

# 5 Data Cleaning and EDA

Code ▾

▶ Code

## Learning Outcomes

- Recognize common file formats
- Categorize data by its variable type
- Build awareness of issues with data faithfulness and develop targeted solutions

In the past few lectures, we've learned that `pandas` is a toolkit to restructure, modify, and explore a dataset. What we haven't yet touched on is *how* to make these data transformation decisions. When we receive a new set of data from the "real world," how do we know what processing we should do to convert this data into a usable form?

**Data cleaning**, also called **data wrangling**, is the process of transforming raw data to facilitate subsequent analysis. It is often used to address issues like:

- Unclear structure or formatting
- Missing or corrupted values
- Unit conversions
- ...and so on

**Exploratory Data Analysis (EDA)** is the process of understanding a new dataset. It is an open-ended, informal analysis that involves familiarizing ourselves with the variables present in the data, discovering potential hypotheses, and identifying possible issues with the data. This last point can often motivate further data cleaning to address any problems with the dataset's format; because of this, EDA and data cleaning are often thought of as an "infinite loop," with each process driving the other.

In this lecture, we will consider the key properties of data to consider when performing data cleaning and EDA. In doing so, we'll develop a "checklist" of sorts for you to consider when approaching a new dataset. Throughout this process, we'll build a deeper understanding of this early (but very important!) stage of the data science lifecycle.

## 5.1 Structure

We often prefer rectangular data for data analysis. Rectangular structures are easy to manipulate and analyze. A key element of data cleaning is about transforming data to be more rectangular.

There are two kinds of rectangular data: tables and matrices. Tables have named columns with different data types and are manipulated using data transformation languages. Matrices contain numeric data of the same type and are manipulated using linear algebra.

### 5.1.1 File Formats

There are many file types for storing structured data: TSV, JSON, XML, ASCII, SAS, etc. We'll only cover CSV, TSV, and JSON in lecture, but you'll likely encounter other formats as you work with different datasets. Reading documentation is your best bet for understanding how to process the multitude of different file types.

### 5.1.1.1 CSV

CSVs, which stand for **Comma-Separated Values**, are a common tabular data format. In the past two **pandas** lectures, we briefly touched on the idea of file format: the way data is encoded in a file for storage. Specifically, our **elections** and **babynames** datasets were stored and loaded as CSVs:

```
pd.read_csv("data/elections.csv").head(5)
```

Year	Candidate	Party	Popular vote	Result	%
0	1824	Andrew Jackson	Democratic-Republican	151271	loss
1	1824	John Quincy Adams	Democratic-Republican	113142	win
2	1828	Andrew Jackson	Democratic	642806	win
3	1828	John Quincy Adams	National Republican	500897	loss
4	1832	Andrew Jackson	Democratic	702735	win

To better understand the properties of a CSV, let's take a look at the first few rows of the raw data file to see what it looks like before being loaded into a **DataFrame**. We'll use the **repr()** function to return the raw string with its special characters:

```
with open("data/elections.csv", "r") as table:
    i = 0
    for row in table:
        print(repr(row))
        i += 1
        if i > 3:
            break
```

```
'Year,Candidate,Party,Popular vote,Result,%
'1824,Andrew Jackson,Democratic–Republican,151271,loss,57.21012204%
'1824,John Quincy Adams,Democratic–Republican,113142,win,42.78987796%
'1828,Andrew Jackson,Democratic,642806,win,56.20392707%
```

Each row, or **record**, in the data is delimited by a newline **\n**. Each column, or **field**, in the data is delimited by a comma **,** (hence, comma-separated!).

### 5.1.1.2 TSV

Another common file type is **TSV (Tab-Separated Values)**. In a TSV, records are still delimited by a newline **\n**, while fields are delimited by **\t** tab character.

Let's check out the first few rows of the raw TSV file. Again, we'll use the **repr()** function so that **print** shows the special characters.

```
with open("data/elections.txt", "r") as table:
    i = 0
    for row in table:
        print(repr(row))
        i += 1
        if i > 3:
            break
```

```
'\ufe0fYear\tCandidate\tParty\tPopular vote\tResult\t%\n'
'1824\tAndrew Jackson\tDemocratic-Republican\t151271\tloss\t57.21012204\n'
'1824\tJohn Quincy Adams\tDemocratic-Republican\t113142\twin\t42.78987796\n'
'1828\tAndrew Jackson\tDemocratic\t642806\twin\t56.20392707\n'
```

TSVs can be loaded into `pandas` using `pd.read_csv`. We'll need to specify the **delimiter** with parameter `sep='\t'` ([documentation](#)).

```
pd.read_csv("data/elections.txt", sep='\t').head(3)
```

Year	Candidate	Party	Popular vote	Result	%
0 1824	Andrew Jackson	Democratic-Republican	151271	loss	57.21
1 1824	John Quincy Adams	Democratic-Republican	113142	win	42.79
2 1828	Andrew Jackson	Democratic	642806	win	56.20

An issue with CSVs and TSVs comes up whenever there are commas or tabs within the records. How does `pandas` differentiate between a comma delimiter vs. a comma within the field itself, for example `8,900`? To remedy this, check out the [`quotechar` parameter](#).

### 5.1.1.3 JSON

**JSON (JavaScript Object Notation)** files behave similarly to Python dictionaries. A raw JSON is shown below.

```
with open("data/elections.json", "r") as table:
    i = 0
    for row in table:
        print(row)
        i += 1
        if i > 8:
            break
```

```
[
```

```
{
```

```
"Year": 1824,
"Candidate": "Andrew Jackson",
```

```
"Party": "Democratic-Republican",
```

```
"Popular vote": 151271,
```

```
"Result": "loss",
```

```
"%": 57.21012204
```

```
},
```

JSON files can be loaded into `pandas` using `pd.read_json`.

```
pd.read_json('data/elections.json').head(3)
```

Year	Candidate	Party	Popular vote	Result	%
0	1824	Andrew Jackson	Democratic-Republican	151271	loss
1	1824	John Quincy Adams	Democratic-Republican	113142	win
2	1828	Andrew Jackson	Democratic	642806	win

### 5.1.1.3.1 EDA with JSON: United States Congress Data

The congress.gov API (Application Programming Interface) provides data about the activities and members of the United States Congress (i.e., the House of Representatives and the Senate).

To get a JSON file containing information about the current members of Congress from California, you could use the following **API call**:

- `https://api.congress.gov/v3/member/CA?api_key=[INSERT_KEY]&limit=250&format=json&currentMember=True`
- You can instantly sign up for a congress.gov **API key** [here](#). Once you have your key, replace `[INSERT_KEY]` above with your key, and enter the API call as a URL in your browser. What happens?
- Once the JSON from the API call is visible in your browser, you can click `File` → `Save Page As` to save the JSON file to your computer.
- Coarsely, API keys are used to track how much a given user engages with the API. There might be limits to the number of API calls (e.g., congress.gov API limits to 5,000 calls per hour), or a cost for API calls (e.g., using the OpenAI API for programmatically using ChatGPT).

After following these steps, we save this JSON file as `data/ca-congress-members.json`.

#### 5.1.1.3.1.1 File Contents

Let's examine this file using Python. We can programmatically view the first couple lines of the file using the same functions we used with CSVs:

```
congress_file = "data/ca-congress-members.json"

# Inspect the first five lines of the file
with open(congress_file, "r") as f:
    for i, row in enumerate(f):
        print(row)
        if i >= 4: break
```

{

```
"members": [
    {
        "bioguideId": "T000491",
        "depiction": {
```

#### 5.1.1.3.1.2 EDA: Digging into JSON with Python

JSON data closely matches the internal Python object model.

- In the following cell, we import the entire JSON datafile into a Python dictionary using the `json` package.

```
import json

# Import the JSON file into Python as a dictionary
with open(congress_file, "rb") as f:
    congress_json = json.load(f)

type(congress_json)
```

dict

The `congress_json` variable is a dictionary encoding the data in the JSON file.

Below, we access the first element of the `members` element of the `congress_json` dictionary.

- This first element is also a dictionary (and there are more dictionaries inside of it!)

```
# Grab the list corresponding to the `members` key in the JSON dictionary,
# and then grab the first element of this list.
# In a moment, we'll see how we knew to use the key `members`, and that
# the resulting object is a list.
congress_json['members'][0]
```

```
{'bioguideId': 'T000491',
'depiction': {'attribution': 'Image courtesy of the Member',
'imageUrl': 'https://www.congress.gov/img/member/6774606d0b34857ecc9091a9_200.jpg'},
'district': 45,
'name': 'Tran, Derek',
```

```
'partyName': 'Democratic',
'state': 'California',
'terms': {'item': [{'chamber': 'House of Representatives',
  'startYear': 2025}]},
'updateDate': '2025-01-21T18:00:52Z',
'url': 'https://api.congress.gov/v3/member/T000491?format=json'}
```

How should we probe a nested dictionary like `congress_json`?

We can start by identifying the top-level **keys** of the dictionary:

```
# Grab the top-level keys of the JSON dictionary
congress_json.keys()
```

```
dict_keys(['members', 'pagination', 'request'])
```

Looks like we have three top-level keys: `members`, `pagination`, and `request`.

You'll often see a top-level `meta` key in JSON files. This does not refer to Meta (formerly Facebook). Instead, it typically refers to metadata (data about the data). Metadata are often maintained alongside the data.

Let's check the type of the `members` element:

```
type(congress_json['members'])
```

```
list
```

Looks like a list! What are the first two elements?

```
congress_json['members'][:2]
```

```
[{'bioguideId': 'T000491',
'depiction': {'attribution': 'Image courtesy of the Member',
'imageUrl': 'https://www.congress.gov/img/member/6774606d0b34857ecc9091a9_200.jpg'},
'district': 45,
'name': 'Tran, Derek',
'partyName': 'Democratic',
'state': 'California',
'terms': {'item': [{'chamber': 'House of Representatives',
  'startYear': 2025}]},
'updateDate': '2025-01-21T18:00:52Z',
'url': 'https://api.congress.gov/v3/member/T000491?format=json'},
{'bioguideId': 'M001241',
'depiction': {'attribution': 'Image courtesy of the Member',
'imageUrl': 'https://www.congress.gov/img/member/67744ed90b34857ecc909155_200.jpg'},
'district': 47,
'name': 'Min, Dave',
'partyName': 'Democratic',
```

```
'state': 'California',
'terms': {'item': [{'chamber': 'House of Representatives',
  'startYear': 2025}]},
'updateDate': '2025-01-21T18:00:52Z',
'url': 'https://api.congress.gov/v3/member/M001241?format=json'}
```

More dictionaries! You can repeat the process above to traverse the nested dictionary.

