

## 3 Pandas II

### Learning Outcomes

- Continue building familiarity with `pandas` syntax.
- Extract data from a `DataFrame` using conditional selection.
- Recognize situations where aggregation is useful and identify the correct technique for performing an aggregation.

Last time, we introduced the `pandas` library as a toolkit for processing data. We learned the `DataFrame` and `Series` data structures, familiarized ourselves with the basic syntax for manipulating tabular data, and began writing our first lines of `pandas` code.

In this lecture, we'll start to dive into some advanced `pandas` syntax. You may find it helpful to follow along with a notebook of your own as we walk through these new pieces of code.

We'll start by loading the `babynames` dataset.

#### ► Code

|   | State | Sex | Year | Name     | Count |
|---|-------|-----|------|----------|-------|
| 0 | CA    | F   | 1910 | Mary     | 295   |
| 1 | CA    | F   | 1910 | Helen    | 239   |
| 2 | CA    | F   | 1910 | Dorothy  | 220   |
| 3 | CA    | F   | 1910 | Margaret | 163   |
| 4 | CA    | F   | 1910 | Frances  | 134   |

### 3.1 Conditional Selection

Conditional selection allows us to select a subset of rows in a `DataFrame` that satisfy some specified condition.

To understand how to use conditional selection, we must look at another possible input of the `.loc` and `[]` methods – a boolean array, which is simply an array or `Series` where each element is either `True` or `False`. This boolean array must have a length equal to the number of rows in the `DataFrame`. It will return all rows that correspond to a value of `True` in the array. We used a very similar technique when performing conditional extraction from a `Series` in the last lecture.

To see this in action, let's select all even-indexed rows in the first 10 rows of our `DataFrame`.

```
# Ask yourself: why is :9 is the correct slice to select the first 10 rows?
babynames_first_10_rows = babynames.loc[:9, :]
```

```
# Notice how we have exactly 10 elements in our boolean array argument
babynames_first_10_rows[[True, False, True, False, True, False, True, False, True, False]]
```

|   | State | Sex | Year | Name     | Count |
|---|-------|-----|------|----------|-------|
| 0 | CA    | F   | 1910 | Mary     | 295   |
| 2 | CA    | F   | 1910 | Dorothy  | 220   |
| 4 | CA    | F   | 1910 | Frances  | 134   |
| 6 | CA    | F   | 1910 | Evelyn   | 126   |
| 8 | CA    | F   | 1910 | Virginia | 101   |

We can perform a similar operation using `.loc`.

```
babynames_first_10_rows.loc[[True, False, True, False, True, False, True, False, True, False]]
```

|   | State | Sex | Year | Name     | Count |
|---|-------|-----|------|----------|-------|
| 0 | CA    | F   | 1910 | Mary     | 295   |
| 2 | CA    | F   | 1910 | Dorothy  | 220   |
| 4 | CA    | F   | 1910 | Frances  | 134   |
| 6 | CA    | F   | 1910 | Evelyn   | 126   |
| 8 | CA    | F   | 1910 | Virginia | 101   |

These techniques worked well in this example, but you can imagine how tedious it might be to list out `True` and `False` for every row in a larger `DataFrame`. To make things easier, we can instead provide a logical condition as an input to `.loc` or `[]` that returns a boolean array with the necessary length.

For example, to return all names associated with `F` sex:

```
# First, use a logical condition to generate a boolean array
logical_operator = (babynames["Sex"] == "F")

# Then, use this boolean array to filter the DataFrame
babynames[logical_operator].head()
```

|   | State | Sex | Year | Name     | Count |
|---|-------|-----|------|----------|-------|
| 0 | CA    | F   | 1910 | Mary     | 295   |
| 1 | CA    | F   | 1910 | Helen    | 239   |
| 2 | CA    | F   | 1910 | Dorothy  | 220   |
| 3 | CA    | F   | 1910 | Margaret | 163   |
| 4 | CA    | F   | 1910 | Frances  | 134   |

Recall from the previous lecture that `.head()` will return only the first few rows in the `DataFrame`. In reality, `babynames[logical_operator]` contains as many rows as there are entries in the original `babynames`

`DataFrame` with sex "F".

