# 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

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- 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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- actively toxic behavior is usually explicitly banned by codes of conduct,
- ▶ 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)

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

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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- What is the expected output? What do you see instead?
- ▶ 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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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

The main slack page for the global community of Python users is **Pyslackers**: https://pyslackers.com/web

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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

Usenet – a distributed discussion system (no central server) – was invented in 1979, and is still in use today. The Python Usenet message boards are at <a href="https://mail.python.org/archives">https://mail.python.org/archives</a>. The comp.lang.python board is for general discussions and questions about Python.

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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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What follows is a set of *guidelines and suggestions*. NOT a definitive list of how to do things.

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.



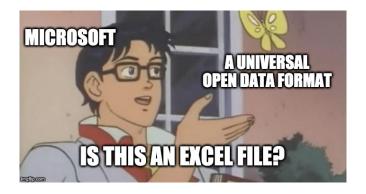
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A comma-separated values (CSV) file:

<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:

import pandas as pd

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

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I put several versions of the ANES data on our class GitHub page.

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I put several versions of the ANES data on our class GitHub page.

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")
```

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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)
```

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- If you catch anything weird, return to 1. and try different parameters for pd.read\_csv()

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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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	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												
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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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```
pd.set_option('display.max_columns', None)
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To always display all of the rows, type

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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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```

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

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

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

```
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.

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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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- ► 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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- ► The default is header=0, which uses the first row as variable names
- 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<sub>th</sub> row for variable names, and deletes all higher rows

```
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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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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.

pd.read\_csv(filepath\_or\_buffer, sep, header, usecols, skiprows, skipfooter, nrows, na\_values)

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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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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.

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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.

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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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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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.

```
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.

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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>.

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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',
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There are two ways to proceed next:

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  '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

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

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

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.

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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io - (string) This argument is the same as the
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- 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:

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

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

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```
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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 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")
```

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")
```

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")
```