You'll notice that each record of `congress_json['members']` looks like it could be a column of a DataFrame.

- The keys look a lot like column names, and the values could be the entries in each row.

But, the two other elements of `congress_json` don't have the same structure as `congress_json['members']`.

- So, they probably don't belong in a DataFrame containing the members of Congress from CA.
  - We'll see the implications of this inconsistency in the next section.

```
print(congress_json['pagination'])  
print(congress_json['request'])
```

```
{'count': 54}  
{'contentType': 'application/json', 'format': 'json'}
```

### 5.1.1.3.1.3 JSON with pandas

`pandas` has a built in function called `pd.read_json` for reading in JSON files. In order to read in this JSON file, you might want to try something like the code in the cell below. However, if we tried to run this code, it would error.

```
#pd.read_json(congress_file)
```

- The code above tries to import the entire JSON file located at `congress_file` (`congress_json`), including `congress_json['pagination']` and `congress_json['request']`.
  - We only want to make a DataFrame out of `congress_json['members']`.

Instead, let's try converting the `members` element of `congress_json` to a DataFrame by using `pd.DataFrame`:

```
# Convert dictionary to DataFrame
congress_df = pd.DataFrame(congress_json['members'])
congress_df.head()
```

bioguideId	depiction	district	name	partyName	state	terms	updateDate	url
0 T000491	{'attribution': 'Image courtesy of', 'url': 'https://api.congress.gov/116/bioguide/T000491/photo'} Tran, Derek			Democratic	California	{'item': ['House of Representatives'], 'start': '2021-01-06', 'end': '2025-01-03'}	2025-01-06T18:00:52Z	https://api.congress.gov/116/bioguide/T000491

bioguideId	depiction	district	name	partyName	state	terms	updateDate	url
	the Member'...							
1 M001241	{'attribution': 47.00 'Image courtesy of the Member'...	Min, Dave	Democratic	California	{'item': [['chamber': 'House of Representative...}}	2025-01- 21T18:00:52Z	for...	https://api.congress.g
2 K000400	{'attribution': 37.00 'Image courtesy of the Member'...	Kamlager- Dove, Sydney	Democratic	California	{'item': [['chamber': 'House of Representative...}}	2025-01- 21T18:00:52Z	for...	https://api.congress.g
3 G000598	{'attribution': 42.00 'Image courtesy of the Member'...	Garcia, Robert	Democratic	California	{'item': [['chamber': 'House of Representative...}}	2025-01- 21T18:00:52Z	for...	https://api.congress.g
4 K000397	{'attribution': 40.00 'Image courtesy of the Member'...	Kim, Young	Republican	California	{'item': [['chamber': 'House of Representative...}}	2025-01- 21T18:00:52Z	for...	https://api.congress.g

We've successfully begun to rectangularize our JSON data!

## 5.1.2 Other Data Formats

So far, we've looked at text data that comes in a quite nice format. Although some data cleaning might be necessary, it has still had all of the components of a rectangular dataset. However, not all data comes like this, and there are also different kinds of data we can use. Some examples include:

- **Image Data:** Used for medical diagnosis
- **Audio Data:** Used for speech recognition, sentiment analysis
- **Video Data:** Used for object tracking, facial recognition
- **Text:** Used for LLMs, document review

Even though this may not look tabular at first, all of these formats can be represented in tabular/matrix form! So by learning how to work with tabular data, you are well equipped to deal with other kinds of data as well.

## 5.1.3 Variable Types

Variables are columns. A variable is a measurement of a particular concept. Variables have two common properties: data type/storage type and variable type/feature type. The data type of a variable indicates how each variable value is stored in memory (integer, floating point, boolean, etc.) and affects which `pandas` functions are used. The variable type is a conceptualized measurement of information (and therefore indicates what values a variable can take on). Variable type is identified through expert knowledge, exploring the data itself, or consulting

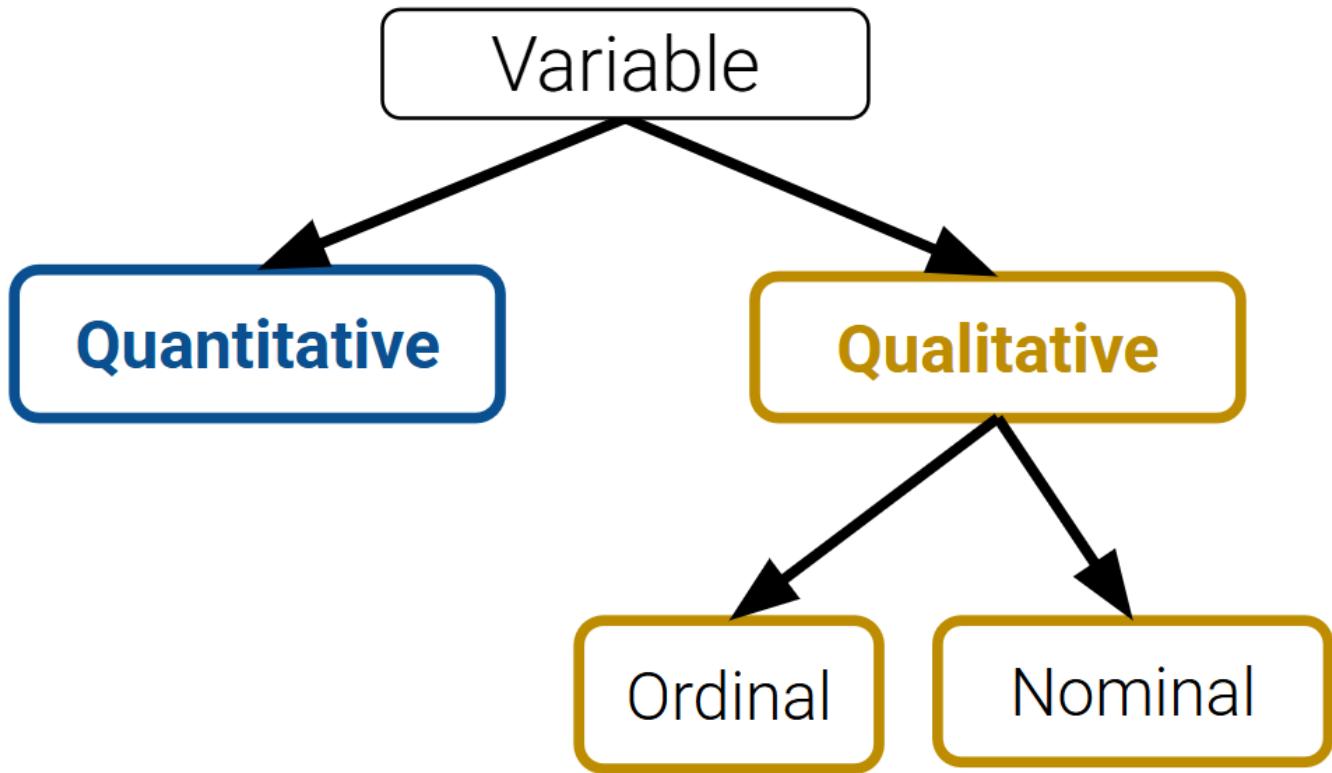
the data codebook. The variable type affects how one visualizes and interprets the data. In this class, “variable types” are conceptual.

After loading data into a file, it’s a good idea to take the time to understand what pieces of information are encoded in the dataset. In particular, we want to identify what variable types are present in our data. Broadly speaking, we can categorize variables into one of two overarching types.

**Quantitative variables** describe some numeric quantity or amount. Some examples include weights, GPA, CO<sub>2</sub> concentrations, someone’s age, or the number of siblings they have.

**Qualitative variables**, also known as **categorical variables**, describe data that isn’t measuring some quantity or amount. The sub-categories of categorical data are:

- **Ordinal qualitative variables**: categories with ordered levels. Specifically, ordinal variables are those where the difference between levels has no consistent, quantifiable meaning. Some examples include levels of education (high school, undergrad, grad, etc.), income bracket (low, medium, high), or Yelp rating.
- **Nominal qualitative variables**: categories with no specific order. For example, someone’s political affiliation or Cal ID number.



Classification of variable types

Note that many variables don’t sit neatly in just one of these categories. Qualitative variables could have numeric levels, and conversely, quantitative variables could be stored as strings.

## 5.2 Granularity and Temporality

After understanding the structure of the dataset, the next task is to determine what exactly the data represents. We'll do so by considering the data's granularity and temporality.

## 5.2.1 Granularity

The **granularity** of a dataset is what a single row represents. You can also think of it as the level of detail included in the data. To determine the data's granularity, ask: what does each row in the dataset represent? Fine-grained data contains a high level of detail, with a single row representing a small individual unit. For example, each record may represent one person. Coarse-grained data is encoded such that a single row represents a large individual unit – for example, each record may represent a group of people.

## 5.2.2 Temporality

The **temporality** of a dataset describes the periodicity over which the data was collected as well as when the data was most recently collected or updated.

Time and date fields of a dataset could represent a few things:

1. when the “event” happened
2. when the data was collected, or when it was entered into the system
3. when the data was copied into the database

To fully understand the temporality of the data, it also may be necessary to standardize time zones or inspect recurring time-based trends in the data (do patterns recur in 24-hour periods? Over the course of a month? Seasonally?). The convention for standardizing time is the Coordinated Universal Time (UTC), an international time standard measured at 0 degrees latitude that stays consistent throughout the year (no daylight savings). We can represent Berkeley's time zone, Pacific Standard Time (PST), as UTC-7 (with daylight savings).

### 5.2.2.1 Temporality with pandas' dt accessors

Let's briefly look at how we can use `pandas' dt` accessors to work with dates/times in a dataset using the dataset you'll see in Lab 3: the Berkeley PD Calls for Service dataset.

