Everything is Better with Friends

Using SAS in Python Applications with SASPy and Open-Source Tooling (Beyond the Basics)

→ Setup for Part 2

Getting setup to use Google Colab

- 1. To execute code cells, you'll need credentials for the following:
 - Google. (If you're not already signed in, you should see a Sign In button in the upper right corner. You can also visit
 https://accounts.google.com/signup to create an account for free.)
- 2. We recommend enabling line numbers using the Tools menu: Tools -> Settings -> Editor -> Show line numbers -> Save
- 3. We also recommend enabling the Table of Contents using the View menu: View -> Table of contents
- 4. To save a copy of this notebook, along with any edits you make, please use the File menu: File -> Save a copy in Drive
- 5. Some useful Zoom Reactions:
 - In the section of the s
 - 🖖 (Raise Hand) when you need tech support
 - I'm Away) to let us know you've stepped away
- 6. Looking for "extra credit"? Please let us know if you spot any typos!

▼ Install and import packages

```
# Install the rich module for colorful printing, limiting the version for compatibility with Colab
    !pip install 'rich<13.3'
    # We'll use IPython to display DataFrames or HTML content
 5
    from IPython.display import display, HTML
    # We'll use the pandas package to create and manipulate DataFrame objects
 8
    import pandas
 9
10 # We'll use the requests package to call a web API
11
    import requests
12
    # We're overwriting the default print function with rich.print
13
    from rich import print
14
15
16
    # We're also setting the maximum line width of rich.print to be a bit wider (to avoid line wrapping)
17
    from rich import get console
    console = get console()
    console.width = 165
```

- ▼ Part 2. Rectangularizing unstructured data in Python applications
- ▼ Section 2.1. Create pharm_class_response

```
1 # Let's explore one of the many endpoints of the openFDA API!
 3 # This might not be the most exciting ways to get data, but it's one of the few public API endpoints
 4 # that doesn't require an access token, and it'll give us a chance to practice working with JSON.
 6 # Use an open web API to get the number of drugs available by pharmacologic class.
 7 # Note: By default, only the first 100 results are provided, sorted in descending order by count.
 8 # To retrieve more than the first 100 results, a combination of limit and skip parameters can be
 9 # used, as described at https://open.fda.gov/apis/paging/
10 pharm class response = requests.get('https://api.fda.gov/drug/ndc.json?count=pharm class.exact')
11
12 # Check the resulting status code to make sure the API call was successful, with 200 = "OK".
13 http status = pharm class response.status code
14 http status info = f'https://httpstatuses.com/{http status}'
15 if http status == 200:
       print('API call successful!\n')
17 print(f'See {http status info} for more information about HTTP status code {http status}.')
    API call successful!
    See <a href="https://httpstatuses.com/200">https://httpstatuses.com/200</a> for more information about HTTP status code 200.
```

- 1. True or False: Changing Line 15 to a single-equals (=) would have the same effect.
- 2. True or False: Removing the indentation on Line 16 would have the same effect.
- Fun Fact: The FDA provides many open APIs. Examples queries can be found at https://open.fda.gov/apis/drug/ndc/example-api-queries/

Solution: False! Single-equals (=) is only used for variable assignment, and double-equals (=) is only used to test for equality.

▼ Section 2.2. Explore pharm_class_response

```
1 # What exactly makes up an HTTP response?
2 print(dir(pharm_class_response))
3 print('\n')
4
5 # Let's take a look at the HTTP headers used behind the scenes
6 print(dict(pharm_class_response.headers))
```

[. .. .

