

6 - Pandas-IngestData

March 22, 2023

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1 Ingest, Inspect & Clean

```
[3]: import numpy as np
import pandas as pd
from numpy.random import randn
```

1.1 Ingest data

https://pandas.pydata.org/docs/user_guide/io.html

NOTE - diabetes1 - 19 columns w/ header - diabetes2 - 19 columns no header

```
[4]: # https://raw.githubusercontent.com/jimcody2014/python-data/main/diabetes1.csv
# https://raw.githubusercontent.com/jimcody2014/python-data/main/diabetes2.csv
# https://raw.githubusercontent.com/jimcody2014/python-data/main/
↳diabetes_inspect.csv
```

1.1.1 Reading csv files

```
[5]: diabetes1 = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/
↳python-data/main/diabetes1.csv')
```

```
[6]: diabetes1.head()
```

```
[6]:
```

	encounter_id	patient_nbr	race	gender	age	weight	\
0	2278392	8222157	Caucasian	Female	[0-10)	?	
1	149190	55629189	Caucasian	Female	[10-20)	?	
2	64410	86047875	AfricanAmerican	Female	[20-30)	?	
3	500364	82442376	Caucasian	Male	[30-40)	?	
4	16680	42519267	Caucasian	Male	[40-50)	?	

	admission_type_id	admission_source_id	time_in_hospital	payer_code	\
0	6	1	1	?	
1	1	7	3	?	
2	1	7	2	?	
3	1	7	2	?	
4	1	7	1	?	

	num_lab_procedures	num_procedures	num_medications	diag_1	A1Cresult	\
0	41	0	1	250.83	None	
1	59	0	18	276	None	
2	11	5	13	648	None	
3	44	1	16	8	None	
4	51	0	8	197	None	

	metformin	miglitol	insulin	readmitted	Date
0	No	No	No	NO	10/20/2021
1	No	No	Up	>30	09/25/2021
2	No	No	No	NO	08/29/2021
3	No	No	Up	NO	10/20/2021
4	No	No	Steady	NO	09/25/2021

1.1.2 read_csv options

```
[7]: df = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/python-data/
↳main/diabetes2.csv')
```

```
[8]: df.head()
```

```
[8]:      2278392   8222157      Caucasian  Female  [0-10)  ?  6  1  1.1  ?  1  41  \
0    149190   55629189      Caucasian  Female  [10-20)  ?  1  7    3  ?  59
1     64410   86047875  AfricanAmerican  Female  [20-30)  ?  1  7    2  ?  11
2    500364   82442376      Caucasian    Male  [30-40)  ?  1  7    2  ?  44
3     16680   42519267      Caucasian    Male  [40-50)  ?  1  7    1  ?  51
4     35754   82637451      Caucasian    Male  [50-60)  ?  2  2    3  ?  31

      0  1.2  250.83  None  No  No.1   No.2   NO  10/20/21
0  0  18    276  None  No  No      Up  >30  9/25/21
1  5  13    648  None  No  No      No  NO   8/29/21
2  1  16     8  None  No  No      Up  NO  10/20/21
3  0   8    197  None  No  No  Steady  NO  9/25/21
4  6  16    414  None  No  No  Steady  >30  8/29/21
```

```
[9]: df = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/python-data/
↳main/diabetes2.csv',
                      header = None)
df.head()
```

```
[9]:      0      1      2      3      4  5  6  7  8  9  10  \
0  2278392   8222157      Caucasian  Female  [0-10)  ?  6  1  1  ?  41
1   149190   55629189      Caucasian  Female  [10-20)  ?  1  7  3  ?  59
2    64410   86047875  AfricanAmerican  Female  [20-30)  ?  1  7  2  ?  11
3   500364   82442376      Caucasian    Male  [30-40)  ?  1  7  2  ?  44
4    16680   42519267      Caucasian    Male  [40-50)  ?  1  7  1  ?  51

      11  12      13      14  15  16      17  18      19
0  0  1  250.83  None  No  No      No  NO  10/20/21
1  0  18    276  None  No  No      Up  >30  9/25/21
2  5  13    648  None  No  No      No  NO  8/29/21
3  1  16     8  None  No  No      Up  NO  10/20/21
4  0   8    197  None  No  No  Steady  NO  9/25/21
```

```
[10]: df = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/python-data/
↳main/diabetes2.csv',
                      header = None,
                      names = ('EncounterId','b','c','d','e','f','g','h','i','j','k',
                              'l','m','n','o','p','q','r','s','EDT'))
df.head()
```

```
[10]:
```