► Code

	CASENO	OFFENSE	EVENTDT	EVENTTM	CVLEGEND	CVDOW	InDbDate	Block_Location	BLKADDR	City	State
0	21014296	THEFT MISD. (UNDER \$950)	04/01/2021 12:00:00	10:58 AM	LARCENY	4	06/15/2021 12:00:00 AM	Berkeley, CA\n(37.869058, -122.270455)	NaN	Berkeley	CA
1	21014391	THEFT MISD. (UNDER \$950)	04/01/2021 12:00:00	10:38 AM	LARCENY	4	06/15/2021 12:00:00 AM	Berkeley, CA\n(37.869058, -122.270455)	NaN	Berkeley	CA

CASENO	OFFENSE	EVENTDT	EVENTTM	CVLEGEND	CVDOW	InDbDate	Block_Location	BLKADDR	City	State
2	21090494	THEFT MISD. (UNDER \$950)	04/19/2021 12:00:00 AM	LARCENY	1	06/15/2021 12:00:00 AM	2100 BLOCK HASTE ST\nBerkeley, CA\n(37.864908,...	2100	Berkeley	CA
3	21090204	THEFT FELONY (OVER \$950)	02/13/2021 12:00:00 AM	LARCENY	6	06/15/2021 12:00:00 AM	2600 BLOCK WARRING ST\nBerkeley, CA\n(37.86393...	2600	Berkeley	CA
4	21090179	BURGLARY AUTO	02/08/2021 12:00:00 AM	BURGLARY - VEHICLE	1	06/15/2021 12:00:00 AM	2700 BLOCK GARBER ST\nBerkeley, CA\n(37.86066,...	2700	Berkeley	CA

Looks like there are three columns with dates/times: `EVENTDT`, `EVENTTM`, and `InDbDate`.

Most likely, `EVENTDT` stands for the date when the event took place, `EVENTTM` stands for the time of day the event took place (in 24-hr format), and `InDbDate` is the date this call is recorded onto the database.

If we check the data type of these columns, we will see they are stored as strings. We can convert them to `datetime` objects using pandas `to_datetime` function.

```
calls["EVENTDT"] = pd.to_datetime(calls["EVENTDT"])
calls.head()
```

CASENO	OFFENSE	EVENTDT	EVENTTM	CVLEGEND	CVDOW	InDbDate	Block_Location	BLKADDR	City	State	
0	21014296	THEFT MISD. (UNDER \$950)	2021-04-01	10:58	LARCENY	4	06/15/2021 12:00:00 AM	Berkeley, CA\n(37.869058, -122.270455)	NaN	Berkeley	CA
1	21014391	THEFT MISD. (UNDER \$950)	2021-04-01	10:38	LARCENY	4	06/15/2021 12:00:00 AM	Berkeley, CA\n(37.869058, -122.270455)	NaN	Berkeley	CA
2	21090494	THEFT MISD. (UNDER \$950)	2021-04-19	12:15	LARCENY	1	06/15/2021 12:00:00 AM	2100 BLOCK HASTE ST\nBerkeley, CA\n(37.864908,...	2100	Berkeley	CA
3	21090204	THEFT FELONY (OVER \$950)	2021-02-13	17:00	LARCENY	6	06/15/2021 12:00:00 AM	2600 BLOCK WARRING ST\nBerkeley, CA\n(37.86393...	2600	Berkeley	CA
4	21090179	BURGLARY AUTO	2021-02-08	6:20	BURGLARY - VEHICLE	1	06/15/2021 12:00:00 AM	2700 BLOCK GARBER ST\nBerkeley, CA\n(37.86066,...	2700	Berkeley	CA

Now, we can use the `dt` accessor on this column.

We can get the month:

```
calls["EVENTDT"].dt.month.head()
```

```
0    4
1    4
2    4
3    2
4    2
Name: EVENTDT, dtype: int32
```

Which day of the week the date is on:

```
calls["EVENTDT"].dt.dayofweek.head()
```

```
0    3
1    3
2    0
3    5
4    0
Name: EVENTDT, dtype: int32
```

Check the minimum values to see if there are any suspicious-looking, 70s dates:

```
calls.sort_values("EVENTDT").head()
```

CASENO	OFFENSE	EVENTDT	EVENTTM	CVLEGEND	CVDOW	InDbDate	Block_Location	BLKADDR	Ci
2513	20057398 BURGLARY COMMERCIAL	2020-12-17	16:05	BURGLARY - COMMERCIAL	4	06/15/2021	600 BLOCK	600	B
					AM	12:00:00	GILMAN	BLOCK	
						ST\nBerkeley,	GILMAN ST		
						CA\n(37.878405,...			
624	20057207 ASSAULT/BATTERY MISD.	2020-12-17	16:50	ASSAULT	4	06/15/2021	2100 BLOCK	2100	B
					AM	12:00:00	SHATTUCK	BLOCK	
						AVE\nBerkeley,	SHATTUCK		
						CA\n(37.871...	AVE		
154	20092214 THEFT FROM AUTO	2020-12-17	18:30	LARCENY - FROM VEHICLE	4	06/15/2021	800 BLOCK	800	B
					AM	12:00:00	SHATTUCK	BLOCK	
						AVE\nBerkeley,	SHATTUCK		
						CA\n(37.8918...	AVE		
659	20057324 THEFT MISD. (UNDER \$950)	2020-12-17	15:44	LARCENY	4	06/15/2021	1800 BLOCK 4TH	1800	B
					AM	12:00:00	ST\nBerkeley,	BLOCK	
						CA\n(37.869888,	4TH ST		
						-...			

CASENO	OFFENSE	EVENTDT	EVENTTM	CVLEGEND	CVDOW	InDbDate	Block_Location	BLKADDR	Ci
993	20057573 BURGLARY RESIDENTIAL	2020-12-17	22:15	BURGLARY - RESIDENTIAL	4	06/15/2021 12:00:00 AM	1700 BLOCK ST\nBerkeley, CA\n(37.857495...	1700 BLOCK STUART	Be

Doesn't look like it! We are good!

We can also do many things with the `dt` accessor like switching time zones and converting time back to UNIX/POSIX time. Check out the documentation on [.dt accessor](#) and [time series/date functionality](#).

## 5.3 Faithfulness

At this stage in our data cleaning and EDA workflow, we've achieved quite a lot: we've identified how our data is structured, come to terms with what information it encodes, and gained insight as to how it was generated. Throughout this process, we should always recall the original intent of our work in Data Science – to use data to better understand and model the real world. To achieve this goal, we need to ensure that the data we use is faithful to reality; that is, that our data accurately captures the “real world.”

Data used in research or industry is often “messy” – there may be errors or inaccuracies that impact the faithfulness of the dataset. Signs that data may not be faithful include:

- Unrealistic or “incorrect” values, such as negative counts, locations that don’t exist, or dates set in the future
- Violations of obvious dependencies, like an age that does not match a birthday
- Clear signs that data was entered by hand, which can lead to spelling errors or fields that are incorrectly shifted
- Signs of data falsification, such as fake email addresses or repeated use of the same names
- Duplicated records or fields containing the same information
- Truncated data, e.g. Microsoft Excel would limit the number of rows to 65536 and the number of columns to 255

We often solve some of these more common issues in the following ways:

- Spelling errors: apply corrections or drop records that aren’t in a dictionary
- Time zone inconsistencies: convert to a common time zone (e.g. UTC)
- Duplicated records or fields: identify and eliminate duplicates (using primary keys)
- Unspecified or inconsistent units: infer the units and check that values are in reasonable ranges in the data

### 5.3.1 Missing Values

Another common issue encountered with real-world datasets is that of missing data. One strategy to resolve this is to simply drop any records with missing values from the dataset. This does, however, introduce the risk of inducing biases – it is possible that the missing or corrupt records may be systemically related to some feature of interest in the data. Another solution is to keep the data as `NaN` values.

A third method to address missing data is to perform **imputation**: infer the missing values using other data available in the dataset. There is a wide variety of imputation techniques that can be implemented; some of the

most common are listed below.

- Average imputation: replace missing values with the average value for that field
- Hot deck imputation: replace missing values with some random value
- Regression imputation: develop a model to predict missing values and replace with the predicted value from the model.
- Multiple imputation: replace missing values with multiple random values

Regardless of the strategy used to deal with missing data, we should think carefully about *why* particular records or fields may be missing – this can help inform whether or not the absence of these values is significant or meaningful.

## 5.4 EDA Demo 1: Tuberculosis in the United States

---

Now, let's walk through the data-cleaning and EDA workflow to see what we can learn about the presence of Tuberculosis in the United States!

We will examine the data included in the [original CDC article](#) published in 2021.

### 5.4.1 CSVs and Field Names

Suppose Table 1 was saved as a CSV file located in `data/cdc_tuberculosis.csv`.

We can then explore the CSV (which is a text file, and does not contain binary-encoded data) in many ways: 1. Using a text editor like emacs, vim, VSCode, etc. 2. Opening the CSV directly in DataHub (read-only), Excel, Google Sheets, etc. 3. The `Python` file object 4. `pandas`, using `pd.read_csv()`

To try out options 1 and 2, you can view or download the Tuberculosis from the [lecture demo notebook](#) under the `data` folder in the left hand menu. Notice how the CSV file is a type of **rectangular data (i.e., tabular data) stored as comma-separated values**.

Next, let's try out option 3 using the `Python` file object. We'll look at the first four lines:

► Code

```
,No. of TB cases,,
```

```
U.S. jurisdiction,2019,2020,2021
```

```
Total,"8,900","7,173","7,860"
```

```
Alabama,87,72,92
```

Whoa, why are there blank lines interspersed between the lines of the CSV?

You may recall that all line breaks in text files are encoded as the special newline character `\n`. Python's `print()` prints each string (including the newline), and an additional newline on top of that.

If you're curious, we can use the `repr()` function to return the raw string with all special characters:

#### ► Code

```
' ,No. of TB cases,,\n'
'U.S. jurisdiction,2019,2020,2021\n'
'Total,"8,900","7,173","7,860"\n'
'Alabama,87,72,92\n'
```

Finally, let's try option 4 and use the tried-and-true Data 100 approach: `pandas`.

```
tb_df = pd.read_csv("data/cdc_tuberculosis.csv")
tb_df.head()
```

	Unnamed: 0	No. of TB cases	Unnamed: 2	Unnamed: 3
0	U.S. jurisdiction	2019	2020	2021
1	Total	8,900	7,173	7,860
2	Alabama	87	72	92
3	Alaska	58	58	58
4	Arizona	183	136	129

You may notice some strange things about this table: what's up with the “Unnamed” column names and the first row?

Congratulations — you're ready to wrangle your data! Because of how things are stored, we'll need to clean the data a bit to name our columns better.