Concept Check 2.2

- Try this, and see what happens: Find the type of dir(pharm_class_response) by using the type function, e.g., using print(type(dir(pharm class response)))
- Try this, and see what happens: Find the type of dict(pharm_class_response.headers) by using the type function, e.g., using print(type(dict(pharm_class_response.headers)))
- Fun Facts:
 - Lists are a fundamental Python data structure and are similar to SAS DATA step arrays.
 - o Dictionaries are another fundamental Python data structure and are related to SAS formats and DATA step hash tables.
 - In the requests module, the headers attribute is a special dictionary with type <class 'requests.structures.CaseInsensitiveDict'>, so the dict function is used to convert it to a regular dictionary.
 - The dir and type functions are examples of Python's introspection features, which allow us to ask an object to tell us about itself.
 Other introspection functions include help (to get an object's built-in documentation) and id (to get an object's behind-the-scenes unique id).

▼ Section 2.3. Create pharm_class_json

```
'connection'

1 # Extract and print the JSON-formatted list of counts of drugs by pharmacologic class.

2 pharm_class_json = pharm_class_response.json()

3 print(pharm_class_json)
```

```
'meta': {
          'disclaimer': 'Do not rely on openFDA to make decisions regarding medical care. While we make eve
all results are unvalidated. We may limit or otherwise restrict your access to the API in line with our T
          'terms': 'https://open.fda.gov/terms/',
          'license': 'https://open.fda.gov/license/',
          'last updated': '2023-02-27'
     'results': [
          {'term': 'Cell-mediated Immunity [PE]', 'count': 6032},
          {'term': 'Increased Histamine Release [PE]', 'count': 6018},
          {'term': 'Allergens [CS]', 'count': 6011},
          {'term': 'Increased IgG Production [PE]', 'count': 3840},
          {'term': 'Anti-Inflammatory Agents', 'count': 3721},
          {'term': 'Cyclooxygenase Inhibitors [MoA]', 'count': 3721},
          {'term': 'Non-Steroidal [CS]', 'count': 3721},
          {'term': 'Nonsteroidal Anti-inflammatory Drug [EPC]', 'count': 3721}, {'term': 'Histamine H1 Receptor Antagonists [MoA]', 'count': 2397},
          {'term': 'Corticosteroid Hormone Receptor Agonists [MoA]', 'count': 2383},
          {'term': 'Corticosteroid [FPC]', 'count': 2383},
{'term': 'Histamine-1 Receptor Antagonist [EPC]', 'count': 2314},
          {'term': 'Osmotic Activity [MoA]', 'count': 2221},
{'term': 'Increased Large Intestinal Motility [PE]', 'count': 2185},
         {'term': 'Osmotic Laxative [EPC]', 'count': 2143},
{'term': 'Sigma-1 Agonist [EPC]', 'count': 2063},
{'term': 'Sigma-1 Receptor Agonists [MoA]', 'count': 2063},
          {'term': 'Uncompetitive N-methyl-D-aspartate Receptor Antagonist [EPC]', 'count': 2063},
          {'term': 'Uncompetitive NMDA Receptor Antagonists [MoA]', 'count': 2063},
          {'term': 'Inhibition Large Intestine Fluid/Electrolyte Absorption [PE]', 'count': 2040},
          {'term': 'Decreased Central Nervous System Disorganized Electrical Activity [PE]', 'count': 1958}
         {'term': 'Adrenergic alphal-Agonists [MoA]', 'count': 1879},
{'term': 'alpha-1 Adrenergic Agonist [EPC]', 'count': 1879},
{'term': 'Opioid Agonist [EPC]', 'count': 1649},
{'term': 'Dietary Proteins [CS]', 'count': 1607},
          {'term': 'Non-Standardized Food Allergenic Extract [EPC]', 'count': 1602},
          {'term': 'Plant Proteins [CS]', 'count': 1598},
          {'term': 'Pollen [CS]', 'count': 1583},
          {'term': 'Antiarrhythmic [EPC]', 'count': 1551},
          {'term': 'Non-Standardized Plant Allergenic Extract [EPC]', 'count': 1519},
```

- Short Answer: What types of standard Python objects appear in the output of pharm class json?
- Fun Fact: In Python, it's common to work with deeply nested objects (like a Russian nested doll, or a Turducken).

```
{'term': 'Atypical Antipsychotic [EPC]', 'count': 1341},
```