	EncounterId	b	c	d	e	f	g	h	i	j	k	\
0	2278392	8222157	Caucasian	Female	[0-10)	?	6	1	1	?	41	
1	149190	55629189	Caucasian	Female	[10-20)	?	1	7	3	?	59	
2	64410	86047875	AfricanAmerican	Female	[20-30)	?	1	7	2	?	11	
3	500364	82442376	Caucasian	Male	[30-40)	?	1	7	2	?	44	
4	16680	42519267	Caucasian	Male	[40-50)	?	1	7	1	?	51	

	l	m	n	o	p	q	r	s	EDT
0	0	1	250.83	None	No	No	No	NO	10/20/21
1	0	18	276	None	No	No	Up	>30	9/25/21
2	5	13	648	None	No	No	No	NO	8/29/21
3	1	16	8	None	No	No	Up	NO	10/20/21
4	0	8	197	None	No	No	Steady	NO	9/25/21

```
[11]: # Override existing column names
```

```
diabetes1 = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/
python-data/main/diabetes1.csv',
                        header = 0,
                        names = [
                            ('EncounterId', 'b', 'c', 'd', 'e', 'f', 'g', 'h', 'i', 'j', 'k',
                             'l', 'm', 'n', 'o', 'p', 'q', 'r', 's', 'EDT'))
diabetes1.head()
```

```
[11]:
```

	EncounterId	b	c	d	e	f	g	h	i	j	k	\
0	2278392	8222157	Caucasian	Female	[0-10)	?	6	1	1	?	41	
1	149190	55629189	Caucasian	Female	[10-20)	?	1	7	3	?	59	
2	64410	86047875	AfricanAmerican	Female	[20-30)	?	1	7	2	?	11	
3	500364	82442376	Caucasian	Male	[30-40)	?	1	7	2	?	44	
4	16680	42519267	Caucasian	Male	[40-50)	?	1	7	1	?	51	

	l	m	n	o	p	q	r	s	EDT
0	0	1	250.83	None	No	No	No	NO	10/20/2021
1	0	18	276	None	No	No	Up	>30	09/25/2021
2	5	13	648	None	No	No	No	NO	08/29/2021
3	1	16	8	None	No	No	Up	NO	10/20/2021
4	0	8	197	None	No	No	Steady	NO	09/25/2021

```
[12]: # Use a column as the index
```

```
df = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/python-data/
main/diabetes2.csv',
                 header = None,
                 names = ('EncounterId', 'b', 'c', 'd', 'e', 'f', 'g', 'h', 'i', 'j', 'k',
                          'l', 'm', 'n', 'o', 'p', 'q', 'r', 's', 'EDT'),
                 index_col = 'EncounterId')
df.head()
```

```
[12]:
```

	b	c	d	e	f	g	h	i	j	k	l	\
EncounterId												
2278392	8222157	Caucasian	Female	[0-10)	?	6	1	1	?	41	0	
149190	55629189	Caucasian	Female	[10-20)	?	1	7	3	?	59	0	
64410	86047875	AfricanAmerican	Female	[20-30)	?	1	7	2	?	11	5	
500364	82442376	Caucasian	Male	[30-40)	?	1	7	2	?	44	1	
16680	42519267	Caucasian	Male	[40-50)	?	1	7	1	?	51	0	

	m	n	o	p	q	r	s	EDT
EncounterId								
2278392	1	250.83	None	No	No	No	NO	10/20/21
149190	18	276	None	No	No	Up	>30	9/25/21
64410	13	648	None	No	No	No	NO	8/29/21
500364	16	8	None	No	No	Up	NO	10/20/21
16680	8	197	None	No	No	Steady	NO	9/25/21

```
[13]: df.loc[64410]
```

```
[13]: b      86047875
      c  AfricanAmerican
      d      Female
      e    [20-30)
      f      ?
      g      1
      h      7
      i      2
      j      ?
      k     11
      l      5
      m     13
      n     648
      o     None
      p      No
      q      No
      r      No
      s      NO
      EDT    8/29/21
      Name: 64410, dtype: object
```

```
[14]: # Only use a subset of the columns

df = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/python-data/
↳main/diabetes2.csv',
                 header = None,
                 names = ('EncounterId','b','c','d','e','EDT'),
                 index_col = 'EncounterId',
                 usecols = [0,1,2,3,7,19])
```

```
#df.head()
```

```
[15]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 101768 entries, 2278392 to 443797076
Data columns (total 5 columns):
#   Column  Non-Null Count  Dtype
---  -
0    b      101768 non-null  int64
1    c      101768 non-null  object
2    d      101768 non-null  object
3    e      101768 non-null  int64
4   EDT    101768 non-null  object
dtypes: int64(2), object(3)
memory usage: 4.7+ MB
```

```
[16]: # Change data types as the data is read in
df = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/python-data/
↳main/diabetes2.csv',
                  header = None,
                  names = ('EncounterId','b','c','d','e','EDT'),
                  index_col = 'EncounterId',
                  usecols = [0,1,2,3,7,19],
                  dtype = {'e':object }
                  )
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 101768 entries, 2278392 to 443797076
Data columns (total 5 columns):
#   Column  Non-Null Count  Dtype
---  -
0    b      101768 non-null  int64
1    c      101768 non-null  object
2    d      101768 non-null  object
3    e      101768 non-null  object
4   EDT    101768 non-null  object
dtypes: int64(1), object(4)
memory usage: 4.7+ MB
```

```
[17]: # Change data types as the data is read in
df = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/python-data/
↳main/diabetes2.csv',
                  header = None,
                  names = ('EncounterId','b','c','d','e','EDT'),
                  index_col = 'EncounterId',
```

```

        usecols = [0,1,2,3,7,19],
        dtype = object)
df.info()

