How to Clean Messy Data in Python



DS 6001: Practice and Applications of Data Science

Online Communities
Using Python's built-in help documentation
Good old Google
Stack Overflow
Interacting with other Python users on PySlackers
Live chats with Python users on Freenode
Python Mailing lists

Loading CSV and ASCII Data into Python

Electronic data files
Changing the working directory
Loading standard CSV files
Looking at the data to see if it loaded correctly
Loading messy CSV and other ASCII files
Writing CSV and ASCII files

Loading Other Kinds of Electronic Data Files

Loading fixed width files
Loading Excel files
Loading SAS, Stata, and SPSS files
Working with JSON files
What is JSON?
Who uses JSON?
Loading JSON data into Python
Converting DataFrames/CSVs to JSON
Writing JSON files to disk

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- Defining and working with objects
- Saving scripts/notebooks
- Using functions

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If you are not comfortable with these skills, that's fine, but speak to us after class so we can help you get these skills.

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And you need to be able to perform these tasks instinctively, without having to think about it too much.

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But before we can teach you all those ninja skills, we have to talk about the most important programming skill of all, which is ...

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Here are at least six places to go for help:

- 1. Python documentation
- 2. Google
- 3. Stack Overflow
- 4. PySlackers
- 5. Internet relay chat (IRC) rooms with other Python users
- 6. Various Python mailing lists

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BUT, like any online community, there's the potential for a **toxic culture** to destroy everything.

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Actively toxic communities are easy to identify. They encourage and are characterized by **overt** sexism, racism, bigotry, and calls for violence or other aggression against individuals.



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- ▶ and individuals are often unaware of when they are acting in a passively toxic way.



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Downvotes without explanation: this can be very upsetting to anyone, especially to people with less experience

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Overzealous curation: Being very quick to tag a question as a "duplicate" without checking to see nuanced ways in which the question comes from a new situation.

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Passive toxicity **shrinks** the community and makes it **more homogeneous**.

Across society, small, homogeneous communities are much more likely to exclude or discriminate against people based on sex, race, class, language and other factors. And that leads to many ethical problems.

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- 3. The type what kind of object is this?

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The most important skill is to know how to read the docstring to quickly find the information you need.

To understand how to read the docstring, call up the docstring for a linear regression class object from the sklearn package:

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import sklearn.linear_model
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The header tells us that the LinearRegression object is a class, stored in the linear_model.base module within the sklearn package.

2. The signature

LinearRegression(fit_intercept=True, normalize=False,
copy_X=True, n_jobs=None)

Some docstrings list the signature, although the signature can be accessed elsewhere. The signature lists all of the parameters of a function.

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3. The short description

Ordinary least squares Linear Regression.

A one-or-two sentence summary of what the function does.

4. The parameters section is the most useful for learning how to use a function:

Parameters

for more details.

```
fit intercept: boolean, optional, default True
    whether to calculate the intercept for this model. If set
    to False, no intercept will be used in calculations
    (e.g. data is expected to be already centered).
normalize: boolean, optional, default False
    This parameter is ignored when ``fit intercept`` is set to False.
    If True, the regressors X will be normalized before regression by
    subtracting the mean and dividing by the l2-norm.
    If you wish to standardize, please use
    :class:`sklearn.preprocessing.StandardScaler` before calling ``fit`` on
    an estimator with ``normalize=False``.
copy X: boolean, optional, default True
    If True, X will be copied; else, it may be overwritten.
n jobs : int or None, optional (default=None)
    The number of jobs to use for the computation. This will only provide
    speedup for n_targets > 1 and sufficient large problems.
    ``None`` means 1 unless in a :obj:`joblib.parallel backend` context.
    ``-1`` means using all processors. See :term:`Glossary <n_jobs>`
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Each parameter is described in a sentence or two to explain what the parameter does.

5. The attributes

Attributes

coef_ : array, shape (n_features,) or (n_targets, n_features)
 Estimated coefficients for the linear regression problem.
 If multiple targets are passed during the fit (y 2D), this
 is a 2D array of shape (n_targets, n_features), while if only
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If the output is saved in an object named regress, to access the coefficients, type regress.coef_, and to access the intercept, type regress.intercept_.

6. The examples

Examples

```
>>> import numpy as np
>>> from sklearn.linear_model import LinearRegression
>>> X = np.array([[1, 1], [1, 2], [2, 2], [2, 3]])
>>> # y = 1 * x_0 + 2 * x_1 + 3
>>> y = np.dot(X, np.array([1, 2])) + 3
>>> reg = LinearRegression().fit(X, y)
>>> reg.score(X, y)
1.0
>>> reg.coef_
array([1, 2.])
>>> reg.intercept_ # doctest: +ELLIPSIS
3.0000...
>>> reg.predict(np.array([[3, 5]]))
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Examples are meant to be run, not just looked at. Copy-and-paste the examples into your notebook or script, run the code. See if you can do more things with the given objects than the examples do.

self: returns an instance of self.

7. The <u>related methods</u> defines methods that <u>expand the</u> <u>functionality</u> of the one you are looking at, along with their own documentation:

```
Methods defined here:
init (self, fit intercept=True, normalize=False, copy X=True, n jobs=None)
    Initialize self. See help(type(self)) for accurate signature.
fit(self, X, y, sample weight=None)
    Fit linear model.
    Parameters
    X : array-like or sparse matrix, shape (n samples, n features)
        Training data
    y : array_like, shape (n_samples, n_targets)
        Target values. Will be cast to X's dtype if necessary
    sample weight: numpy array of shape [n samples]
        Individual weights for each sample
        .. versionadded:: 0.17
           parameter *sample weight* support to LinearRegression.
    Returns
```

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Stack Overflow is the most popular and most useful website for help with programming of all kinds. Google searching a Python problem will usually lead to a Stack Overflow post on the same issue.

Python is now the **most frequent** tag for posts on Stack Overflow: see the video embedded on this blog post.

Finding a Stack Overflow post that's relevant to your problem can give you both the code and intuition to solve your problem.

Or maybe not! Small differences in the situation can make the solution irrelevant to you. **Be cautious** and don't treat a Stack Overflow post as automatically a definitive answer.

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Going for reputation is an **entirely optional** activity. If you don't want to worry about it, don't.



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If you do post to Stack Overflow, you are likely to get some very useful responses if you follow some guidelines. There's **a strategy for getting good responses**: stackoverflow.com/help/how-to-ask

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So spend a significant amount of time digging through the internet. If there's something similar, but not quite what you need, you can say so in your post.



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Good: How to place labels on top of points in a matplotlib scatterplot?

Step 3: Start the post with a paragraph describing the problem in more detail.