A reasonable first step is to identify the row with the right header. The `pd.read_csv()` function ([documentation](#)) has the convenient `header` parameter that we can set to use the elements in row 1 as the appropriate columns:

```
tb_df = pd.read_csv("data/cdc_tuberculosis.csv", header=1) # row index
tb_df.head(5)
```

	U.S. jurisdiction	2019	2020	2021
0	Total	8,900	7,173	7,860
1	Alabama	87	72	92
2	Alaska	58	58	58
3	Arizona	183	136	129
4	Arkansas	64	59	69

## 5.4.2 Record Granularity

You might already be wondering: what's up with that first record?

Row 0 is what we call a **rollup record**, or summary record. It's often useful when displaying tables to humans. The **granularity** of record 0 (Totals) vs the rest of the records (States) is different.

Okay, EDA step two. How was the rollup record aggregated?

Let's check if Total TB cases is the sum of all state TB cases. If we sum over all rows, we should get **2x** the total cases in each of our TB cases by year (why do you think this is?).

► Code

```
U.S. jurisdiction      TotalAlabamaAlaskaArizonaArkansasCaliforniaCol...
2019                  8,9008758183642,111666718245583029973261085237...
2020                  7,1737258136591,706525417194122219282169239376...
2021                  7,8609258129691,750585443194992281064255127494...
dtype: object
```

Whoa, what's going on with the TB cases in 2019, 2020, and 2021? Check out the column types:

► Code

```
U.S. jurisdiction      object
2019                  object
2020                  object
2021                  object
dtype: object
```

Since there are commas in the values for TB cases, the numbers are read as the **object** datatype, or **storage type** (close to the **Python** string datatype), so **pandas** is concatenating strings instead of adding integers (recall that Python can “sum”, or concatenate, strings together: `"data" + "100"` evaluates to `"data100"` ).

Fortunately `read_csv` also has a `thousands` parameter ([documentation](#)):

```
# improve readability: chaining method calls with outer parentheses/line breaks
tb_df = (
    pd.read_csv("data/cdc_tuberculosis.csv", header=1, thousands=',')
)
tb_df.head(5)
```

	U.S. jurisdiction	2019	2020	2021
0	Total	8900	7173	7860
1	Alabama	87	72	92
2	Alaska	58	58	58
3	Arizona	183	136	129
4	Arkansas	64	59	69

```
tb_df.sum()
```

```
U.S. jurisdiction    TotalAlabamaAlaskaArizonaArkansasCaliforniaCol...
2019                  17800
2020                  14346
2021                  15720
dtype: object
```

The total TB cases look right. Phew!

Let's just look at the records with **state-level granularity**:

#### ► Code

U.S. jurisdiction		2019	2020	2021
1	Alabama	87	72	92
2	Alaska	58	58	58
3	Arizona	183	136	129
4	Arkansas	64	59	69
5	California	2111	1706	1750

### 5.4.3 Gather Census Data

U.S. Census population estimates [source](#) (2019), [source](#) (2020-2024).

Running the below cells cleans the data. There are a few new methods here:

- `df.convert_dtypes()` ([documentation](#)) conveniently converts all float dtypes into ints and is out of scope for the class.
- `df.drop_na()` ([documentation](#)) will be explained in more detail next time.

#### ► Code

Geographic											
Area	2010	2011	2012	2013	2014	2015	2016	2017	2018	2019	
0 United States	309321666	311556874	313830990	315993715	318301008	320635163	322941311	324985539	326687501	3282	
1 Northeast	55380134	55604223	55775216	55901806	56006011	56034684	56042330	56059240	56046620	5598	
2 Midwest	66974416	67157800	67336743	67560379	67745167	67860583	67987540	68126781	68236628	6832	
3 South	114866680	116006522	117241208	118364400	119624037	120997341	122351760	123542189	124569433	1255	
4 West	72100436	72788329	73477823	74167130	74925793	75742555	76559681	77257329	77834820	7834	

Occasionally, you will want to modify code that you have imported. To reimport those modifications you can either use `python`'s `importlib` library:

```
from importlib import reload
```

```
reload(utils)
```

or use `iPython` magic which will intelligently import code when files change:

```
%load_ext autoreload
%autoreload 2
```

## ► Code

	<b>Geographic Area</b>	<b>2020</b>	<b>2021</b>	<b>2022</b>	<b>2023</b>	<b>2024</b>
<b>0</b>	United States	331577720	332099760	334017321	336806231	340110988
<b>1</b>	Northeast	57431458	57252533	57159597	57398303	57832935
<b>2</b>	Midwest	68984258	68872831	68903297	69186401	69596584
<b>3</b>	South	126476549	127368010	129037849	130893358	132665693
<b>4</b>	West	78685455	78606386	78916578	79328169	80015776

## 5.4.4 Joining Data (Merging DataFrame s)

Time to `merge`! Here we use the `DataFrame` method `df1.merge(right=df2, ...)` on `DataFrame df1` ([documentation](#)). Contrast this with the function `pd.merge(left=df1, right=df2, ...)` ([documentation](#)). Feel free to use either.

```
# merge TB DataFrame with two US census DataFrames
tb_census_df = (
    tb_df
    .merge(right=census_2010s_df,
          left_on="U.S. jurisdiction", right_on="Geographic Area")
    .merge(right=census_2020s_df,
          left_on="U.S. jurisdiction", right_on="Geographic Area")
)
tb_census_df.head(5)
```

U.S. jurisdiction	Geographic										
	2019_x	2020_x	2021_x	Area_x	2010	2011	2012	2013	2014	2015	2016
<b>0</b> Alabama	87	72	92	Alabama	4785437	4799069	4815588	4830081	4841799	4852347	4863525
<b>1</b> Alaska	58	58	58	Alaska	713910	722128	730443	737068	736283	737498	741456
<b>2</b> Arizona	183	136	129	Arizona	6407172	6472643	6554978	6632764	6730413	6829676	6941072
<b>3</b> Arkansas	64	59	69	Arkansas	2921964	2940667	2952164	2959400	2967392	2978048	2989918
<b>4</b> California	2111	1706	1750	California	37319502	37638369	37948800	38260787	38596972	38918045	3916711

We're only interested in the population for the years 2019, 2020, and 2021, so let's select just those columns:

```
census_2019_df = census_2010s_df[['Geographic Area', '2019']]
census_2020_2021_df = census_2020s_df[['Geographic Area', '2020', '2021']]
```

```
display(tb_df.tail(2))
display(census_2019_df.tail(2))
display(census_2020_2021_df.tail(2))
```

	<b>U.S. jurisdiction</b>	<b>2019</b>	<b>2020</b>	<b>2021</b>
<b>50</b>	Wisconsin	51	35	66
<b>51</b>	Wyoming	1	0	3

	<b>Geographic Area</b>	<b>2019</b>
<b>55</b>	Wyoming	578759
<b>57</b>	Puerto Rico	3193694

	<b>Geographic Area</b>	<b>2020</b>	<b>2021</b>
<b>55</b>	Wyoming	577681	579636
<b>57</b>	Puerto Rico	3281590	3262711

All three dataframes have a column containing U.S. states, along with some other geographic areas. These columns are our **join keys**.

- Below, we use `df1.merge(right=df2, ...)` to carry out the merge ([documentation](#)).
- We could have alternatively used the function `pd.merge(left=df1, right=df2, ...)` ([documentation](#)).

```
# merge TB dataframe with two US census dataframes
tb_census_df = (
    tb_df
    .merge(right=census_2019_df,
          left_on="U.S. jurisdiction", right_on="Geographic Area")
    .merge(right=census_2020_2021_df,
          left_on="U.S. jurisdiction", right_on="Geographic Area")
)
tb_census_df.tail(2)
```

	<b>U.S. jurisdiction</b>	<b>2019_x</b>	<b>2020_x</b>	<b>2021_x</b>	<b>Geographic Area_x</b>	<b>2019_y</b>	<b>Geographic Area_y</b>	<b>2020_y</b>	<b>2021_y</b>
<b>49</b>	Wisconsin	51	35	66	Wisconsin	5822434	Wisconsin	5897375	5881608
<b>50</b>	Wyoming	1	0	3	Wyoming	578759	Wyoming	577681	579636

To see what's going on with the duplicate columns, and the `_x` and `_y`, let's do the just the first merge:

```
tb_df.merge(right=census_2019_df,
            left_on="U.S. jurisdiction",
            right_on="Geographic Area").head()
```

<b>U.S. jurisdiction</b>	<b>2019_x</b>	<b>2020</b>	<b>2021</b>	<b>Geographic Area</b>	<b>2019_y</b>
0 Alabama	87	72	92	Alabama	4903185
1 Alaska	58	58	58	Alaska	731545
2 Arizona	183	136	129	Arizona	7278717
3 Arkansas	64	59	69	Arkansas	3017804
4 California	2111	1706	1750	California	39512223

Notice that the columns containing the **join keys** have all been retained, and all contain the same values.

- Furthermore, notice that the duplicated columns are appended with `_x` and `_y` to keep the column names unique.
- In the TB case count data, column `2019` represents the number of TB cases in 2019, but in the Census data, column `2019` represents the U.S. population.

We can use the `suffixes` argument to modify the `_x` and `_y` defaults to our liking ([documentation](#)).

```
# Specify the suffixes to use for duplicated column names
tb_df.merge(right=census_2019_df,
            left_on="U.S. jurisdiction",
            right_on="Geographic Area",
            suffixes('_cases', '_population')).head()
```

<b>U.S. jurisdiction</b>	<b>2019_cases</b>	<b>2020</b>	<b>2021</b>	<b>Geographic Area</b>	<b>2019_population</b>
0 Alabama	87	72	92	Alabama	4903185
1 Alaska	58	58	58	Alaska	731545
2 Arizona	183	136	129	Arizona	7278717
3 Arkansas	64	59	69	Arkansas	3017804
4 California	2111	1706	1750	California	39512223

Notice the `_x` and `_y` have changed to `_cases` and `_population`, just like we specified.