```

```

<class 'pandas.core.frame.DataFrame'>
Int64Index: 101768 entries, 2278392 to 443797076
Data columns (total 5 columns):
#   Column  Non-Null Count  Dtype
---  -
0    b      101768 non-null  object
1    c      101768 non-null  object
2    d      101768 non-null  object
3    e      101768 non-null  object
4    EDT     101768 non-null  object
dtypes: object(5)
memory usage: 4.7+ MB

```

1.1.3 Ingesting Dates

```

[18]: # Change data types as the data is read in
df = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/python-data/
↳main/diabetes2.csv',
                header = None,
                names = ('EncounterId','b','c','d','e','EDT'),
                index_col = 'EncounterId',
                usecols = [0,1,2,3,7,19],
                dtype = {'e':object, },
                parse_dates=['EDT']
                )
df.info()

```

```

<class 'pandas.core.frame.DataFrame'>
Int64Index: 101768 entries, 2278392 to 443797076
Data columns (total 5 columns):
#   Column  Non-Null Count  Dtype
---  -
0    b      101768 non-null  int64
1    c      101768 non-null  object
2    d      101768 non-null  object
3    e      101768 non-null  object
4    EDT     101768 non-null  datetime64[ns]
dtypes: datetime64[ns](1), int64(1), object(3)
memory usage: 4.7+ MB

```

```

[19]: df

```

```
[19]:
```

	b	c	d	e	EDT
EncounterId					
2278392	8222157	Caucasian	Female	1	2021-10-20
149190	55629189	Caucasian	Female	7	2021-09-25
64410	86047875	AfricanAmerican	Female	7	2021-08-29
500364	82442376	Caucasian	Male	7	2021-10-20
16680	42519267	Caucasian	Male	7	2021-09-25
...
443854148	41088789	Caucasian	Male	7	2021-10-20
443857166	31693671	Caucasian	Female	7	2021-09-25
443867222	175429310	Caucasian	Male	7	2021-08-29
62256	49726791	AfricanAmerican	Female	2	2021-10-20
443797076	183766055	Caucasian	Male	1	2021-09-25

[101768 rows x 5 columns]

```
[20]: df = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/python-data/
↳main/diabetes2.csv',
                        header = None,
                        names = ('EncounterId', 'b', 'c', 'd', 'e', 'EDT'),
                        index_col = 'EncounterId',
                        usecols = [0,1,2,3,7,19],
                        dtype = {'e':object, },
                        parse_dates=['EDT'],
                        dayfirst=True                                # Why isn't this working?
                        )
df
```

```
[20]:
```

	b	c	d	e	EDT
EncounterId					
2278392	8222157	Caucasian	Female	1	2021-10-20
149190	55629189	Caucasian	Female	7	2021-09-25
64410	86047875	AfricanAmerican	Female	7	2021-08-29
500364	82442376	Caucasian	Male	7	2021-10-20
16680	42519267	Caucasian	Male	7	2021-09-25
...
443854148	41088789	Caucasian	Male	7	2021-10-20
443857166	31693671	Caucasian	Female	7	2021-09-25
443867222	175429310	Caucasian	Male	7	2021-08-29
62256	49726791	AfricanAmerican	Female	2	2021-10-20
443797076	183766055	Caucasian	Male	1	2021-09-25

[101768 rows x 5 columns]

```
[21]: df.shape
```

```
[21]: (101768, 5)
```



```
[22]: # Only bring in X number of rows
df = pd.read_csv('https://raw.githubusercontent.com/jimcody2014/python-data/
↳main/diabetes2.csv',
                header = None,
                names = ('EncounterId','b','c','d','e'),
                index_col = 'EncounterId',
                usecols = [0,1,2,3,7],
                nrows = 1000)

df.shape
```

[22]: (1000, 4)