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Some good things to include in this paragraph:

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- ▶ You can write the version of Python you are using, the version of the modules, and the operating system on your computer, in case the problem turns out to be specific to one of those

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If the code needs to run on data, can you use something pre-loaded in Python that everyone can access? (There are example datasets included with psykitlearn, for example.)

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If the code needs to run on data, can you use something pre-loaded in Python that everyone can access? (There are example datasets included with psykitlearn, for example.)

Make the code as short as possible, and use comments, to help people understand the code more quickly.

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Don't ask about **homework problems**. (Here's an example of someone getting called out on this)

Interacting with other Python users on PySlackers

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Some useful channels:

- data_science
- python_
- job_advice

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Internet chatrooms can be rough places, but the #python channel claims to enforce this Code of Conduct:

https://www.python.org/psf/codeofconduct/.

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- 3. To use the #python channel, you need to register your nickname. To check if your nickname is unique, click on the "freenode" tab on the left-hand sidebar. A text box will appear on the bottom of the screen. Type:

/msg NickServ info

4. Step 3 will open a new tab. Switch to that tab. If no one else already has your nickname, you will see

```
NickServ: (notice) <nickname> is not registered.
```

If you see something else, it means someone already has your nickname. You can change your nickname right here by typing /nick followed by another nickname. Then type /msg NickServ info again. Repeat until you see the message listed above.

<u>Important note</u>: DON'T use a password here that you use for important things like **email**, **bank accounts**, **etc**.

We shouldn't have the same faith in the security of Freenode's servers as we can have in Google's.

Also, this is the kind of platform that tends to attract hackers. And for people used to a graphical user interface, it might be easy to mistype in a way that accidentally displays your password in the chat.

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Use a unique, throwaway password!

5. To register this nickname, type

```
/msg NickServ register <password> <email-address>
```

where <password> is a password you will use in the future, and <email address> is the email you want associated with this account.

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You are free to chat away. Pay attention to the guidelines that appear as links on the top of the screen.



Python mailing lists and message boards

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If you have a question for the Python core development team, send an email to help@python.org. The team is pretty busy, so be sure to check other resources and lists for an answer first.

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It's OKAY to mix styles, packages, and approaches. Use whatever works, but **keep track of what you do**.

Through the 1970s, data was stored on punch cards and fed directly to a mainframe computer capable of regression analysis.



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ASCII – American Standard Code for Information Interchange pronounced "As-Key"

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- Data points usually delimited by commas, spaces, or tabs. Might require a data dictionary to read.



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We will go over individual data files today, and databases soon.



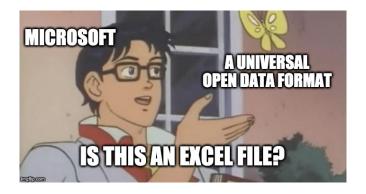
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A comma-separated values (CSV) file:

<u>Note</u>: Although the CSV format is universal, Excel sometimes opens by default when you double-click on the CSV file. But, CSV files are NOT exclusive to Excel.

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A tab delimited file:

sez	k r	ace	r	egio	on h	appy	7 1	ife	s	ibs		child	ls	age	ec	luc
pae				_					occcat80			tax		~		obey
poi	popular th		thnkself		wor			hel	poth	oth hlth1		hlth2		hlth3		•
	th4		th5	hlth6		hlth7		hlth8		hlth9		work1		work2		
WO	work3		work4		work5		work6		work7		work8		work9		prob1	
prob2		prob3		prob4										•		
2	1	1	1	1	1	2	61	12	97	12	97	22	3	1	1	0
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
0	0	0	0	0												
2	1	1	2	1	2	1	32	20	20	18	20	75	1	1	0	5
4	1	2	3	1	1	2	2	2	2	2	2	2	2	2	1	2
2	1	2	1	1	2	4	5									
1	1	1	1	0	2	1	35	20	16	14	17	59	1	0	1	5
4	1	2	3	2	2	2	2	2	2	2	2	2	2	2	2	2
2	2	2	2	2												
2	1	1	9	2	2	0	26	20	20	20	97	48	1	1	0	4
5	1	3	2	1	2	2	2	2	2	2	2	2	2	1	2	2
2	2	2	2	2	2	2										
2	2	1	2	1	4	0	25	12	98	98	97	42	3	1	1	0
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
0	0	0	0	0												

A fixed-width ASCII file with no delimination. Files like these minimize memory (no need to store a bunch of commas), but require a dictionary file to read them.

Dictionary:

- Variable 1: sex, column 1
- Variable 2: race, column 2
- **.** . . .
- ▶ Variable 8: age, columns 8-9

Before we go over the functions, it is useful to set the working directory at the start of your script or notebook.

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To set the working directory:

- ▶ Load the os package: import os
- ► Type the folder's address into os.chdir("folder")

To **check** on the path Python is currently using as a default, type os.getcwd() into the console.

Changing the working directory

To **check** on the path Python is currently using as a default, type os.getcwd() into the console.

If you want to change the working directory back after you've run the relevant code:

```
import os
oldpath = os.getcwd()
os.chdir("folder")

#(Your code goes here)
os.chdir(oldpath)
```

We will be using the Pandas package:

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- 3. The URL of a data file that's accessible online

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You can load the anes_example.csv data by either **downloading** and unzipping the file, or by using the URL:

https://raw.githubusercontent.com/NovaVolunteer/ Practice_Application_DS/master/Week%205/anes_example.csv

If you download and unzip the ANES data, and you've already changed your working directory, then to load the ANES data, type

```
anes = pd.read_csv("anes_example.csv")
```

If you download and unzip the ANES data, and you've already changed your working directory, then to load the ANES data, type

```
anes = pd.read_csv("anes_example.csv")
```

If you want to load the data directly from the URL, save the URL as a separate object, then pass this to the function:

```
url = "https://raw.githubusercontent.com/NovaVolunteer/
    Practice_Application_DS/master/Week%205/
    anes_example.csv"
anes = pd.read_csv(url)
```

Before we get to the other parameters of the pd.read_csv() function, let's talk about the workflow of loading data.

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- Examine the loaded dataframe object to make sure the data was correctly read
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There's an important set of functions in Python that let you quickly explore a dataframe.

If you are using a <u>Jupyter Notebook</u>, typing the name of the data frame **in its own cell** will produce a good-looking HTML table illustrating the data frame.