Here, `logical_operator` evaluates to a `Series` of boolean values with length 407428.

#### ► Code

There are a total of 407428 values in 'logical\_operator'

Rows starting at row 0 and ending at row 239536 evaluate to `True` and are thus returned in the `DataFrame`. Rows from 239537 onwards evaluate to `False` and are omitted from the output.

#### ► Code

The 0th item in this 'logical\_operator' is: True

The 239536th item in this 'logical\_operator' is: True

The 239537th item in this 'logical\_operator' is: False

Passing a `Series` as an argument to `babynames[]` has the same effect as using a boolean array. In fact, the `[]` selection operator can take a boolean `Series`, array, and list as arguments. These three are used interchangeably throughout the course.

We can also use `.loc` to achieve similar results.

```
babynames.loc[babynames["Sex"] == "F"].head()
```

|   | State | Sex | Year | Name     | Count |
|---|-------|-----|------|----------|-------|
| 0 | CA    | F   | 1910 | Mary     | 295   |
| 1 | CA    | F   | 1910 | Helen    | 239   |
| 2 | CA    | F   | 1910 | Dorothy  | 220   |
| 3 | CA    | F   | 1910 | Margaret | 163   |
| 4 | CA    | F   | 1910 | Frances  | 134   |

Boolean conditions can be combined using various bitwise operators, allowing us to filter results by multiple conditions. In the table below, p and q are boolean arrays or `Series`.

| Symbol | Usage | Meaning                |
|--------|-------|------------------------|
| ~      | ~p    | Returns negation of p  |
|        | p   q | p OR q                 |
| &      | p & q | p AND q                |
| ^      | p ^ q | p XOR q (exclusive or) |

When combining multiple conditions with logical operators, we surround each individual condition with a set of parenthesis `()`. This imposes an order of operations on `pandas` evaluating your logic and can avoid code

erroring.

For example, if we want to return data on all names with sex "F" born before the year 2000, we can write:

```
babynames[(babynames["Sex"] == "F") & (babynames["Year"] < 2000)].head()
```

|   | State | Sex | Year | Name     | Count |
|---|-------|-----|------|----------|-------|
| 0 | CA    | F   | 1910 | Mary     | 295   |
| 1 | CA    | F   | 1910 | Helen    | 239   |
| 2 | CA    | F   | 1910 | Dorothy  | 220   |
| 3 | CA    | F   | 1910 | Margaret | 163   |
| 4 | CA    | F   | 1910 | Frances  | 134   |

Note that we’re working with Series, so using and in place of &, or or in place | will error.

```
# This line of code will raise a ValueError
# babynames[(babynames["Sex"] == "F") and (babynames["Year"] < 2000)].head()
```

If we want to return data on all names with sex "F" or all born before the year 2000, we can write:

```
babynames[(babynames["Sex"] == "F") | (babynames["Year"] < 2000)].head()
```

|   | State | Sex | Year | Name     | Count |
|---|-------|-----|------|----------|-------|
| 0 | CA    | F   | 1910 | Mary     | 295   |
| 1 | CA    | F   | 1910 | Helen    | 239   |
| 2 | CA    | F   | 1910 | Dorothy  | 220   |
| 3 | CA    | F   | 1910 | Margaret | 163   |
| 4 | CA    | F   | 1910 | Frances  | 134   |

Boolean array selection is a useful tool, but can lead to overly verbose code for complex conditions. In the example below, our boolean condition is long enough to extend for several lines of code.

```
# Note: The parentheses surrounding the code make it possible to break the code on to mul
(
    babynames[(babynames["Name"] == "Bella") |
                (babynames["Name"] == "Alex") |
                (babynames["Name"] == "Ani") |
                (babynames["Name"] == "Lisa")]
).head()
```

|      | State | Sex | Year | Name  | Count |
|------|-------|-----|------|-------|-------|
| 6289 | CA    | F   | 1923 | Bella | 5     |

|       | State | Sex | Year | Name  | Count |
|-------|-------|-----|------|-------|-------|
| 7512  | CA    | F   | 1925 | Bella | 8     |
| 12368 | CA    | F   | 1932 | Lisa  | 5     |
| 14741 | CA    | F   | 1936 | Lisa  | 8     |
| 17084 | CA    | F   | 1939 | Lisa  | 5     |

Fortunately, `pandas` provides many alternative methods for constructing boolean filters.