Putting it all together, and dropping the duplicated `Geographic Area` columns:

```
# Redux: merge TB dataframe with two US census dataframes
tb_census_df = (
    tb_df

    .merge(right=census_2019_df,
           left_on="U.S. jurisdiction", right_on="Geographic Area",
           suffixes('_cases', '_population'))
    .drop(columns="Geographic Area")

    .merge(right=census_2020_2021_df,
           left_on="U.S. jurisdiction", right_on="Geographic Area",
```

```

        suffixes=('_cases', '_population'))
        .drop(columns="Geographic Area")

)
tb_census_df.tail(2)

```

	U.S. jurisdiction	2019_cases	2020_cases	2021_cases	2019_population	2020_population	2021_population
49	Wisconsin	51	35	66	5822434	5897375	5881608
50	Wyoming	1	0	3	578759	577681	579636

## 5.4.5 Reproducing Data: Compute Incidence

Let's see if we can reproduce the original CDC numbers from our augmented dataset of TB case counts and state populations.

- Recall that the nationwide TB incidence was **2.7 in 2019, 2.2 in 2020, and 2.4 in 2021**.
- Along the way, we'll also compute state-level incidence.

From the [CDC report](#): TB incidence is computed as “Cases per 100,000 persons using mid-year population estimates from the U.S. Census Bureau.”

Let's start with a simpler question: What is the per person incidence?

- In other words, what is the probability that a randomly selected person in the population had TB within a given year?

$$\text{TB incidence per person} = \frac{\# \text{ TB cases in population}}{\text{Total population size}}$$

Let's calculate per person incidence for 2019:

```

# Calculate per person incidence for 2019
tb_census_df["per person incidence 2019"] =
    tb_census_df["2019_cases"]/tb_census_df["2019_population"]
)
tb_census_df

```

	U.S. jurisdiction	2019_cases	2020_cases	2021_cases	2019_population	2020_population	2021_population	2019	per person incidence
0	Alabama	87	72	92	4903185	5033094	5049196	0.00	
1	Alaska	58	58	58	731545	733017	734420	0.00	
2	Arizona	183	136	129	7278717	7187135	7274078	0.00	
3	Arkansas	64	59	69	3017804	3014546	3026870	0.00	
4	California	2111	1706	1750	39512223	39521958	39142565	0.00	

U.S.								per person incidence
	jurisdiction	2019_cases	2020_cases	2021_cases	2019_population	2020_population	2021_population	2019
...	...	...	...	...	...	...	...	...
46	Virginia	191	169	161	8535519	8637615	8658910	0.00
47	Washington	221	163	199	7614893	7727209	7743760	0.00
48	West Virginia	9	13	7	1792147	1791646	1785618	0.00
49	Wisconsin	51	35	66	5822434	5897375	5881608	0.00
50	Wyoming	1	0	3	578759	577681	579636	0.00

51 rows × 8 columns

TB is really rare in the United States, so per person TB incidence is really low, as expected.

- But, if we were to consider 100,000 people, the probability of seeing a TB case is higher.
- In fact, it would be 100,000 times higher!

$$\text{TB incidence per 100,000} = 100,000 * \text{TB incidence per person}$$

```
# To help read bigger numbers in Python, you can use _ to separate thousands,
# akin to using commas. 100_000 is the same as writing 100000, but more readable.
tb_census_df["per 100k incidence 2019"] = (
    100_000 * tb_census_df["per person incidence 2019"])
)
tb_census_df
```

U.S.								per person incidence	per inc
	jurisdiction	2019_cases	2020_cases	2021_cases	2019_population	2020_population	2021_population	2019	2019
0	Alabama	87	72	92	4903185	5033094	5049196	0.00	1.7
1	Alaska	58	58	58	731545	733017	734420	0.00	7.9
2	Arizona	183	136	129	7278717	7187135	7274078	0.00	2.5
3	Arkansas	64	59	69	3017804	3014546	3026870	0.00	2.1
4	California	2111	1706	1750	39512223	39521958	39142565	0.00	5.3
...	...	...	...	...	...	...	...	...	...
46	Virginia	191	169	161	8535519	8637615	8658910	0.00	2.2
47	Washington	221	163	199	7614893	7727209	7743760	0.00	2.9
48	West Virginia	9	13	7	1792147	1791646	1785618	0.00	0.5
49	Wisconsin	51	35	66	5822434	5897375	5881608	0.00	0.8
50	Wyoming	1	0	3	578759	577681	579636	0.00	0.1

51 rows × 9 columns

Now we're seeing more human-readable values.

- For example, there 5.3 tuberculosis cases for every 100,000 California residents in 2019.

To wrap up this exercise, let's calculate the nationwide incidence of TB in 2019.

```
# Recall that the CDC reported an incidence of 2.7 per 100,000 in 2019.
tot_tb_cases_50_states = tb_census_df["2019_cases"].sum()
tot_pop_50_states = tb_census_df["2019_population"].sum()
tb_per_100k_50_states = 100_000 * tot_tb_cases_50_states / tot_pop_50_states
tb_per_100k_50_states
```

2.7114346007625656

We can use a `for` loop to compute the incidence for 2019, 2020, and 2021.

- You'll notice that we get the same numbers reported by the CDC!

```
# f strings (f"...") are a handy way to pass in variables to strings.
for year in [2019, 2020, 2021]:
    tot_tb_cases_50_states = tb_census_df[f"{year}_cases"].sum()
    tot_pop_50_states = tb_census_df[f"{year}_population"].sum()
    tb_per_100k_50_states = 100_000 * tot_tb_cases_50_states / tot_pop_50_states
    print(tb_per_100k_50_states)
```

2.7114346007625656

2.163293721906285

2.366758711298075

## 5.5 Summary

We went over a lot of content this lecture; let's summarize the most important points:

### 5.5.1 Dealing with Missing Values

There are a few options we can take to deal with missing data:

- Drop missing records
- Keep `Nan` missing values
- Impute using an interpolated column

### 5.5.2 EDA and Data Wrangling

There are several ways to approach EDA and Data Wrangling:

- Examine the **data and metadata**: what is the date, size, organization, and structure of the data?
- Examine each **field/attribute/dimension** individually.

- Examine pairs of related dimensions (e.g. breaking down grades by major).
- Along the way, we can:
  - **Visualize** or summarize the data.
  - **Validate assumptions** about data and its collection process. Pay particular attention to when the data was collected.
  - Identify and **address anomalies**.
  - Apply data transformations and corrections (we'll cover this in the upcoming lecture).
  - **Record everything you do!** Developing in Jupyter Notebook promotes *reproducibility* of your own work!

## 5.6 [BONUS] EDA Demo 2: Mauna Loa CO<sub>2</sub> Data – A Lesson in Data Faithfulness

We no longer cover the following demo in lecture, but we provide the following section in the course notes for interested students.

[Mauna Loa Observatory](#) has been monitoring CO<sub>2</sub> concentrations since 1958.

```
co2_file = "data/co2_mm_mlo.txt"
```

Let's do some **EDA!!**

### 5.6.1 Reading this file into Pandas ?

Let's instead check out this **.txt** file. Some questions to keep in mind: Do we trust this file extension? What structure is it?

Lines 71-78 (inclusive) are shown below:

line number	file contents						
71	#	decimal	average	interpolated	trend	#days	
72	#	date			(season corr)		
73	1958	3	1958.208	315.71	315.71	314.62	-1
74	1958	4	1958.292	317.45	317.45	315.29	-1
75	1958	5	1958.375	317.50	317.50	314.71	-1
76	1958	6	1958.458	-99.99	317.10	314.85	-1
77	1958	7	1958.542	315.86	315.86	314.98	-1
78	1958	8	1958.625	314.93	314.93	315.94	-1

Notice how:

- The values are separated by white space, possibly tabs.
- The data line up down the rows. For example, the month appears in 7th to 8th position of each line.
- The 71st and 72nd lines in the file contain column headings split over two lines.

We can use `read_csv` to read the data into a `pandas DataFrame`, and we provide several arguments to specify that the separators are white space, there is no header (**we will set our own column names**), and to skip the first

72 rows of the file.

```
co2 = pd.read_csv(
    co2_file, header = None, skiprows = 72,
    sep = r'\s+'      #delimiter for continuous whitespace (stay tuned for regex next le
)
co2.head()
```

0	1	2	3	4	5	6
0	1958	3	1958.21	315.71	315.71	314.62
1	1958	4	1958.29	317.45	317.45	315.29
2	1958	5	1958.38	317.50	317.50	314.71
3	1958	6	1958.46	-99.99	317.10	314.85
4	1958	7	1958.54	315.86	315.86	314.98

Congratulations! You've wrangled the data!

...But our columns aren't named. **We need to do more EDA.**

## 5.6.2 Exploring Variable Feature Types

The NOAA [webpage](#) might have some useful tidbits (in this case it doesn't).

Using this information, we'll rerun `pd.read_csv`, but this time with some **custom column names**.