3]:	anes								
3]:		caseid	turnout12	turnout12b	vote12	percent16	meet	givefut	info
	0	1.0	1	NaN	2.0	100	1	3	4
	1	2.0	2	NaN	NaN	50	4	5	4
	2	3.0	1	NaN	1.0	100	1	1	1
	3	4.0	1	NaN	2.0	100	5	4	5
	4	5.0	1	NaN	1.0	100	2	1	3

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	0	1.0	1	NaN	2.0	100	1	3	4
	1	2.0	2	NaN	NaN	50	4	5	4
	2	3.0	1	NaN	1.0	100	1	1	1
	3	4.0	1	NaN	2.0	100	5	4	5
	4	5.0	1	NaN	1.0	100	2	1	3

If you are using <u>Spyder</u>, look in the <u>upper-right window</u> and select the "Variable explorer" tab. Clicking on the data frame will open a separate window for viewing the data.

One annoying thing about Jupyter's interactive viewer is that it omits the columns in the middle for data frames with more than about 20 columns:

ane	es												
		caseid	turnout12	turnout12b	vote12	percent16	meet	givefut	info	march	sign	 votereg	pid3
	0	1.0	1	NaN	2.0	100	1	3	4	1	2	 1	1
	1	2.0	2	NaN	NaN	50	4	5	4	2	2	 2	3
	2	3.0	1	NaN	1.0	100	1	1	1	1	1	 1	2
	3	4.0	1	NaN	2.0	100	5	4	5	2	2	 1	1
4	4	5.0	1	NaN	1.0	100	2	1	3	1	2	 1	4
	5	6.0	1	NaN	3.0	100	3	3	2	2	1	 1	3

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anes												
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1	2.0	2	NaN	NaN	50	4	5	4	2	2	 2	3
2	3.0	1	NaN	1.0	100	1	1	1	1	1	 1	2
3	4.0	1	NaN	2.0	100	5	4	5	2	2	 1	1
4	5.0	1	NaN	1.0	100	2	1	3	1	2	 1	4
5	6.0	1	NaN	3.0	100	3	3	2	2	1	 1	3

The columns it skipped (about **148** in this case) are replaced by a column of dots.

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anes												
	caseid	turnout12	turnout12b	vote12	percent16	meet	givefut	info	march	sign	 votereg	pid3
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1	2.0	2	NaN	NaN	50	4	5	4	2	2	 2	3
2	3.0	1	NaN	1.0	100	1	1	1	1	1	 1	2
3	4.0	1	NaN	2.0	100	5	4	5	2	2	 1	1
4	5.0	1	NaN	1.0	100	2	1	3	1	2	 1	4
5	6.0	1	NaN	3.0	100	3	3	2	2	1	 1	3

The columns it skipped (about **148** in this case) are replaced by a column of dots.

To keep Python from skipping columns, you can change this behavior **globally** (for all subsequent code) or **locally** (for each line of code individually).



To always display all of the columns, type

pd.set_option('display.max_columns', None)

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```
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```
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Caution: If you are working with large dataframes, it's probably not a good idea to always display ALL of the rows and columns.

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```
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```

Caution: If you are working with large dataframes, it's probably not a good idea to always display ALL of the rows and columns.

To keep a specific line of code from skipping variables, use the anes.loc and anes.iloc functions. (Replace "anes" with the name of your dataframe object.)

anes.loc allows you to select columns of a data frame by name, and anes.iloc allows you to select columns by column number.

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To see the "sign", "give12mo", and "ftobama" variables, type

anes.loc[:, ['sign', 'give12mo', 'ftobama']]

	sign	give12mo	ftobama
0	2	2	100.0
1	2	2	39.0
2	1	1	1.0
3	2	2	89.0
4	2	1	1.0
5	1	1	0.0
6	2	1	73.0
7	1	2	0.0
8	2	1	12.0

To see all variables in between "sign", and "fthisp", type

anes.loc[:, 'sign':'fthisp']

	sign	give12mo	compromise	ftobama	ftblack	ftwhite	fthisp
0	2	2	1	100.0	100.0	100	100.0
1	2	2	1	39.0	6.0	74	6.0
2	1	1	2	1.0	50.0	50	50.0
3	2	2	1	89.0	61.0	64	61.0
4	2	1	2	1.0	61.0	58	71.0
5	1	1	2	0.0	50.0	51	51.0
6	2	1	1	73.0	100.0	70	100.0
7	1	2	1	0.0	70.0	70	69.0
8	2	1	2	12.0	50.0	50	50.0

To select columns and rows numerically, use anes.iloc. To see rows 254 through 262 and all columns, type

```
anes.iloc[254:262, :]
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To see all rows, columns 21 through 30, type

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anes.iloc[:, 21:30]
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To select columns and rows numerically, use anes.iloc. To see rows 254 through 262 and all columns, type

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anes.iloc[254:262, :]
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To see all rows, columns 21 through 30, type

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anes.iloc[:, 21:30]
```

To see rows 254 through 262, columns 21 through 30, type

```
anes.iloc[254:262, 21:30]
```

To see only the first 10 rows of the data, type anes.head(10). Replace 10 with however many rows you want to see.

To see only the first 10 rows of the data, type anes.head(10). Replace 10 with however many rows you want to see.

To see only the last 10 rows of the data, type anes.tail(10).

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To see only the last 10 rows of the data, type anes.tail(10).

Typing anes.info() tells us the dimensions of the data, the number of variables of each type, and the size of the dataframe in memory:

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1200 entries, 0 to 1199

Columns: 168 entries, caseid to ever_vs_12mo_rand

dtypes: float64(76), int64(86), object(6)

memory usage: 1.5+ MB

anes.columns lists all the variable names.

If there are too many variables, Python will abbreviate the list with "..." To see the omitted items, change the maximum number of items that can display in a list with:

```
pd.set_option('display.max_seq_items', None)
```

(Again, be careful about removing this limit for data frames with a large number of columns)

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```

(Again, be careful about removing this limit for data frames with a large number of columns)

anes.dtypes lists the variables along with their types (int64 for integers, float64 for numbers with decimals, object for variables that might be either categorical or string).

anes.describe() shows basic summary statistics for every variable in the dataframe.

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There are different summary statistics for different types of variables. By default, anes.describe() displays stats only for the float and int types:

- count number of non-missing observations
- mean the sample mean
- ▶ std the sample standard deviation
- ▶ min the minimum value
- ▶ 25% the 25th percentile
- ▶ 50% the median value
- ▶ 75% the 75th percentile
- ▶ max the maximum value

Use the percentiles argument to display different percentiles. To see the 20th, 37.5th, and 74.23th percentiles, type

```
anes.describe(percentiles = [.20, .375, .7423])
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anes.describe(include = "int"), and to see just the float
variables, type anes.describe(include = "float").

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anes.describe(include = "object"). These variables have
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variables, type anes.describe(include = "float").

To see object variables, type

anes.describe(include = "object"). These variables have
different stats:

- count number of non-missing observations
- unique number of unique observations
- ▶ top the most frequent value
- ▶ freq the frequency of the top value



To see all of the variables, type anes.describe(include = "all"), but this will result in NA values for stats that aren't relevant to the variable.

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Use these tools a lot! After loading the data, you need to quickly be able to see if there were any problems with loading the data. Ask:

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- Are the dimensions what I expect?
- ► Are the variable names set to what they are supposed to be?
- Are there any bizarrely high/low means or other stats?

There are many reasons why a load might have failed. Fortunately, there are parameters within the pd.read_csv() function to deal with many of these issues.