The `.isin` function is one such example. This method evaluates if the values in a `Series` are contained in a different sequence (list, array, or `Series`) of values. In the cell below, we achieve equivalent results to the `DataFrame` above with far more concise code.

```
names = ["Bella", "Alex", "Narges", "Lisa"]
babynames["Name"].isin(names).head()
```

```
0    False
1    False
2    False
3    False
4    False
Name: Name, dtype: bool
```

```
babynames[babynames["Name"].isin(names)].head()
```

|       | State | Sex | Year | Name  | Count |
|-------|-------|-----|------|-------|-------|
| 6289  | CA    | F   | 1923 | Bella | 5     |
| 7512  | CA    | F   | 1925 | Bella | 8     |
| 12368 | CA    | F   | 1932 | Lisa  | 5     |
| 14741 | CA    | F   | 1936 | Lisa  | 8     |
| 17084 | CA    | F   | 1939 | Lisa  | 5     |

The function `str.startswith` can be used to define a filter based on string values in a `Series` object. It checks to see if string values in a `Series` start with a particular character.

```
# Identify whether names begin with the letter "N"
babynames["Name"].str.startswith("N").head()
```

```
0    False
1    False
2    False
3    False
4    False
Name: Name, dtype: bool
```

```
# Extracting names that begin with the letter "N"
babynames[babynames["Name"].str.startswith("N")].head()
```

|     | State | Sex | Year | Name   | Count |
|-----|-------|-----|------|--------|-------|
| 76  | CA    | F   | 1910 | Norma  | 23    |
| 83  | CA    | F   | 1910 | Nellie | 20    |
| 127 | CA    | F   | 1910 | Nina   | 11    |
| 198 | CA    | F   | 1910 | Nora   | 6     |
| 310 | CA    | F   | 1911 | Nellie | 23    |

## 3.2 Adding, Removing, and Modifying Columns

In many data science tasks, we may need to change the columns contained in our `DataFrame` in some way. Fortunately, the syntax to do so is fairly straightforward.

To add a new column to a `DataFrame`, we use a syntax similar to that used when accessing an existing column. Specify the name of the new column by writing `df["column"]`, then assign this to a `Series` or array containing the values that will populate this column.

```
# Create a Series of the length of each name.
babynames["name_lengths"] = babynames["Name"].str.len()

# Add a column named "name_lengths" that includes the length of each name
babynames["name_lengths"] = babynames["name_lengths"]
babynames.head(5)
```

|   | State | Sex | Year | Name     | Count | name_lengths |
|---|-------|-----|------|----------|-------|--------------|
| 0 | CA    | F   | 1910 | Mary     | 295   | 4            |
| 1 | CA    | F   | 1910 | Helen    | 239   | 5            |
| 2 | CA    | F   | 1910 | Dorothy  | 220   | 7            |
| 3 | CA    | F   | 1910 | Margaret | 163   | 8            |
| 4 | CA    | F   | 1910 | Frances  | 134   | 7            |

If we need to later modify an existing column, we can do so by referencing this column again with the syntax `df["column"]`, then re-assigning it to a new `Series` or array of the appropriate length.

```
# Modify the "name_lengths" column to be one less than its original value
babynames["name_lengths"] = babynames["name_lengths"] - 1
babynames.head()
```

|   | State | Sex | Year | Name     | Count | name_lengths |
|---|-------|-----|------|----------|-------|--------------|
| 0 | CA    | F   | 1910 | Mary     | 295   | 3            |
| 1 | CA    | F   | 1910 | Helen    | 239   | 4            |
| 2 | CA    | F   | 1910 | Dorothy  | 220   | 6            |
| 3 | CA    | F   | 1910 | Margaret | 163   | 7            |
| 4 | CA    | F   | 1910 | Frances  | 134   | 6            |