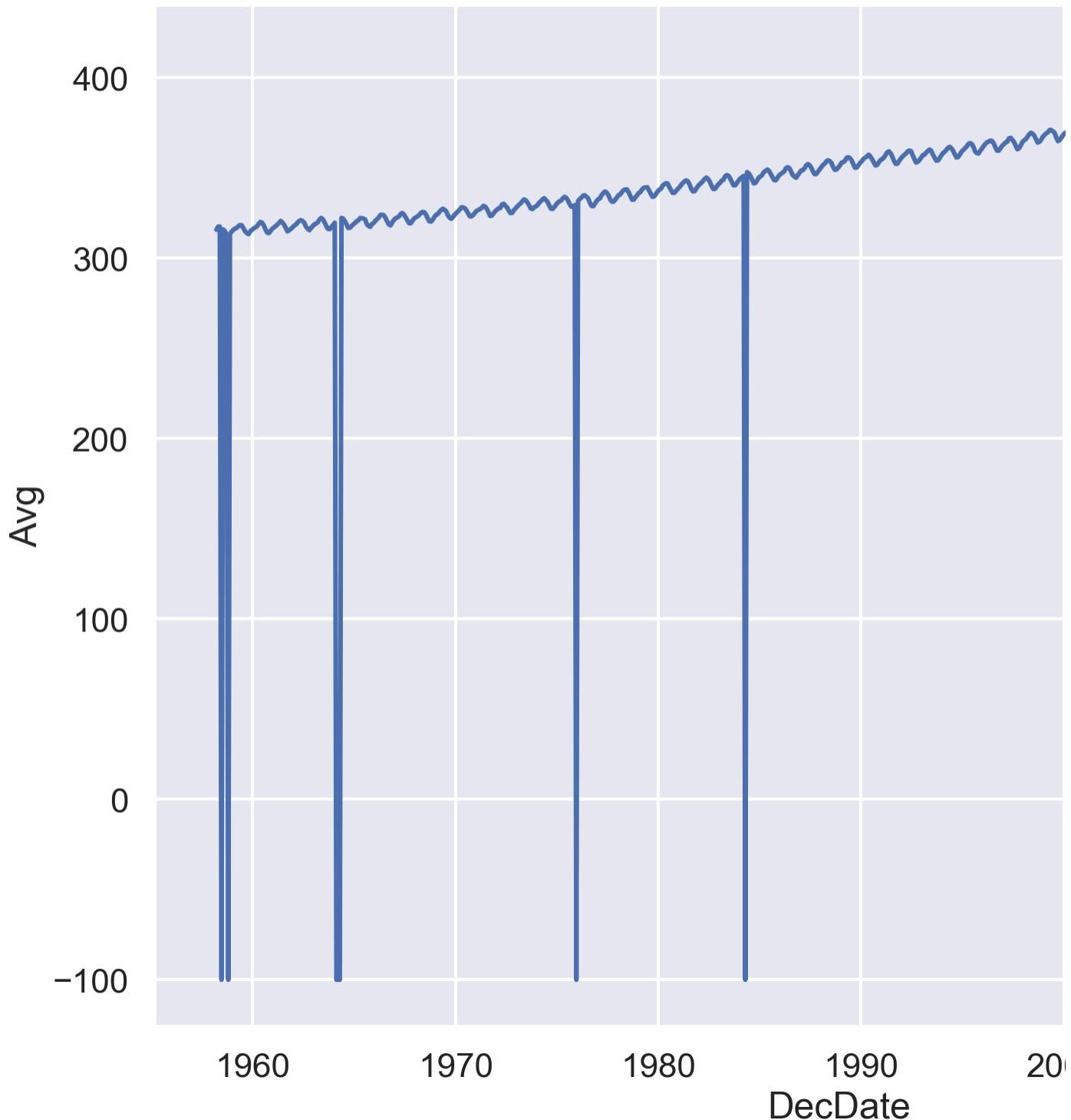
```
co2 = pd.read_csv(
    co2_file, header = None, skiprows = 72,
    sep = r'\s+', #regex for continuous whitespace (next lecture)
    names = ['Yr', 'Mo', 'DecDate', 'Avg', 'Int', 'Trend', 'Days']
)
co2.head()
```

	Yr	Mo	DecDate	Avg	Int	Trend	Days
0	1958	3	1958.21	315.71	315.71	314.62	-1
1	1958	4	1958.29	317.45	317.45	315.29	-1
2	1958	5	1958.38	317.50	317.50	314.71	-1
3	1958	6	1958.46	-99.99	317.10	314.85	-1
4	1958	7	1958.54	315.86	315.86	314.98	-1

## 5.6.3 Visualizing CO<sub>2</sub>

Scientific studies tend to have very clean data, right...? Let's jump right in and make a time series plot of CO<sub>2</sub> monthly averages.

► Code



The code above uses the `seaborn` plotting library (abbreviated `sns`). We will cover this in the Visualization lecture, but now you don't need to worry about how it works!

Yikes! Plotting the data uncovered a problem. The sharp vertical lines suggest that we have some **missing values**. What happened here?

```
co2.head()
```

	<b>Yr</b>	<b>Mo</b>	<b>DecDate</b>	<b>Avg</b>	<b>Int</b>	<b>Trend</b>	<b>Days</b>
<b>0</b>	1958	3	1958.21	315.71	315.71	314.62	-1
<b>1</b>	1958	4	1958.29	317.45	317.45	315.29	-1
<b>2</b>	1958	5	1958.38	317.50	317.50	314.71	-1
<b>3</b>	1958	6	1958.46	-99.99	317.10	314.85	-1
<b>4</b>	1958	7	1958.54	315.86	315.86	314.98	-1

```
co2.tail()
```

	<b>Yr</b>	<b>Mo</b>	<b>DecDate</b>	<b>Avg</b>	<b>Int</b>	<b>Trend</b>	<b>Days</b>
<b>733</b>	2019	4	2019.29	413.32	413.32	410.49	26
<b>734</b>	2019	5	2019.38	414.66	414.66	411.20	28
<b>735</b>	2019	6	2019.46	413.92	413.92	411.58	27
<b>736</b>	2019	7	2019.54	411.77	411.77	411.43	23
<b>737</b>	2019	8	2019.62	409.95	409.95	411.84	29

Some data have unusual values like -1 and -99.99.

Let's check the description at the top of the file again.

- -1 signifies a missing value for the number of days **Days** the equipment was in operation that month.
- -99.99 denotes a missing monthly average **Avg**

How can we fix this? First, let's explore other aspects of our data. Understanding our data will help us decide what to do with the missing values.

## 5.6.4 Sanity Checks: Reasoning about the data

First, we consider the shape of the data. How many rows should we have?

- If chronological order, we should have one record per month.
- Data from March 1958 to August 2019.
- We should have \$ 12 (2019-1957) - 2 - 4 = 738 \$ records.

```
co2.shape
```

(738, 7)

Nice!! The number of rows (i.e. records) match our expectations.

Let's now check the quality of each feature.

## 5.6.5 Understanding Missing Value 1: Days

**Days** is a time field, so let's analyze other time fields to see if there is an explanation for missing values of days of operation.

Let's start with **months**, **Mo**.

Are we missing any records? The number of months should have 62 or 61 instances (March 1957-August 2019).

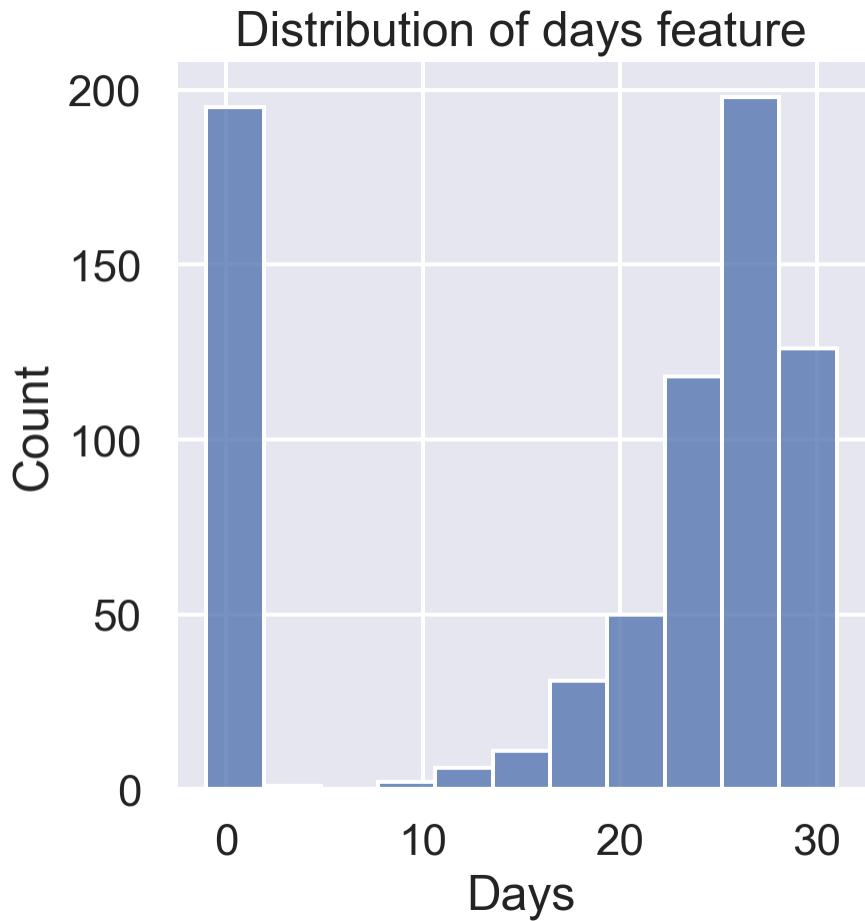
```
co2["Mo"].value_counts().sort_index()
```

```
Mo
1    61
2    61
3    62
4    62
5    62
6    62
7    62
8    62
9    61
10   61
11   61
12   61
Name: count, dtype: int64
```

As expected Jan, Feb, Sep, Oct, Nov, and Dec have 61 occurrences and the rest 62.

Next let's explore **days** **Days** itself, which is the number of days that the measurement equipment worked.

► Code



In terms of data quality, a handful of months have averages based on measurements taken on fewer than half the days. In addition, there are nearly 200 missing values—**that's about 27% of the data!**

Finally, let's check the last time feature, **year Yr**.

Let's check to see if there is any connection between missing-ness and the year of the recording.

► **Code**



#### Observations:

- All of the missing data are in the early years of operation.
- It appears there may have been problems with equipment in the mid to late 80s.

#### Potential Next Steps:

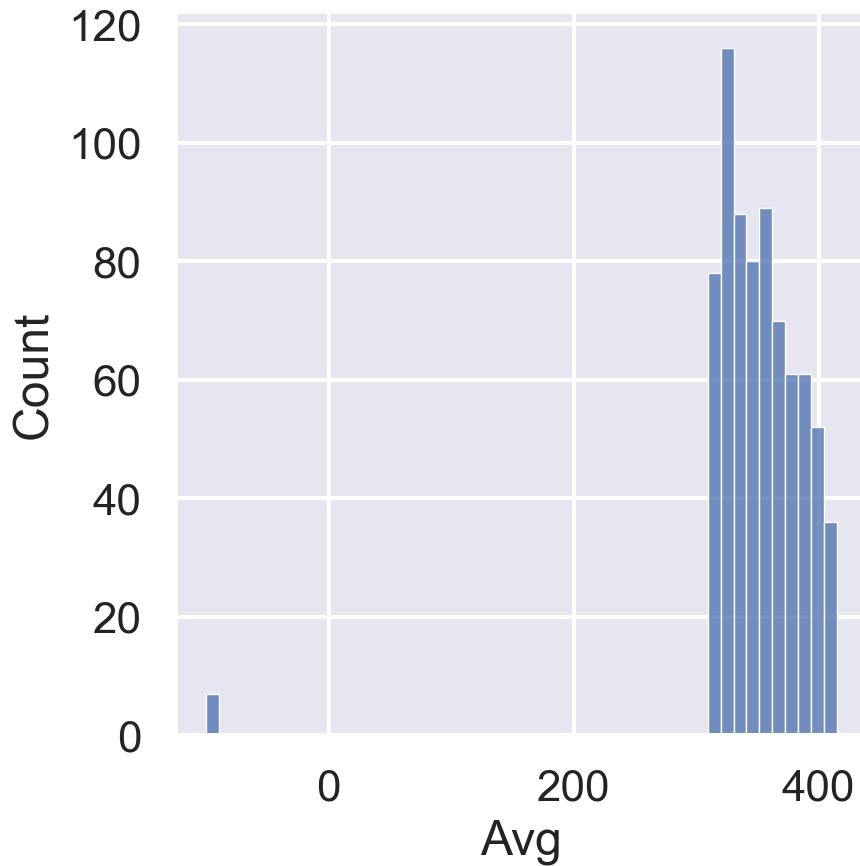
- Confirm these explanations through documentation about the historical readings.