```
pd.read_csv(filepath_or_buffer, sep, header)
```

sep or **delimiter** – (string) The symbol that is used in the file to separate one datapoint from the next on the same row. By default, it looks for commas.

► For tab-delimited, use sep="\t"

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- header=None assumes there are no variable names and that the first row is data. It labels the columns with numbers, but if you also type prefix="X" the variables will be X0, X1, ...

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- ► header=j uses the j_{th} row for variable names, and deletes all higher rows

```
pd.read_csv(filepath_or_buffer, sep, header, usecols)
```

usecols – (a list of strings or integers) Use this if you only want some of the columns to be loaded from the outset:

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- usecols = ["caseid", "vote12", "meet"] only loads the variables named "caseid", "vote12", and "meet", as recognized by whatever Python thinks is the header

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In general, don't use this parameter unless the data file is too large to load in its entirety. You can delete columns later.

```
pd.read_csv(filepath_or_buffer, sep, header, usecols,
skiprows, skipfooter, nrows)
```

skiprows – (integer, or a list of integers) Likewise, which rows to skip when loading the data:

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skiprows=3 skips the first three rows of the data. If header is left to its default, the 4th row is assumed to contain the variable names

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- skiprows=[0,3,5] skips the 1st, 4th, and 6th rows

skipfooter - same as skiprows but counts up from the
bottom row

nrows - (integer) only loads the first several rows, as specified by the user



```
pd.read_csv(filepath_or_buffer, sep, header, usecols,
skiprows, skipfooter, nrows, na_values)
```

na_values - (list of strings or numeric) Sometimes data authors
use codes other than NA to indicate a missing value.

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 $\frac{\text{Example:}}{-7, -8, -9}$, and 998, as well as blank cells and NA to represent missing values.

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na_values - (list of strings or numeric) Sometimes data authors
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Example: the American National Election Study (ANES) data uses $\overline{-7, -8, -9}$, and 998, as well as blank cells and NA to represent missing values.

To replace all these values with NA across the whole data frame, type $na_values = [-7, -8, -9, 998]$.

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Caution: Only specify missing codes in the pd.read_csv() function if the code ALWAYS means a missing value. If 998 is a valid datapoint for some variables, you can replace the missing codes for relevant variables later.

```
pd.read_csv(filepath_or_buffer, sep, header, usecols,
skiprows, skipfooter, nrows, na_values, comment)
```

comment - (string) If there are comments in the data file itself (it shouldn't happen but it does!), what character to read as indicating a commented-out row.

```
pd.read_csv(filepath_or_buffer, sep, header, usecols,
skiprows, skipfooter, nrows, na_values, comment)
```

comment - (string) If there are comments in the data file itself (it shouldn't happen but it does!), what character to read as indicating a commented-out row.

If the data authors wrote "# Collected on Mon 9/23" before some rows, then "# Collected on Tues 9/24" further down, you can ignore these by typing <code>comment="#"</code>.

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If the data authors wrote "# Collected on Mon 9/23" before some rows, then "# Collected on Tues 9/24" further down, you can ignore these by typing <code>comment="#"</code>.

Careful: if the comment-symbol appears ANYWHERE on the row, the remainder of the row is not read. That's a problem if, for example, the data contain tweets and one tweet reads "UVA is #1!".

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We'll go over all of that in detail soon. But after having cleaned the data, you might want to **save the cleaned dataframe as a CSV** or as a different ASCII file.

Once the data are loaded into Python, there are many tools, techniques, and functions to know to get the data into a clean state.

We'll go over all of that in detail soon. But after having cleaned the data, you might want to **save the cleaned dataframe as a CSV** or as a different ASCII file.

Suppose the anes object contains a cleaned dataframe. To save it as a CSV, use anes.to_csv(). There are several parameters, you can see with help(anes.to_csv).

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Suppose the anes object contains a cleaned dataframe. To save it as a CSV, use anes.to_csv(). There are several parameters, you can see with help(anes.to_csv).

Let's talk about two important parameters: anes.to_csv(path_or_buf, sep)

```
anes.to_csv(path_or_buf, sep)
```

path_or_buf - (string) the name of the file to save, with the
appropriate file extension (.csv, .txt, etc.)

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appropriate file extension (.csv, .txt, etc.)

You can write an entire file path here if you want. But if you set the working directory, and write the file name alone, it will save in the working directory.

```
anes.to_csv(path_or_buf, sep)
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path_or_buf - (string) the name of the file to save, with the
appropriate file extension (.csv, .txt, etc.)

You can write an entire file path here if you want. But if you set the working directory, and write the file name alone, it will save in the working directory.

sep - (string) the character to use as a delimiter. A comma by default. Use $sep="\t"$ for a tab-delimited file.

```
anes.to_csv(path_or_buf, sep)
```

path_or_buf - (string) the name of the file to save, with the appropriate file extension (.csv, .txt, etc.)

You can write an entire file path here if you want. But if you set the working directory, and write the file name alone, it will save in the working directory.

sep – (string) the character to use as a delimiter. A comma by default. Use sep="\t" for a tab-delimited file.

To save the anes dataframe as a standard CSV file, type:

```
anes.to_csv("anes_cleaned.csv", sep=",")
```



A fixed-width file contains no delimiters. Instead, it aligns all of the data for one variable in the **same position** on each row. These files might use less memory than CSV.

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But that makes the data impossible to parse without an <u>external list</u> of which variable is stored where. The first and most important step is to <u>get this list</u>.