Solution: We see instances of int, str, dict, and list.

```
Clarents (Administration Association Appendicts (Man))
```

▼ Section 2.4. Create pharm_class_list

```
Date of API results: 2023-02-27

[
    ['Cell-mediated Immunity [PE]', 6032],
    ['Increased Histamine Release [PE]', 6018],
    ['Allergens [CS]', 6011],
    ['Increased IgG Production [PE]', 3840],
    ['Anti-Inflammatory Agents', 3721],
    ['Cyclopygapase Inhibitors [MoAl', 3721]
```

- 1. True or False: Changing Line 8 to pharm_class_json['RESULTS'] (i.e., changing the dictionary key to all caps) would have the same effect.
- 2. Short Answer: What types of standard Python objects appear in the definition of pharm class list?
- Fun Fact: Instead of bothering with a list of lists, we could have instead built a DataFrame row-by-row inside the for-loop. However, DataFrame operations inside a for-loop tend to be slow.

```
['Uncompetitive NMDA Receptor Antagonists [MOA]', 2063],
```

Solution: False! In general, dictionary keys are case-sensitive in Python, just like variable names.

```
[Jolobo 1 Adronoraic Agonict [EDC]] 10701
```

▼ Section 2.5. Create pharm_class_df

```
I'Plant Proteins [CS]'. 15981.

1 # Now that we've finish looping, we can put the definitions in a DataFrame called pharm_class_df.

2 pharm_class_df = pandas.DataFrame(pharm_class_list, columns = ['term', 'count'])

3

4 # We can also inspect the size of pharm_class_df.

5 print(f'The size of pharm_class_df: {pharm_class_df.shape}')

6 print('\n')

7

8 # In addition, we can get a sense of the average size pharmacologic class.

9 print(f"The median size pharmacologic class in pharm_class_df: {pharm_class_df['count'].median()}")

10 print('\n')

11

12 # Finally, we can display pharm_class_df.

13 print(f'The contents of pharm_class_df:')

14 display(pharm class_df)
```

```
The size of pharm_class_df: (100, 2)

The median size pharmacologic class in pharm_class_df: 1079.5

The contents of pharm_class_df:

term count

Cell-mediated Immunity [PE] 6032

1 Increased Histamine Release [PE] 6018
```

- Short Answer: Other than the median, what are some descriptive statistics we might consider using to better understand the contents of pharm class df?
- Fun Fact: JSON-formatted data is useful because of how flexibly information can be nested. However, to actually work with the information inside, it's common to first rectangularize the JSON object.

Solution: The following section of the "Getting Started" guide for pandas gives a good overview: https://pandas.pydata.org/docs/getting_started/intro_tutorials/06_calculate_statistics.html

```
JU Hoyolio Altituepressalit [Li O] 000
```

▼ Section 2.6. Additional Exercises

For practice, we recommend the following:

