to this function. A simple way to make an array with a few elements.
    
    As with any array, all arguments should have the same type.

    >>> make_array(0)
    array([0])
    >>> make_array(2, 3, 4)
    array([2, 3, 4])
    >>> make_array("foo", "bar")
    array(['foo', 'bar'],
## 2.4. Utility Functions (datascience.util)

**dtype='\'<U3'\')**

```python
>>> make_array()
array([], dtype=float64)
```

**datascience.util.percentile\( (p, arr=None)\)**

Returns the \(p\)th 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, \texttt{percentile} returns itself curried with \(p\)

```python
>>> percentile(74.9, [1, 3, 5, 9])
5
>>> percentile(75, [1, 3, 5, 9])
5
>>> percentile(75.1, [1, 3, 5, 9])
9
>>> f = percentile(75)
>>> f([1, 3, 5, 9])
5
```

**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
- \(lbound\) (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.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 \texttt{apply} under the hood, then converts back to a Table

**Args:**
- \(table\) [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.proportions_from_distribution\( (table, label, sample_size, column_name='Random Sample')\)**

Adds a column named \texttt{column_name} containing the proportions of a random draw using the distribution in \texttt{label}.
This method uses `np.random.multinomial` to draw `sample_size` samples from the distribution in `table.column(label)`, then divides by `sample_size` to create the resulting column of proportions.

Returns a new `Table` and does not modify `table`.

Args: `table`: An instance of `Table`.

  - `label`: Label of column in `table`. This column must contain a distribution (the values must sum to 1).
  - `sample_size`: The size of the sample to draw from the distribution.
  - `column_name`: The name of the new column that contains the sampled proportions. Defaults to ‘Random Sample’.

Returns: A copy of `table` with a column `column_name` containing the sampled proportions. The proportions will sum to 1.

Throws:

  - `ValueError`: If the `label` is not in the table, or if `table.column(label)` does not sum to 1.

`datascience.util.minimize(f, start=None, smooth=False, log=None, array=False, **vargs)`

Minimize a function `f` of one or more arguments.

Args: `f`: A function that takes numbers and returns a number

  - `start`: A starting value or list of starting values
  - `smooth`: Whether to assume that `f` is smooth and use first-order info
  - `log`: Logging function called on the result of optimization (e.g. print)
  - `vargs`: Other named arguments passed to scipy.optimize.minimize

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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