Example: the National Journal conducted a public opinion poll and saved the data in fixed-width format. I saved the codebook on GitHub, and the data here:

https://raw.githubusercontent.com/NovaVolunteer/ Practice_Application_DS/master/Week%205/njcc33850.dat

In this codebook, find the variable names and save them in a list, for example:

```
datanames = ['psraid', 'sample', 'int_date', 'area',
  'state', 'cregion', 'density', 'usr', 'cc1', 'cc1a',
  'cc2', 'cc3', 'cc4', 'cc5', 'cc6', 'cc7', 'ql1', 'ql1a',
  'qc1', 'hh1', 'employ', 'par', 'sex', 'age', 'educ2',
  'hisp', 'race', 'inc', 'income', 'reg', 'party',
  'partyln', 'iphoneus', 'hphoneus', 'recage', 'receduc',
  'racethn', 'standwt', 'raceos']
```

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  'cc2', 'cc3', 'cc4', 'cc5', 'cc6', 'cc7', 'q11', 'q11a',
  'qc1', 'hh1', 'employ', 'par', 'sex', 'age', 'educ2',
  'hisp', 'race', 'inc', 'income', 'reg', 'party',
  'partyln', 'iphoneus', 'hphoneus', 'recage', 'receduc',
  'racethn', 'standwt', 'raceos']
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There are two ways to proceed next:

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  'qc1', 'hh1', 'employ', 'par', 'sex', 'age', 'educ2',
  'hisp', 'race', 'inc', 'income', 'reg', 'party',
  'partyln', 'iphoneus', 'hphoneus', 'recage', 'receduc',
  'racethn', 'standwt', 'raceos']
```

There are **two ways** to proceed next:

Method 1: If you know how many characters each variable takes, at maximum, save these widths as a list:

<u>Method 2</u>: if you know the starting and ending position of each variable, create a **list of length 2** for each variable, where

- the first element is the column the previous variable ends on (or 0 for the first variable)
- ▶ and the second element is the column the current variable ends on.

<u>Method 2</u>: if you know the starting and ending position of each variable, create a **list of length 2** for each variable, where

- the first element is the column the previous variable ends on (or 0 for the first variable)
- ▶ and the second element is the column the current variable ends on.

For example, if a variable occupies columns 34, 35, and 36, its list of length 2 is [33,36] .

<u>Method 2</u>: if you know the starting and ending position of each variable, create a **list of length 2** for each variable, where

- the first element is the column the previous variable ends on (or 0 for the first variable)
- ▶ and the second element is the column the current variable ends on.

For example, if a variable occupies columns 34, 35, and 36, its list of length 2 is [33,36].

Create a list-of-lists, which can look like:

```
datapos = [[0,6], [6,7], [7,13], [13,16], [16,18], [18,19], [19,20], [20,23], [23,24], [24,25], [25,26], [26,27], [27,28], [28,29], [29,30], [30,31], [31,32], [32,33], [33,34], [34,35], [35,36], [36,37], [37,38], [38,40], [40,41], [41,42], [42,43], [43,45], [45,46], [46,47], [47,48], [48,49], [49,50], [50,51], [51,52], [52,53], [53,54], [54,58], [58,88]]
```

To read the fixed-width file, use the pd.read_fwf() function.

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To parse the data using variable widths, save the URL, the variable names, and widths in separate objects (as on the previous slides), and type:

```
njcc = pd.read_fwf(url, widths=datawidths,
    header=None, names=datanames)
```

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```

To parse the data using variable positions, save the URL, the variable names, and positions in separate objects, and type:

```
njcc = pd.read_fwf(url, colspecs=datapos,
    header=None, names=datanames)
```



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But that's not a good strategy because it requires me to have access to Excel. To work entirely with Python, use the pd.read_excel() function.

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My usual strategy: Open Excel, select the sheet I want, and save it as a CSV before loading it in Python.

But that's not a good strategy because it requires me to have access to Excel. To work entirely with Python, use the pd.read_excel() function.

Many of the parameters that work for pd.read_csv() work for pd.read_excel() too, including: header, names, usecols, skiprows, skipfooter, nrows, na_values, and comment.

There are two arguments we should go over: pd.read_excel(io, sheet_name)

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- path and filename,
- filename alone (if you've set the working directory),
- or a URL where the Excel file is stored online.

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sheet_name - (string, int, or list) If the Excel file has sheets with
names, you can type the name of the sheet here. Or type a
number: 0 refers to the first sheet, 1 to the second, etc.

If you specify a list, pd.read_excel() will produce a list of dataframes, one for each sheet you specify. Typing sheet_name = None produces a list with all of the sheets.



Example: I saved an Excel sheet on GitHub with NBA statistics, here:

https://github.com/NovaVolunteer/ Practice_Application_DS/blob/master/Week%205/ NBA-Team-Sample-BoxScore-Dataset.xlsx?raw=true

This Excel file has four sheets:

- ▶ NBA-TEAM-SAMPLE has team stats for every game last season;
- METADATA defines variables;
- TEAMS provides team names and locations;
- PROVIDE DATE FORMAT has information abouty date formats.

I save the URL as an object. Then, to load the NBA-TEAM-SAMPLE sheet, I type one of these lines:

```
nba = pd.read_excel(url, sheet_name="NBA-TEAM-SAMPLE")
nba = pd.read_excel(url, sheet_name=0)
```

I save the URL as an object. Then, to load the NBA-TEAM-SAMPLE sheet, I type one of these lines:

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nba = pd.read_excel(url, sheet_name="NBA-TEAM-SAMPLE")
nba = pd.read_excel(url, sheet_name=0)
```

To load the **TEAMS** sheet, I type one of these lines:

```
nba = pd.read_excel(url, sheet_name="TEAMS")
nba = pd.read_excel(url, sheet_name=2)
```

I save the URL as an object. Then, to load the NBA-TEAM-SAMPLE sheet, I type one of these lines:

```
nba = pd.read_excel(url, sheet_name="NBA-TEAM-SAMPLE")
nba = pd.read_excel(url, sheet_name=0)
```

To load the **TEAMS** sheet, I type one of these lines:

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nba = pd.read_excel(url, sheet_name="TEAMS")
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```

To load both sheets:

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BUT until recently most data science was conducted using proprietary software: SAS, Stata, or SPSS. Many researchers still use these platforms. So **you will likely have to work with these files.**

Like with Excel, opening SAS/Stata/SPSS and saving as CSV is a bad solution because you need the software to do that, and the software is expensive.

Regular **SAS** files have the extension .sas7bdat, and compressed SAS files ("transport files") have the extension .xport. We'll work with a dataset on inflation, here:

https://github.com/NovaVolunteer/Practice_Application_DS/blob/master/Week%205/inflation.sas7bdat?raw=true

Regular **SAS** files have the extension .sas7bdat, and compressed SAS files ("transport files") have the extension .xport. We'll work with a dataset on inflation, here:

```
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```

Stata files all have the extension .dta. We'll work with a CBS news poll, here:

```
https://github.com/NovaVolunteer/Practice\_Application\_DS/\\blob/master/Week\%205/cbspoll.dta?raw=true
```

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```

Stata files all have the extension .dta. We'll work with a CBS news poll, here:

```
https://github.com/NovaVolunteer/Practice_Application_DS/blob/master/Week%205/cbspoll.dta?raw=true
```

SPSS files have the extension .sav, or .zsav for compressed files. We'll work with the ANES in SPSS format:

```
https://github.com/NovaVolunteer/Practice_Application_DS/blob/master/Week%205/anes_timeseries_2016.sav?raw=true
```

You can load SAS and Stata files with Pandas using the pd.read_sas() and pd.read_stata() functions.

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To load an SPSS file, you need to install the pyreadstat package,

pip install pyreadstat

and import this package

import pyreadstat

Then you can use the pyreadstat.read_sav() function.

These functions are very similar to pd.read_csv(), but one important difference is they can't read a URL. So you have to download a local copy of the files.

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To load the SAS inflation data: inflation = pd.read_sas("inflation.sas7bdat")

These functions are very similar to <code>pd.read_csv()</code>, but one important difference is they can't read a URL. So you have to download a **local copy** of the files.