We can rename a column using the `.rename()` method. It takes in a dictionary that maps old column names to their new ones.

```
# Rename "name_lengths" to "Length"
babynames = babynames.rename(columns={"name_lengths": "Length"})
babynames.head()
```

|   | State | Sex | Year | Name     | Count | Length |
|---|-------|-----|------|----------|-------|--------|
| 0 | CA    | F   | 1910 | Mary     | 295   | 3      |
| 1 | CA    | F   | 1910 | Helen    | 239   | 4      |
| 2 | CA    | F   | 1910 | Dorothy  | 220   | 6      |
| 3 | CA    | F   | 1910 | Margaret | 163   | 7      |
| 4 | CA    | F   | 1910 | Frances  | 134   | 6      |

If we want to remove a column or row of a `DataFrame`, we can call the `.drop` ([documentation](#)) method. Use the `axis` parameter to specify whether a column or row should be dropped. Unless otherwise specified, `pandas` will assume that we are dropping a row by default.

```
# Drop our new "Length" column from the DataFrame
babynames = babynames.drop("Length", axis="columns")
babynames.head(5)
```

|   | State | Sex | Year | Name     | Count |
|---|-------|-----|------|----------|-------|
| 0 | CA    | F   | 1910 | Mary     | 295   |
| 1 | CA    | F   | 1910 | Helen    | 239   |
| 2 | CA    | F   | 1910 | Dorothy  | 220   |
| 3 | CA    | F   | 1910 | Margaret | 163   |
| 4 | CA    | F   | 1910 | Frances  | 134   |

Notice that we *re-assigned* `babynames` to the result of `babynames.drop(...)`. This is a subtle but important point: `pandas` table operations **do not occur in-place**. Calling `df.drop(...)` will output a *copy* of `df` with the row/column of interest removed without modifying the original `df` table.

In other words, if we simply call:

```
# This creates a copy of `babynames` and removes the column "Name"...
babynames.drop("Name", axis="columns")

# ...but the original `babynames` is unchanged!
# Notice that the "Name" column is still present
babynames.head(5)
```

|   | State | Sex | Year | Name     | Count |
|---|-------|-----|------|----------|-------|
| 0 | CA    | F   | 1910 | Mary     | 295   |
| 1 | CA    | F   | 1910 | Helen    | 239   |
| 2 | CA    | F   | 1910 | Dorothy  | 220   |
| 3 | CA    | F   | 1910 | Margaret | 163   |
| 4 | CA    | F   | 1910 | Frances  | 134   |

### 3.3 Useful Utility Functions

`pandas` contains an extensive library of functions that can help shorten the process of setting and getting information from its data structures. In the following section, we will give overviews of each of the main utility functions that will help us in Data 100.

Discussing all functionality offered by `pandas` could take an entire semester! We will walk you through the most commonly-used functions and encourage you to explore and experiment on your own.

- `NumPy` and built-in function support
- `.shape`
- `.size`
- `.describe()`
- `.sample()`
- `.value_counts()`
- `.unique()`
- `.sort_values()`

The [pandas documentation](#) will be a valuable resource in Data 100 and beyond.

#### 3.3.1 NumPy

`pandas` is designed to work well with `NumPy`, the framework for array computations you encountered in [Data 8](#). Just about any `NumPy` function can be applied to `pandas DataFrame`s and `Series`.

```
# Pull out the number of babies named Yash each year
yash_count = babynames[babynames["Name"] == "Yash"]["Count"]
yash_count.head()
```



```
331824      8
334114      9
336390     11
338773     12
341387     10
Name: Count, dtype: int64
```

```
# Average number of babies named Yash each year
np.mean(yash_count)
```

```
17.142857142857142
```

```
# Max number of babies named Yash born in any one year
np.max(yash_count)
```

```
29
```

### 3.3.2 `.shape` and `.size`

`.shape` and `.size` are attributes of `Series` and `DataFrame`s that measure the “amount” of data stored in the structure. Calling `.shape` returns a tuple containing the number of rows and columns present in the `DataFrame` or `Series`. `.size` is used to find the total number of elements in a structure, equivalent to the number of rows times the number of columns.