- Run the code cell below.
- Repeat the steps in Sections 2.4-5 with the following two changes:
 - Form a DataFrame whose first column is count.
 - Calculate a statistic other than median.

```
1 # Let's try a different openFDA endpoint.
2 generic_name_response = requests.get('https://api.fda.gov/drug/ndc.json?count=generic_name.exact')
3
4 # Check the resulting status code to make sure the API call was successful, with 200 = "OK".
5 if generic_name_response.status_code == 200:
6     print('API call successful!\n')
7
8 # Finally, let's extract and print the JSON-formatted return value.
9 generic_name_json = generic_name_response.json()
10 print('Here\'s the resulting data structure:')
11 print(generic name json)
```

```
API call successful!
      Here's the resulting data structure:
             'meta': {
                   'disclaimer': 'Do not rely on openFDA to make decisions regarding medical care. While we make every effort to ensure that data is accurate, yo
      all results are unvalidated. We may limit or otherwise restrict your access to the API in line with our Terms of Service.',
                  'terms': 'https://open.fda.gov/terms/',
                  'license': 'https://open.fda.gov/license/',
                  'last_updated': '2023-02-27'
           },
'results': [
                 {'term': 'ALCOHOL', 'count': 1756},
{'term': 'Alcohol', 'count': 1058},
{'term': 'Ibuprofen', 'count': 940},
                  {'term': 'Acetaminophen', 'count': 921},
                 {'term': 'Zinc Oxide', 'count': 783},
{'term': 'Ethyl Alcohol', 'count': 713},
                 {'term': 'Benzalkonium Chloride', 'count': 636},
{'term': 'BENZALKONIUM CHLORIDE', 'count': 584},
{'term': 'Menthol', 'count': 582},
                 {'term': 'Sodium Fluoride', 'count': 511},
{'term': 'Salicylic Acid', 'count': 464},
{'term': 'Isopropyl Alcohol', 'count': 428},
                 {'term': 'Oxygen', 'count': 406},
{'term': 'ZINC OXIDE', 'count': 391},
{'term': 'Benzocaine', 'count': 381},
                  {'term': 'Nicotine Polacrilex', 'count': 381},
                  {'term': 'Avobenzone, Homosalate, Octisalate, Octocrylene', 'count': 372},
                  {'term': 'Gabapentin', 'count': 355},
                 {'term': 'Aspirin', 'count': 336}, {'term': 'MENTHOL', 'count': 320},
                  {'term': 'SALICYLIC ACID', 'count': 309},
                  {'term': 'Diphenhydramine HCl', 'count': 302},
                  {'term': 'Titanium Dioxide', 'count': 302},
                  {'term': 'Titanium Dioxide, Zinc Oxide', 'count': 274},
                  {'term': 'Lisinopril', 'count': 268},
                 {'term': 'Hydrocortisone', 'count': 263}, {'term': 'Naproxen Sodium', 'count': 260},
                  {'term': 'Loratadine', 'count': 258}, {'term': 'Famotidine', 'count': 257},
                 {'term': 'TITANIUM DIOXIDE', 'count': 257},
{'term': 'TITANIUM DIOXIDE', 'count': 256},
{'term': 'ETHYL ALCOHOL', 'count': 256},
{'term': 'Levothyroxine Sodium', 'count': 249},
{'term': 'Simethicone', 'count': 243},
{'term': 'Aripiprazole', 'count': 242},
{'term': 'Levetiracetam', 'count': 234},
{'term': 'Bearalkopium chloride', 'count': 234},
                 {'term': 'Benzalkonium chloride', 'count': 233},
{'term': 'Guaifenesin', 'count': 233},
                  {'term': 'OCTINOXATE and TITANIUM DIOXIDE', 'count': 231},
 1 # Accumulate generic names and counts in a list of lists called generic name list.
 2 generic name list = []
 3 for generic name count in generic name json['results']:
          generic name list.append(
                     generic name count['count'],
                     generic name count['term'],
11 # In case we want to track when these API results were obtained, let's also extract the date.
12 generic name date = generic name json['meta']['last updated']
14 # Now let's print the date.
15 print(f'Date of API results: {generic name date}')
```

5 6

7

8 9 1.0

13

```
16 print('\n')
17
18 # And then let's print generic_name_list.
19 print(generic_name_list)
20
21 # Now that we've finish looping, we can put the definitions in a DataFrame called generic_name_df.
22 generic_name_df = pandas.DataFrame(generic_name_list, columns = ['count','term'])
24 # We can also inspect the size of generic_name_df.
25 print(f'The size of generic_name_df: {generic_name_df.shape}')
26 print('\n')
28 # In addition, we can get a sense of the average size generic type.
29 print(f"The mean generics count in generic_name_df: {generic_name_df['count'].mean()}")
30 print('\n')
32 # Finally, we can display generic name df.
33 print(f'The contents of generic_name_df:')
34 display(generic_name_df)
```

```
Date of API results: 2023-02-27

[
    [1756, 'ALCOHOL'],
    [1058, 'ALcohol'],
    [940, 'Ibuprofen'],
    [921, 'Acetaminophen'],
    [783, 'Zinc Oxide'],
    [713, 'Ethyl Alcohol'],
    [636, 'Benzalkonium Chloride'],
    [584, 'BENZALKONIUM CHLORIDE'],
    [582, 'Menthol'],
    [511, 'Sodium Fluoride'],
    [464, 'Salicylic Acid'],
    [428, 'Isopropyl Alcohol'],
    [406, 'Oxygen'],
```