```
To load the SAS inflation data:
inflation = pd.read_sas("inflation.sas7bdat")

To load the Stata CBS poll data:
cbspoll = pd.read_stata("cbspoll.dta")
```

These functions are very similar to pd.read_csv(), but one important difference is they can't read a URL. So you have to download a local copy of the files.

```
To load the SAS inflation data:
inflation = pd.read_sas("inflation.sas7bdat")
```

```
To load the Stata CBS poll data:
cbspoll = pd.read_stata("cbspoll.dta")
```

Loading SPSS data is trickier. You have to define **two objects**, separated by a comma. The first object will contain the dataframe, and the second object will contain the SPSS metadata:

```
anes_spss, anes_spss_meta =
    pyreadstat.read_sav("anes_timeseries_2016.sav")
```

All of the electronic data files we've discussed so far are sometimes called flat files or rectangular files.

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- ► All entries in a column have the same type (all string, for example, or all numeric, but not a mix of string and numeric)

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- On each row, there's an entry for every field (column), even if that entry is a missing value.
- ► All entries in a column have the same type (all string, for example, or all numeric, but not a mix of string and numeric)

But what if we had non-rectangular data, where every record might have different fields, or a different data type for a field than another record? A JSON file is designed for this situation.

 $\mathsf{JSON} = \mathsf{JavaScript}\ \mathsf{Object}\ \mathsf{Notation}$

Although JSON notation is based on JavaScript, it's **portable** to many other programming languages, like Python and R.

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CSV/ASCII is a text file with specific rules about how datapoints are typed into the file (i.e. separated by commas, column names on the first row, rows separated by carriage returns.)

JSON is just another text file, with a different method for writing the data.

If this is our CSV file:

```
caseid,fttrump,fthrc,birthyr,gender
1,1.0,76.0,1960,1
2,28.0,52.0,1957,2
3,100.0,1.0,1963,1
4,0.0,69.0,1980,1
5,13.0,1.0,1974,1
```

If this is our CSV file:

```
caseid,fttrump,fthrc,birthyr,gender
1,1.0,76.0,1960,1
2,28.0,52.0,1957,2
3,100.0,1.0,1963,1
4,0.0,69.0,1980,1
5,13.0,1.0,1974,1
```

We can represent the same data using the JSON format:

```
[{'caseid': 1.0, 'fttrump': 1.0, 'fthrc': 76.0,
    'birthyr': 1960, 'gender': 1},
{'caseid': 2.0, 'fttrump': 28.0, 'fthrc': 52.0,
    'birthyr': 1957, 'gender': 2},
{'caseid': 3.0, 'fttrump': 100.0, 'fthrc': 1.0,
    'birthyr': 1963, 'gender': 1},
{'caseid': 4.0, 'fttrump': 0.0, 'fthrc': 69.0,
    'birthyr': 1980, 'gender': 1},
{'caseid': 5.0, 'fttrump': 13.0, 'fthrc': 1.0,
    'birthyr': 1974, 'gender': 1}]
```

Let's look at just the first observation (in JSON-lingo, observations are called records):

```
{
    'caseid': 1.0,
    'fttrump': 1.0,
    'fthrc': 76.0,
    'birthyr': 1960,
    'gender': 1
}
```

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}
```

This style is called object literal syntax. In Python, the curly braces { and } represent a **set**. A set is like a list (which we create using square braces [and]) with two differences: elements cannot be repeated, and there's no ordering of the elements.

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This style is called object literal syntax. In Python, the curly braces { and } represent a **set**. A set is like a list (which we create using square braces [and]) with two differences: **elements** cannot be repeated, and there's no ordering of the elements.

Typing [1,2,3] and $\{1,2,3\}$ return the same output. But [3,3,2,1] and $\{3,3,2,1\}$ do not, because the $\{\text{set}\}$ removes the extra 3 and sorts the elements.

```
{
    'caseid': 1.0,
    'fttrump': 1.0,
    'fthrc': 76.0,
    'birthyr': 1960,
    'gender': 1
}
```

Also, elements of a list cannot be named, but elements of a set can be given names: caseid is the name, 1.0 is the element.

```
{
    'caseid': 1.0,
    'fttrump': 1.0,
    'fthrc': 76.0,
    'birthyr': 1960,
    'gender': 1
}
```

Also, elements of a list cannot be named, but elements of a set can be given names: caseid is the name, 1.0 is the element.

A Python set that contains variable names and data points is called a **dictionary**. The fact that the dictionary is a set means that variable names cannot be repeated, and the order the variables are entered doesn't matter.

```
{
    'caseid': 1.0,
    'fttrump': 1.0,
    'fthrc': 76.0,
    'birthyr': 1960,
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A Python set that contains variable names and data points is called a **dictionary**. The fact that the dictionary is a set means that variable names cannot be repeated, and the order the variables are entered doesn't matter.

A JSON file is a <u>list of dictionaries</u>: one dictionary for every record.

JSON and CSV are just two ways to save data in a text file. There are pros and cons to each method.

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▶ Data in which variables are stored with different data types from record to record

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- ▶ Data in which variables are stored with different data types from record to record
- ▶ Data in which different records have different variables

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But <u>JSON</u> is more <u>flexible</u>, and can store data structures that are awkward or impossible for CSV, such as:

- ▶ Data in which variables are stored with different data types from record to record
- Data in which different records have different variables
- Data with a tree-based nesting structure

JSON can store data in which variables are stored with different data types from record to record:

```
[{'caseid': 1.0,
  'fttrump': 'Awful',
  'fthrc': 'Pretty good',
  'birthyr': 1960,
  'gender': 1},
  {'caseid': 2.0,
  'fttrump': 28.0,
  'fthrc': 52.0,
  'birthyr': 1957,
  'gender': 2}]
```

JSON can store data in which different records have different variables:

JSON can store data with a **tree-based** nesting structure:

```
[{'id': 1,
  'name': 'Leanne Graham',
  'username': 'Bret',
  'email': 'Sincere@april.biz',
  'address': {'street': 'Kulas Light',
   'suite': 'Apt. 556',
   'city': 'Gwenborough',
   'zipcode': '92998-3874',
   'geo': {'lat': '-37.3159', 'lng': '81.1496'}},
  'phone': '1-770-736-8031 x56442',
  'website': 'hildegard.org',
  'company': {'name': 'Romaguera-Crona',
   'catchPhrase': 'Multi-layered client-server neural-net',
   'bs': 'harness real-time e-markets'}}]
```

In a tree-based nesting structure, individual fields can themselves contain dictionaries. Nested parts of the JSON are highlighted below:

```
[{'id': 1,
  'name': 'Leanne Graham',
  'username': 'Bret',
  'email': 'Sincere@april.biz',
  'address': {'street': 'Kulas Light',
  'suite': 'Apt. 556',
  'city': 'Gwenborough',
  'zipcode': '92998-3874',
  'geo': {'lat': '-37.3159', 'lng': '81.1496'}},
  'phone': '1-770-736-8031 x56442',
  'website': 'hildegard.org',
  'company': {'name': 'Romaguera-Crona',
  'catchPhrase': 'Multi-layered client-server neural-net',
  'bs': 'harness real-time e-markets'}}]
```

Who uses JSON?

You are most likely to encounter JSON files when working with APIs (Application Programming Interface), which are servers that you can use to retrieve and send data to using code (more on that soon). Many APIs find it convenient to send data in this format.

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You are most likely to encounter JSON files when working with APIs (Application Programming Interface), which are servers that you can use to retrieve and send data to using code (more on that soon). Many APIs find it convenient to send data in this format.

Some databases (especially "NoSQL" types such as MongoDB) work best with JSON formatted data.

JSON is sometimes referred to as a "readable" or "lightweight" version of other structured data formats: XML (Extensible Markup Language) and YAML (Yet Another Markup Language)

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1. If the data contain **no nesting structure**, then read the file directly to a data frame with pd.read_json().

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The trick is getting Python to recognize the data organized in the file, instead of reading the file as one long string.

There are two methods to loading JSON data into Python:

- 1. If the data contain **no nesting structure**, then read the file directly to a data frame with pd.read_json().
- 2. If the data contain nested structures, use <code>json_normalize()</code> from the <code>pandas.io.json</code> module to create a data frame.

1. If the data contain **no nesting structure**, then read the file directly to a data frame with pd.read_json().

```
df = pd.read_json(path_or_buf, orient, typ)
```

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```
df = pd.read_json(path_or_buf, orient, typ)
```

path_or_buf - (str) Either an existing string object with
JSON-formatted data, the filename (if the working directory is
set), the filename and path, or the URL that stores the data

1. If the data contain **no nesting structure**, then read the file directly to a data frame with pd.read_json().

```
df = pd.read_json(path_or_buf, orient, typ)
```

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typ – (str) what kind of output to produce?

- ► A dataframe: typ="frame"
- Or a list of dictionaries: typ="series"



If typ = "series", then pd.read_json() stores the output as a
list of dictionaries - and we can call specific elements of that list.

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list of dictionaries - and we can call specific elements of that list.

This example JSON contains data on 10 users:

We can look at just the first dictionary by calling element 0:

```
data_json[0]
{'id': 1.
 'name': 'Leanne Graham'.
 'username': 'Bret',
 'email': 'Sincere@april.biz',
 'address': {'street': 'Kulas Light',
  'suite': 'Apt. 556',
  'city': 'Gwenborough'.
  'zipcode': '92998-3874',
  'geo': {'lat': '-37.3159', 'lng': '81.1496'}},
 'phone': '1-770-736-8031 x56442',
 'website': 'hildegard.org',
 'company': {'name': 'Romaguera-Crona',
  'catchPhrase': 'Multi-layered client-server neural-net',
  'bs': 'harness real-time e-markets'}}
```

Or we can look at elements within this dictionary by calling the name of the element we want:

```
data_json[0]['address']
{'street': 'Kulas Light',
   'suite': 'Apt. 556',
   'city': 'Gwenborough',
   'zipcode': '92998-3874',
   'geo': {'lat': '-37.3159', 'lng': '81.1496'}}
```

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 'city': 'Gwenborough',
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 'geo': {'lat': '-37.3159', 'lng': '81.1496'}}
data_json[0]['address']['city']
'Gwenborough'
```

orient – (str) how is the JSON data organized? There are five options: records, columns, split, index, and values.

orient="records" works with JSON files organized as a list-of-sets, where each set is an entire record (or a row in flat data):

```
'[{"caseid":1.0,"fttrump":1.0,"fthrc":76.0,"birthyr":1960,"gender":1}, {"caseid":2.0,"fttrump":28.0,"fthrc":52.0,"birthyr":1957,"gender":2}, {"caseid":3.0,"fttrump":100.0,"fthrc":1.0,"birthyr":1963,"gender":1}, {"caseid":4.0,"fttrump":0.0,"fthrc":69.0,"birthyr":1980,"gender":1}, {"caseid":5.0,"fttrump":13.0,"fthrc":1.0,"birthyr":1974,"gender":1}]'
```

orient – (str) how is the JSON data organized? There are five options: records, columns, split, index, and values.

orient="columns" works with JSON files organized as a list-of-sets, where each set is an entire column (the names are the row-names in the flat data):

```
'{"caseid":{"0":1.0,"1":2.0,"2":3.0,"3":4.0,"4":5.0},
"fttrump":{"0":1.0,"1":28.0,"2":100.0,"3":0.0,"4":13.0},
"fthrc":{"0":76.0,"1":52.0,"2":1.0,"3":69.0,"4":1.0},
"birthyr":{"0":1960,"1":1957,"2":1963,"3":1980,"4":1974},
"gender":{"0":1,"1":2,"2":1,"3":1,"4":1}}'
```

orient – (str) how is the JSON data organized? There are five options: records, columns, split, index, and values.

orient="split" works with JSON files organized as set with three lists: columns lists the column names, index lists the row names, and data is a list-of-lists of data points, one list for each row.