Many functions strictly require the dimensions of the arguments along certain axes to match. Calling these dimension-finding functions is much faster than counting all of the items by hand.

```
# Return the shape of the DataFrame, in the format (num_rows, num_columns)
babynames.shape
```

```
(407428, 5)
```

```
# Return the size of the DataFrame, equal to num_rows * num_columns
babynames.size
```

```
2037140
```

### 3.3.3 `.describe()`

If many statistics are required from a `DataFrame` (minimum value, maximum value, mean value, etc.), then `.describe()` ([documentation](#)) can be used to compute all of them at once.

```
babynames.describe()
```

|       | Year          | Count         |
|-------|---------------|---------------|
| count | 407428.000000 | 407428.000000 |
| mean  | 1985.733609   | 79.543456     |
| std   | 27.007660     | 293.698654    |
| min   | 1910.000000   | 5.000000      |
| 25%   | 1969.000000   | 7.000000      |
| 50%   | 1992.000000   | 13.000000     |
| 75%   | 2008.000000   | 38.000000     |
| max   | 2022.000000   | 8260.000000   |

A different set of statistics will be reported if `.describe()` is called on a [Series](#).

```
babynames["Sex"].describe()
```

```
count    407428
unique      2
top        F
freq    239537
Name: Sex, dtype: object
```

### 3.3.4 `.sample()`

As we will see later in the semester, random processes are at the heart of many data science techniques (for example, train-test splits, bootstrapping, and cross-validation). `.sample()` ([documentation](#)) lets us quickly select random entries (a row if called from a [DataFrame](#), or a value if called from a [Series](#)).

By default, `.sample()` selects entries *without* replacement. Pass in the argument `replace=True` to sample with replacement.

```
# Sample a single row
babynames.sample()
```

|        | State | Sex | Year | Name      | Count |
|--------|-------|-----|------|-----------|-------|
| 195242 | CA    | F   | 2011 | Genavieve | 15    |

Naturally, this can be chained with other methods and operators (`iloc`, etc.).

```
# Sample 5 random rows, and select all columns after column 2
babynames.sample(5).iloc[:, 2:]
```

|       | Year | Name      | Count |
|-------|------|-----------|-------|
| 61407 | 1970 | Rosalinda | 47    |

|        | Year | Name     | Count |
|--------|------|----------|-------|
| 93738  | 1983 | Katerina | 5     |
| 49544  | 1964 | Becky    | 248   |
| 378105 | 2012 | Timmy    | 6     |
| 19625  | 1942 | Shelia   | 5     |

```
# Randomly sample 4 names from the year 2000, with replacement, and select all columns of
babynames[babynames["Year"] == 2000].sample(4, replace = True).iloc[:, 2:]
```

|        | Year | Name   | Count |
|--------|------|--------|-------|
| 343780 | 2000 | Ted    | 11    |
| 344678 | 2000 | Cris   | 5     |
| 149393 | 2000 | Carina | 128   |
| 344379 | 2000 | Arsen  | 6     |

### 3.3.5 .value\_counts()

The `Series.value_counts()` ([documentation](#)) method counts the number of occurrence of each unique value in a `Series`. In other words, it *counts* the number of times each unique *value* appears. This is often useful for determining the most or least common entries in a `Series`.

In the example below, we can determine the name with the most years in which at least one person has taken that name by counting the number of times each name appears in the `"Name"` column of `babynames`. Note that the return value is also a `Series`.

```
babynames["Name"].value_counts().head()
```

```
Name
Jean      223
Francis   221
Guadalupe 218
Jessie    217
Marion    214
Name: count, dtype: int64
```

### 3.3.6 .unique()

If we have a `Series` with many repeated values, then `.unique()` ([documentation](#)) can be used to identify only the *unique* values. Here we return an array of all the names in `babynames`.

```
babynames["Name"].unique()
```

```
array(['Mary', 'Helen', 'Dorothy', ..., 'Zae', 'Zai', 'Zayvier'],  
      dtype=object)
```