- Maybe drop the earliest recordings? However, we would want to delay such action until after we have examined the time trends and assess whether there are any potential problems.

## 5.6.6 Understanding Missing Value 2: Avg

Next, let's return to the -99.99 values in `Avg` to analyze the overall quality of the CO<sub>2</sub> measurements. We'll plot a histogram of the average CO<sub>2</sub> measurements

► Code



The non-missing values are in the 300-400 range (a regular range of CO<sub>2</sub> levels).

We also see that there are only a few missing `Avg` values (**<1% of values**). Let's examine all of them:

```
co2[co2["Avg"] < 0]
```

	Yr	Mo	DecDate	Avg	Int	Trend	Days
3	1958	6	1958.46	-99.99	317.10	314.85	-1
7	1958	10	1958.79	-99.99	312.66	315.61	-1
71	1964	2	1964.12	-99.99	320.07	319.61	-1
72	1964	3	1964.21	-99.99	320.73	319.55	-1

<b>Yr</b>	<b>Mo</b>	<b>DecDate</b>	<b>Avg</b>	<b>Int</b>	<b>Trend</b>	<b>Days</b>	
<b>73</b>	1964	4	1964.29	-99.99	321.77	319.48	-1
<b>213</b>	1975	12	1975.96	-99.99	330.59	331.60	0
<b>313</b>	1984	4	1984.29	-99.99	346.84	344.27	2

There doesn't seem to be a pattern to these values, other than that most records also were missing **Days** data.

### 5.6.7 Drop, NaN , or Impute Missing Avg Data?

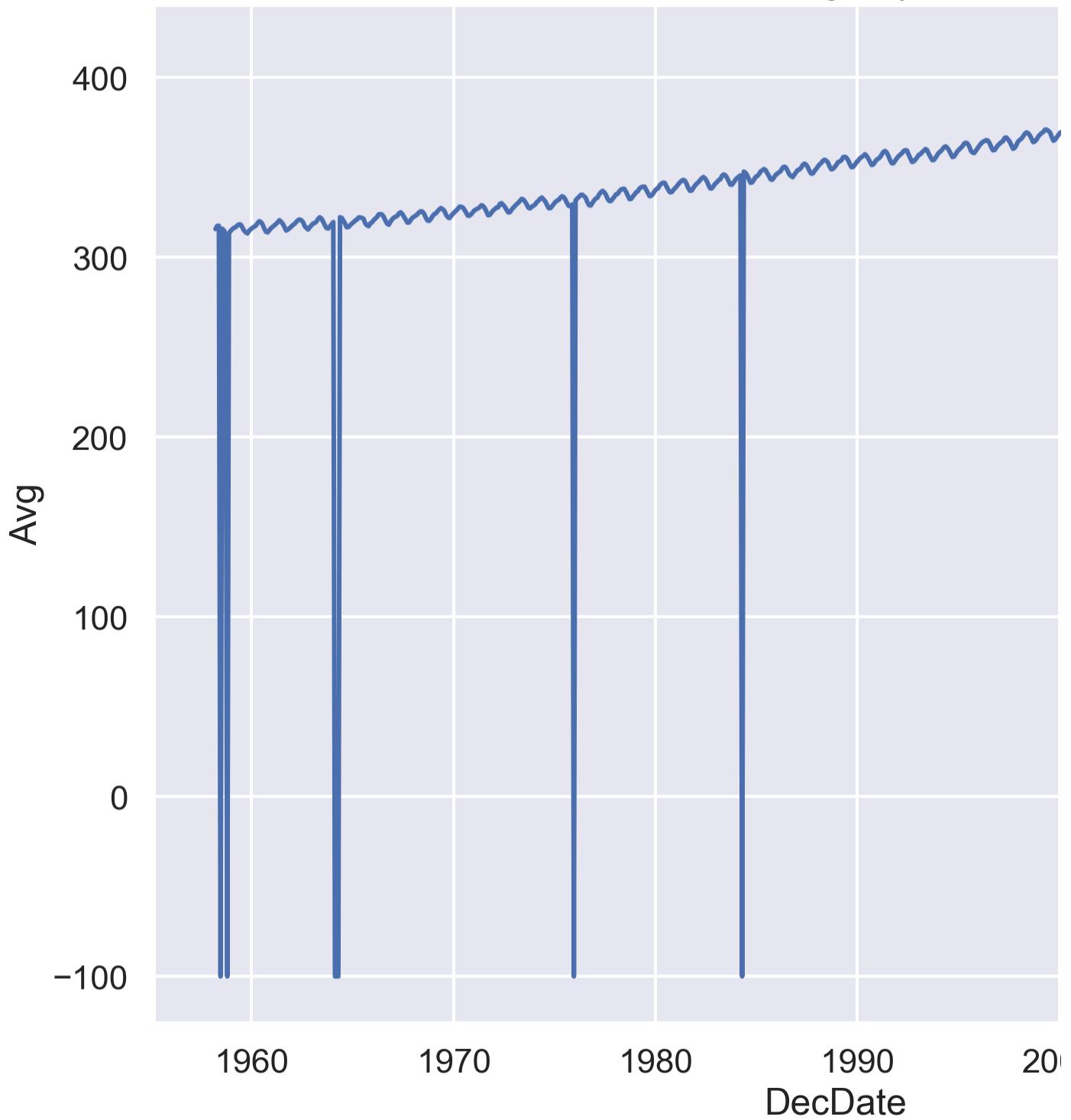
How should we address the invalid **Avg** data?

1. Drop records
2. Set to NaN
3. Impute using some strategy

Remember we want to fix the following plot:

► Code

## CO2 Average By Month



Since we are plotting `Avg` vs `DecDate`, we should just focus on dealing with missing values for `Avg`.

Let's consider a few options: 1. Drop those records 2. Replace -99.99 with NaN 3. Substitute it with a likely value for the average CO<sub>2</sub>?

What do you think are the pros and cons of each possible action?

Let's examine each of these three options.

```
# 1. Drop missing values
co2_drop = co2[co2['Avg'] > 0]
co2_drop.head()
```

	<b>Yr</b>	<b>Mo</b>	<b>DecDate</b>	<b>Avg</b>	<b>Int</b>	<b>Trend</b>	<b>Days</b>
<b>0</b>	1958	3	1958.21	315.71	315.71	314.62	-1
<b>1</b>	1958	4	1958.29	317.45	317.45	315.29	-1
<b>2</b>	1958	5	1958.38	317.50	317.50	314.71	-1
<b>4</b>	1958	7	1958.54	315.86	315.86	314.98	-1
<b>5</b>	1958	8	1958.62	314.93	314.93	315.94	-1

```
# 2. Replace NaN with -99.99
co2_NA = co2.replace(-99.99, np.nan)
co2_NA.head()
```

	<b>Yr</b>	<b>Mo</b>	<b>DecDate</b>	<b>Avg</b>	<b>Int</b>	<b>Trend</b>	<b>Days</b>
<b>0</b>	1958	3	1958.21	315.71	315.71	314.62	-1
<b>1</b>	1958	4	1958.29	317.45	317.45	315.29	-1
<b>2</b>	1958	5	1958.38	317.50	317.50	314.71	-1
<b>3</b>	1958	6	1958.46	NaN	317.10	314.85	-1
<b>4</b>	1958	7	1958.54	315.86	315.86	314.98	-1

We'll also use a third version of the data.

First, we note that the dataset already comes with a **substitute value** for the -99.99.

From the file description:

The **interpolated** column includes average values from the preceding column (**average**) and **interpolated values** where data are missing. Interpolated values are computed in two steps...

The **Int** feature has values that exactly match those in **Avg**, except when **Avg** is -99.99, and then a **reasonable** estimate is used instead.

So, the third version of our data will use the **Int** feature instead of **Avg**.

```
# 3. Use interpolated column which estimates missing Avg values
co2_impute = co2.copy()
co2_impute['Avg'] = co2['Int']
co2_impute.head()
```

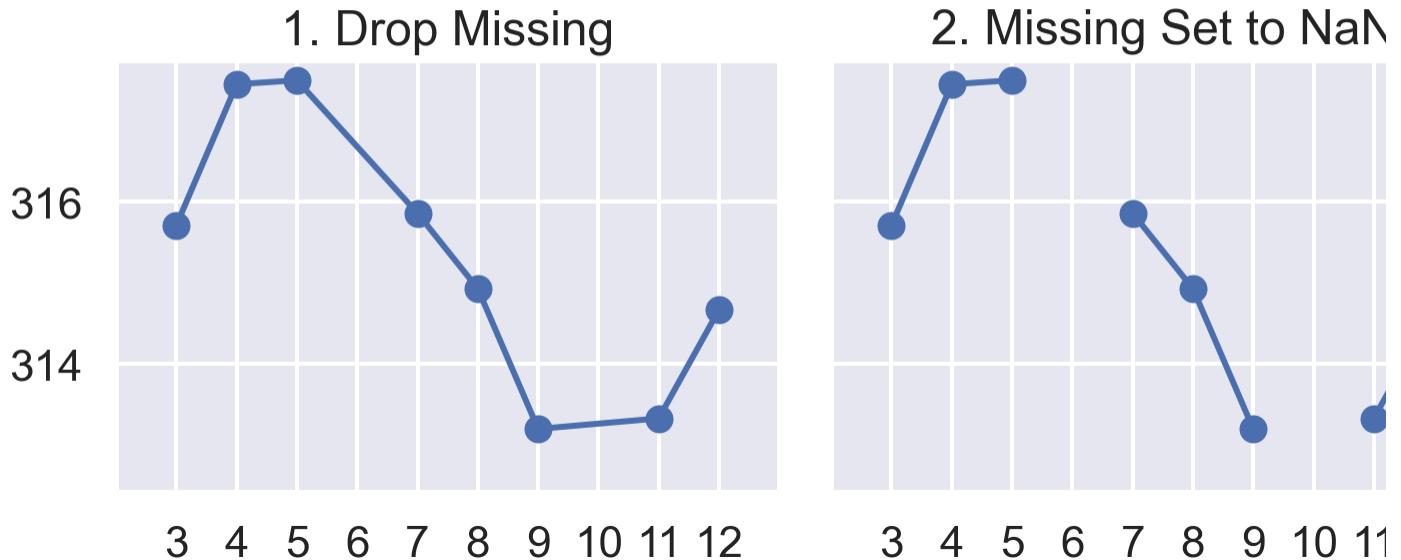
<b>Yr</b>	<b>Mo</b>	<b>DecDate</b>	<b>Avg</b>	<b>Int</b>	<b>Trend</b>	<b>Days</b>
<b>0</b>	1958	3	1958.21	315.71	315.71	-1
<b>1</b>	1958	4	1958.29	317.45	317.45	-1
<b>2</b>	1958	5	1958.38	317.50	317.50	-1
<b>3</b>	1958	6	1958.46	317.10	317.10	-1
<b>4</b>	1958	7	1958.54	315.86	315.86	-1

What's a **reasonable** estimate?