```
'{"columns":["caseid","fttrump","fthrc","birthyr","gender"],
"index":[0,1,2,3,4],
"data":[[1.0,1.0,76.0,1960,1],
        [2.0,28.0,52.0,1957,2],
        [3.0,100.0,1.0,1963,1],
        [4.0,0.0,69.0,1980,1],
        [5.0,13.0,1.0,1974,1]]}'
```

orient – (str) how is the JSON data organized? There are five options: records, columns, split, index, and values.

orient="index" is like orient="records" but includes the name of each row in the data:

```
"("0":{"caseid":1.0,"fttrump":1.0,"fthrc":76.0,"birthyr":1960,"gender":1}, "1":{"caseid":2.0,"fttrump":28.0,"fthrc":52.0,"birthyr":1957,"gender":2}, "2":{"caseid":3.0,"fttrump":100.0,"fthrc":1.0,"birthyr":1963,"gender":1}, "3":{"caseid":4.0,"fttrump":0.0,"fthrc":69.0,"birthyr":1980,"gender":1}, "4":{"caseid":5.0,"fttrump":13.0,"fthrc":1.0,"birthyr":1974,"gender":1}}
```

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orient="index" is like orient="records" but includes the name of each row in the data:

```
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"4":{"caseid":5.0,"fttrump":13.0,"fthrc":1.0,"birthyr":1974,"gender":1}}
```

orient="values" only contains the datapoints:

```
[[1.0,1.0,76.0,1960,1],
[2.0,28.0,52.0,1957,2],
[3.0,100.0,1.0,1963,1],
[4.0,0.0,69.0,1980,1],
[5.0,13.0,1.0,1974,1]]
```

If the JSON data contains different data types for the same variable in different records, such as:

```
[{'caseid': 1.0,
   'fttrump': 'Awful',
   'fthrc': 'Pretty good',
   'birthyr': 1960,
   'gender': 1},
   {'caseid': 2.0,
   'fttrump': 28.0,
   'fthrc': 52.0,
   'birthyr': 1957,
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```

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   'gender': 1},
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   'fttrump': 28.0,
   'fthrc': 52.0,
   'birthyr': 1957,
   'gender': 2}]
```

then the pd.read_json() function stores the variables (fttrump and fthrc in this case) as "object" type data — meaning that it is agnostic about whether the variable contains strings or categories.

If different JSON records contain data on different variables:

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then the pd.read_json() function creates columns for every variable that appears even once, and places NaN values for records that do not address these variables.

	caseid	comment	${ t fttrump}$	turnout	vote
0	1	NaN	1	1	1.0
1	2	NaN	28	0	NaN
2	3	big fan of Trump	100	1	0.0

2. If the data contain nested structures, use <code>json_normalize()</code> from the <code>pandas.io.json</code> module to create a data frame.

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First, import the json_normalize() function:

from pandas.io.json import json_normalize

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First, import the json_normalize() function:

```
from pandas.io.json import json_normalize
```

Second, load the data into Python using pd.read_json() as a list, not a dataframe. For example:

```
url = "https://jsonplaceholder.typicode.com/users"
users = pd.read_json(url, typ="series")
```

If you load the data directly as a data frame using pd.read_json(), and there is nesting in the data, the data frame will keep these structures as strings in the output data:

	address	company	email
0	{'street': 'Kulas Light', 'suite': 'Apt. 556',	{'name': 'Romaguera-Crona', 'catchPhrase': 'Mu	Sincere@april.biz
1	{'street': 'Victor Plains', 'suite': 'Suite 87	{'name': 'Deckow-Crist', 'catchPhrase': 'Proac	Shanna@melissa.tv
2	{'street': 'Douglas Extension', 'suite': 'Suit	{'name': 'Romaguera-Jacobson', 'catchPhrase':	Nathan@yesenia.net
3	{'street': 'Hoeger Mall', 'suite': 'Apt. 692',	{'name': 'Robel-Corkery', 'catchPhrase': 'Mult	Julianne.OConner@kory.org
4	{'street': 'Skiles Walks', 'suite': 'Suite 351	{'name': 'Keebler LLC', 'catchPhrase': 'User-c	Lucio_Hettinger@annie.ca
5	{'street': 'Norberto Crossing', 'suite': 'Apt	{'name': 'Considine-Lockman', 'catchPhrase': '	Karley_Dach@jasper.info
6	('street': 'Rex Trail', 'suite': 'Suite 280',	{'name': 'Johns Group', 'catchPhrase': 'Config	Telly.Hoeger@billy.biz
7	{'street': 'Ellsworth Summit', 'suite': 'Suite	{'name': 'Abernathy Group', 'catchPhrase': 'Im	Sherwood@rosamond.me
8	('street': 'Dayna Park', 'suite': 'Suite 449',	{'name': 'Yost and Sons', 'catchPhrase': 'Swit	Chaim_McDermott@dana.io
9	{'street': 'Kattie Turnpike', 'suite': 'Suite	{'name': 'Hoeger LLC', 'catchPhrase': 'Central	Rey.Padberg@karina.biz

This makes it hard to access the data inside the address or company columns, for example.



Instead, use json_normalize() like this:

```
url = "https://jsonplaceholder.typicode.com/users"
users = pd.read_json(url, typ="series")
users = json_normalize(users)
users
```

	address.city	address.geo.lat	address.geo.lng	address.street	address.suite	address.zipcode	company.bs	company.catchPhrase	company.name
0	Gwenborough	-37.3159	81.1496	Kulas Light	Apt. 556	92998-3874	harness real-time e-markets	Multi-layered client- server neural-net	Romaguera- Crona
1	Wisokyburgh	-43.9509	-34.4618	Victor Plains	Suite 879	90566-7771	synergize scalable supply- chains	Proactive didactic contingency	Deckow-Crist
2	McKenziehaven	-68.6102	-47.0653	Douglas Extension	Suite 847	59590-4157	e-enable strategic applications	Face to face bifurcated interface	Romaguera- Jacobson
3	South Elvis	29.4572	-164.2990	Hoeger Mall	Apt. 692	53919-4257	transition cutting- edge web services	Multi-tiered zero tolerance productivity	Robel-Corkery
4	Roscoeview	-31.8129	62.5342	Skiles Walks	Suite 351	33263	revolutionize end-to-end systems	User-centric fault- tolerant solution	Keebler LLC

The column names are fairly ugly, but every variable is now stored in a separate column.



To convert a dataframe to a JSON file, use: df2 = df.to_json(orient)

Replace df with the name of the dataframe object you are converting, and replace df2 with object you are creating.

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orient - (str) Just like with pd.read_json(), the orient
parameter can be set to

orient="records" - a list of sets, each one referring to one record

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- orient="columns" a list of sets, each one referring to
 one column
- orient="split" three lists, one for column names, one for row names, one for data
- ▶ orient="index" like "records" but including row names
- orient="values" just data in a list-of-lists



```
To write a JSON file to disk, use: df.to_json(path_or_buf, orient)
```

Replace df with the object you want to save as a JSON-encoded text file. If df is a dataframe, then this function both converts it to JSON and writes it to disk.

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orient – (str) Works the same as with reading JSON, or converting a dataframe to JSON.