### 3.3.7 .sort\_values()

Ordering a `DataFrame` can be useful for isolating extreme values. For example, the first 5 entries of a row sorted in descending order (that is, from highest to lowest) are the largest 5 values. `.sort_values` ([documentation](#)) allows us to order a `DataFrame` or `Series` by a specified column. We can choose to either receive the rows in `ascending` order (default) or `descending` order.

```
# Sort the "Count" column from highest to lowest  
babynames.sort_values(by="Count", ascending=False).head()
```

|        | State | Sex | Year | Name    | Count |
|--------|-------|-----|------|---------|-------|
| 268041 | CA    | M   | 1957 | Michael | 8260  |
| 267017 | CA    | M   | 1956 | Michael | 8258  |
| 317387 | CA    | M   | 1990 | Michael | 8246  |
| 281850 | CA    | M   | 1969 | Michael | 8245  |
| 283146 | CA    | M   | 1970 | Michael | 8196  |

Unlike when calling `.value_counts()` on a `DataFrame`, we do not need to explicitly specify the column used for sorting when calling `.value_counts()` on a `Series`. We can still specify the ordering paradigm – that is, whether values are sorted in ascending or descending order.

```
# Sort the "Name" Series alphabetically  
babynames["Name"].sort_values(ascending=True).head()
```

```
366001    Aadan  
384005    Aadan  
369120    Aadan  
398211    Aadarsh  
370306    Aaden  
Name: Name, dtype: object
```

## 3.4 Custom Sorts

Now, let's try to solve a sorting problem using different approaches. Assume we want to find the longest baby names and sort our data accordingly.

We'll start by loading the `babynames` dataset. Note that this dataset is filtered to only contain data from California.

► Code

|        | State | Sex | Year | Name    | Count |
|--------|-------|-----|------|---------|-------|
| 407418 | CA    | M   | 2022 | Zach    | 5     |
| 407419 | CA    | M   | 2022 | Zadkiel | 5     |
| 407420 | CA    | M   | 2022 | Zae     | 5     |
| 407421 | CA    | M   | 2022 | Zai     | 5     |
| 407422 | CA    | M   | 2022 | Zay     | 5     |
| 407423 | CA    | M   | 2022 | Zayvier | 5     |
| 407424 | CA    | M   | 2022 | Zia     | 5     |
| 407425 | CA    | M   | 2022 | Zora    | 5     |
| 407426 | CA    | M   | 2022 | Zuriel  | 5     |
| 407427 | CA    | M   | 2022 | Zylo    | 5     |

### 3.4.1 Approach 1: Create a Temporary Column

One method to do this is to first start by creating a column that contains the lengths of the names.

```
# Create a Series of the length of each name
babynames["name_lengths"] = babynames["Name"].str.len()

# Add a column named "name_lengths" that includes the length of each name
babynames["name_lengths"] = babynames["name_lengths"]
babynames.head(5)
```

|   | State | Sex | Year | Name     | Count | name_lengths |
|---|-------|-----|------|----------|-------|--------------|
| 0 | CA    | F   | 1910 | Mary     | 295   | 4            |
| 1 | CA    | F   | 1910 | Helen    | 239   | 5            |
| 2 | CA    | F   | 1910 | Dorothy  | 220   | 7            |
| 3 | CA    | F   | 1910 | Margaret | 163   | 8            |
| 4 | CA    | F   | 1910 | Frances  | 134   | 7            |

We can then sort the `DataFrame` by that column using `.sort_values()`:

```
# Sort by the temporary column
babynames = babynames.sort_values(by="name_lengths", ascending=False)
babynames.head(5)
```

|        | State | Sex | Year | Name            | Count | name_lengths |
|--------|-------|-----|------|-----------------|-------|--------------|
| 334166 | CA    | M   | 1996 | Franciscojavier | 8     | 15           |
| 337301 | CA    | M   | 1997 | Franciscojavier | 5     | 15           |
| 339472 | CA    | M   | 1998 | Franciscojavier | 6     | 15           |

|        | State | Sex | Year | Name            | Count | name_lengths |
|--------|-------|-----|------|-----------------|-------|--------------|
| 321792 | CA    | M   | 1991 | Ryanchristopher | 7     | 15           |
| 327358 | CA    | M   | 1993 | Johnchristopher | 5     | 15           |