To answer this question, let's zoom in on a short time period, say the measurements in 1958 (where we know we have two missing values).

► Code

## Monthly Averages for 1958



In the big picture since there are only 7 **Avg** values missing (<1% of 738 months), any of these approaches would work.

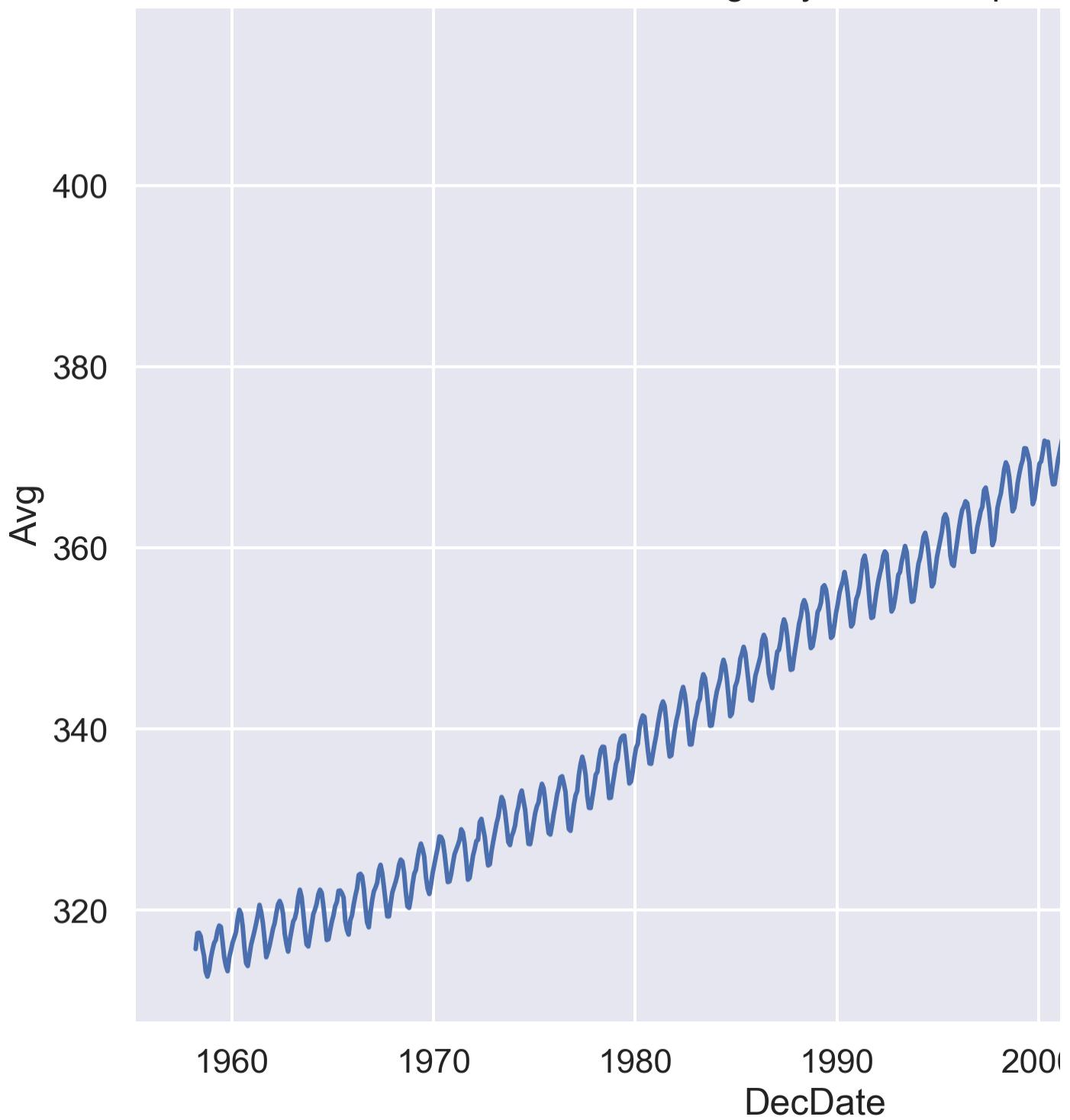
However there is some appeal to **option C, Imputing:**

- Shows seasonal trends for CO<sub>2</sub>
- We are plotting all months in our data as a line plot

Let's replot our original figure with option 3:

► Code

## CO2 Average By Month, Impute



Looks pretty close to what we see on the NOAA [website!](#)

### 5.6.8 Presenting the Data: A Discussion on Data Granularity

From the description:

- Monthly measurements are averages of average day measurements.
- The NOAA GML website has datasets for daily/hourly measurements too.

The data you present depends on your research question.

### How do CO<sub>2</sub> levels vary by season?

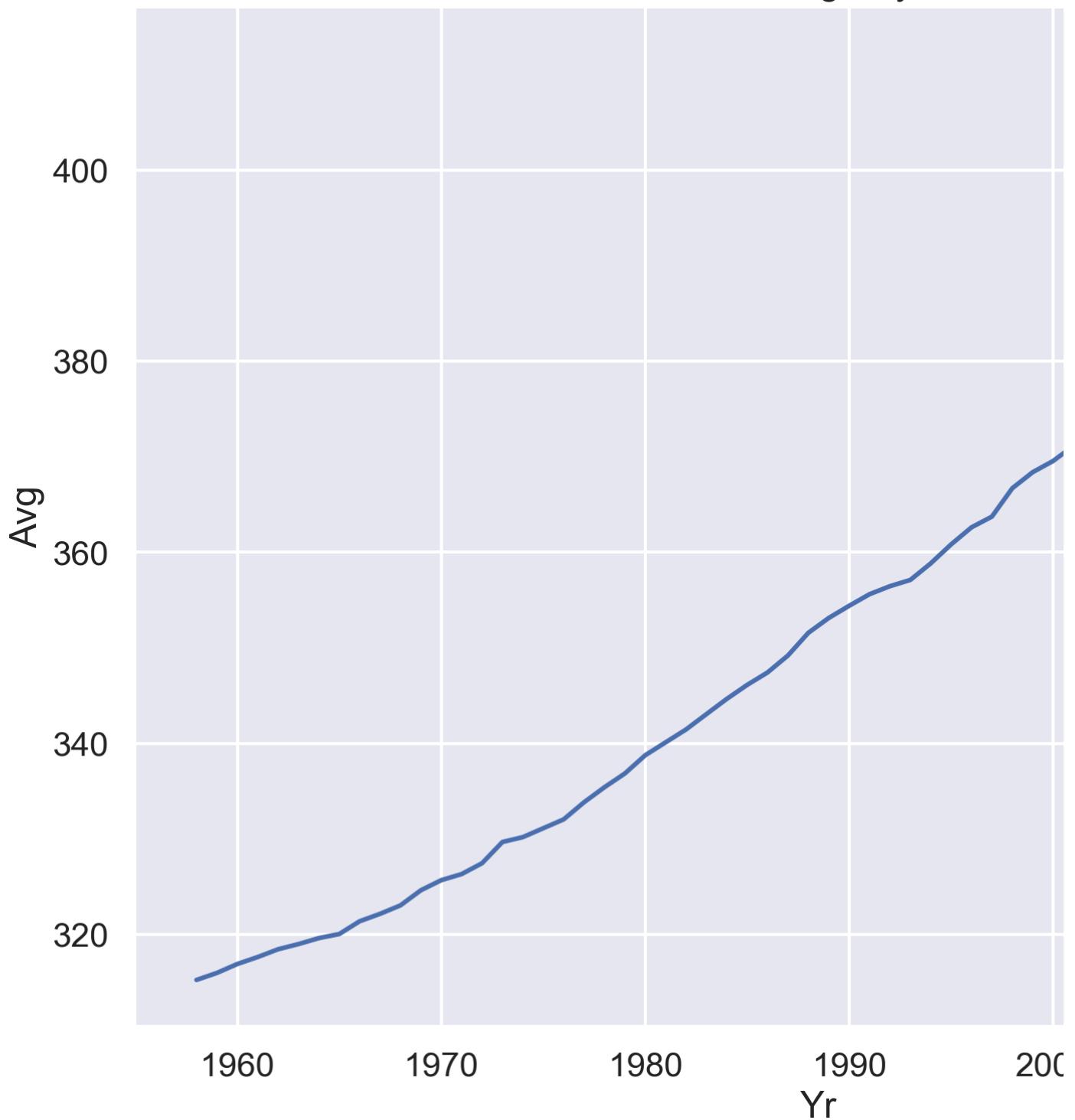
- You might want to keep average monthly data.

### Are CO<sub>2</sub> levels rising over the past 50+ years, consistent with global warming predictions?

- You might be happier with a **coarser granularity** of average year data!

► Code

## CO2 Average By Year



Indeed, we see a rise by nearly 100 ppm of CO<sub>2</sub> since Mauna Loa began recording in 1958.