Notes and Resources

```
[355 'Gahanentin']
```

Want some ideas for what to do next? Here are our suggestions:

- 1. For more about the pandas package, including the methods used above, see the following:
 - https://pandas.pydata.org/docs/reference/api/pandas.DataFrame.shape.html
 - https://pandas.pydata.org/docs/reference/api/pandas.Series.median.html
 - https://pandas.pydata.org/docs/reference/api/pandas.Series.value_counts.html
- 2. For more about the requests package, see https://docs.python-requests.org/
- 3. For more about the rich package, see https://rich.readthedocs.io/
- 4. For more about some of the Python features used, such as dictionaries, lists, and control flow with if-then-else conditionals and for-loops, we recommend the following chapters of <u>A Whirlwind Tour of Python</u>:
 - https://jakevdp.github.io/WhirlwindTourOfPython/06-built-in-data-structures.html
 - https://jakevdp.github.io/WhirlwindTourOfPython/07-control-flow-statements.html
- 5. For more information on f-strings (i.e., Python strings like f'https://httpstatuses.com/{http_status}'), see https://realpython.com/python-f-strings/.
- 6. For background on the HTTP Request/Response Cycle, we recommend the following:
 - Brief Overview: https://backend.turing.edu/module2/lessons/how_the_web_works_http
 - Deeper Overview: https://developer.mozilla.org/en-US/docs/Web/HTTP/Overview
 - Summary of HTTP Status Codes: https://httpstatuses.com/
 - Google's Implementation of HTTP Status Code 418: https://www.google.com/teapot
- 7. For more practice with open web APIs, we recommend looking through <a href="https://github.com/public-apis/publ
- 8. For more about the complexity of parsing JSON in SAS, see https://blogs.sas.com/content/sasdummy/2016/12/02/json-libname-engine-sas/
- 9. We welcome follow-up conversations. You can connect with us on LinkedIn or email us at isaiah.lankham@gmail.com and matthew.t.slaughter@gmail.com

10. If you have a GitHub account (or don't mind creating one), you can also chat with us on Gitter at https://gitter.im/saspy-bffs/community

```
[155, 'Quetiapine Fumarate'],
[155, 'Titanium Dioxide and Zinc Oxide'],
[154, 'Carvedilol'],
[154, 'Clotrimazole'],
[154, 'Minoxidil'],
[153, 'Dimethicone'],
[153, 'Lidocaine Hydrochloride'],
[152, 'Diclofenac Sodium'],
[152, 'Rosuvastatin Calcium'],
[146, 'Diltiazem Hydrochloride'],
[144, 'DiPHENHYDRAMINE HYDROCHLORIDE'],
[144, 'Naproxen'],
[144, 'ACETAMINOPHEN'],
[136, 'Hydrochlorothiazide'],
[136, 'nicotine polacrilex'],
[134, 'Amoxicillin and Clavulanate Potassium'],
[134, 'Glycopyrrolate'],
[133, 'Buspirone Hydrochloride'],
[132, 'Gclipizide'],
[132, 'Glipizide'],
[131, 'LIDOCAINE HYDROCHLORIDE'],
[129, 'CHLOROXYLENOL'],
[129, 'CHLOROXYLENOL'],
[129, 'Cephalexin'],
[128, 'ATORVASTATIN CALCIUM'],
[128, 'Labetalol Hydrochloride'],
[127, 'Aluminum Zirconium Tetrachlorohydrex GLY'],
[127, 'Doxycycline Hyclate'],
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Colab paid products - Cancel contracts here
        [127, 'Doxycycline Hyclate'],
```