Finally, we can drop the `name_length` column from `babynames` to prevent our table from getting cluttered.

```
# Drop the 'name_length' column
babynames = babynames.drop("name_lengths", axis='columns')
babynames.head(5)
```

|        | State | Sex | Year | Name            | Count |
|--------|-------|-----|------|-----------------|-------|
| 334166 | CA    | M   | 1996 | Franciscojavier | 8     |
| 337301 | CA    | M   | 1997 | Franciscojavier | 5     |
| 339472 | CA    | M   | 1998 | Franciscojavier | 6     |
| 321792 | CA    | M   | 1991 | Ryanchristopher | 7     |
| 327358 | CA    | M   | 1993 | Johnchristopher | 5     |

### 3.4.2 Approach 2: Sorting using the `key` Argument

Another way to approach this is to use the `key` argument of `.sort_values()`. Here we can specify that we want to sort `"Name"` values by their length.

```
babynames.sort_values("Name", key=lambda x: x.str.len(), ascending=False).head()
```

|        | State | Sex | Year | Name            | Count |
|--------|-------|-----|------|-----------------|-------|
| 334166 | CA    | M   | 1996 | Franciscojavier | 8     |
| 327472 | CA    | M   | 1993 | Ryanchristopher | 5     |
| 337301 | CA    | M   | 1997 | Franciscojavier | 5     |
| 337477 | CA    | M   | 1997 | Ryanchristopher | 5     |
| 312543 | CA    | M   | 1987 | Franciscojavier | 5     |

### 3.4.3 Approach 3: Sorting using the `map` Function

We can also use the `map` function on a `Series` to solve this. Say we want to sort the `babynames` table by the number of `"dr"` 's and `"ea"` 's in each `"Name"`. We'll define the function `dr_ea_count` to help us out.

```
# First, define a function to count the number of times "dr" or "ea" appear in each name
def dr_ea_count(string):
    return string.count('dr') + string.count('ea')

# Then, use `map` to apply `dr_ea_count` to each name in the "Name" column
babynames["dr_ea_count"] = babynames["Name"].map(dr_ea_count)
```

```
# Sort the DataFrame by the new "dr_ea_count" column so we can see our handiwork
babynames = babynames.sort_values(by="dr_ea_count", ascending=False)
babynames.head()
```

|        | State | Sex | Year | Name     | Count | dr_ea_count |
|--------|-------|-----|------|----------|-------|-------------|
| 115957 | CA    | F   | 1990 | Deandrea | 5     | 3           |
| 101976 | CA    | F   | 1986 | Deandrea | 6     | 3           |
| 131029 | CA    | F   | 1994 | Leandrea | 5     | 3           |
| 108731 | CA    | F   | 1988 | Deandrea | 5     | 3           |
| 308131 | CA    | M   | 1985 | Deandrea | 6     | 3           |

We can drop the `dr_ea_count` once we're done using it to maintain a neat table.

```
# Drop the `dr_ea_count` column
babynames = babynames.drop("dr_ea_count", axis = 'columns')
babynames.head(5)
```

|        | State | Sex | Year | Name     | Count |
|--------|-------|-----|------|----------|-------|
| 115957 | CA    | F   | 1990 | Deandrea | 5     |
| 101976 | CA    | F   | 1986 | Deandrea | 6     |
| 131029 | CA    | F   | 1994 | Leandrea | 5     |
| 108731 | CA    | F   | 1988 | Deandrea | 5     |
| 308131 | CA    | M   | 1985 | Deandrea | 6     |

## 3.5 Parting Note

Manipulating `DataFrames` is not a skill that is mastered in just one day. Due to the flexibility of `pandas`, there are many different ways to get from point A to point B. We recommend trying multiple different ways to solve the same problem to gain even more practice and reach that point of mastery sooner.

Next, we will start digging deeper into the mechanics behind grouping data.