1.1.4 Reading other file types

- excel
- json
- APIs
- database

```
[35]: # Read in an excel spreadsheet
# I had to install openpyxl in my anaconda environment for this to work.

imm = pd.read_excel('/Users/jamescody/Documents/python-data/immunotherapy.xlsx')
imm.head()
```

```
[35]:
```

	sex	age	Time	Number_of_Warts	Type	Area	induration_diameter	\
0	1	22	2.25	14	3	51		50
1	1	15	3.00		3	900		70
2	1	16	10.50		1	100		25
3	1	27	4.50		3	80		30
4	1	20	8.00		1	45		8

	Result_of_Treatment
0	1
1	1
2	1
3	1
4	1

```
[ ]: # Saving a dataframe to json format
imm.to_json('')
```

```
[34]: # Read in a json file

imm2 = pd.read_json('/Users/jamescody/Documents/python-data/immjson.json')
imm2.head()
```

```
[34]: sex age Time Number_of_Warts Type Area induration_diameter \
0 1 22 2.25 14 3 51 50
1 1 15 3.00 2 3 900 70
2 1 16 10.50 2 1 100 25
3 1 27 4.50 9 3 80 30
4 1 20 8.00 6 1 45 8

Result_of_Treatment
0 1
1 1
2 1
3 1
4 1
```

1.1.5 Using an API as input - - - This will not run on Kaggle

```
[27]: import requests
```

```
[28]: # Send the request and receive a response
# Get the content of the response (there are other parts of the response (e.g.,
↳header))
# Display the content in json format

response = requests.get("https://data.cdc.gov/resource/saz5-9hgg.json")
jsonhold = response.json()
jsonhold
```

```
[28]: [{'jurisdiction': 'Connecticut',
'week_of_allocations': '2021-06-21T00:00:00.000',
'_1st_dose_allocations': '54360',
'_2nd_dose_allocations': '54360'},
{'jurisdiction': 'Maine',
'week_of_allocations': '2021-06-21T00:00:00.000',
'_1st_dose_allocations': '21420',
'_2nd_dose_allocations': '21420'},
{'jurisdiction': 'Massachusetts',
'week_of_allocations': '2021-06-21T00:00:00.000',
'_1st_dose_allocations': '104580',
'_2nd_dose_allocations': '104580'},
{'jurisdiction': 'New Hampshire',
'week_of_allocations': '2021-06-21T00:00:00.000',
'_1st_dose_allocations': '21420',
'_2nd_dose_allocations': '21420'},
{'jurisdiction': 'Rhode Island',
'week_of_allocations': '2021-06-21T00:00:00.000',
'_1st_dose_allocations': '17280',
'_2nd_dose_allocations': '17280'},
```

```

{'jurisdiction': 'Vermont',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '11070',
 '_2nd_dose_allocations': '11070'},
{'jurisdiction': 'New Jersey',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '132210',
 '_2nd_dose_allocations': '132210'},
{'jurisdiction': 'New York',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '167670',
 '_2nd_dose_allocations': '167670'},
{'jurisdiction': 'New York City',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '128070',
 '_2nd_dose_allocations': '128070'},
{'jurisdiction': 'Puerto Rico',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '52290',
 '_2nd_dose_allocations': '52290'},
{'jurisdiction': 'U.S. Virgin Islands',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '3240',
 '_2nd_dose_allocations': '3240'},
{'jurisdiction': 'Delaware',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '15210',
 '_2nd_dose_allocations': '15210'},
{'jurisdiction': 'District of Columbia',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '11520',
 '_2nd_dose_allocations': '11520'},
{'jurisdiction': 'Maryland',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '89370',
 '_2nd_dose_allocations': '89370'},
{'jurisdiction': 'Pennsylvania',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '170460',
 '_2nd_dose_allocations': '170460'},
{'jurisdiction': 'Philadelphia',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '24660',
 '_2nd_dose_allocations': '24660'},
{'jurisdiction': 'Virginia',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '126000',

```

```

    '_2nd_dose_allocations': '126000'},
{'jurisdiction': 'West Virginia',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '28800',
 '_2nd_dose_allocations': '28800'},
{'jurisdiction': 'Alabama',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '72090',
 '_2nd_dose_allocations': '72090'},
{'jurisdiction': 'Florida',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '314640',
 '_2nd_dose_allocations': '314640'},
{'jurisdiction': 'Georgia',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '149490',
 '_2nd_dose_allocations': '149490'},
{'jurisdiction': 'Kentucky',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '65880',
 '_2nd_dose_allocations': '65880'},
{'jurisdiction': 'Mississippi',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '44460',
 '_2nd_dose_allocations': '44460'},
{'jurisdiction': 'North Carolina',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '149940',
 '_2nd_dose_allocations': '149940'},
{'jurisdiction': 'South Carolina',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '74160',
 '_2nd_dose_allocations': '74160'},
{'jurisdiction': 'Tennessee',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '98820',
 '_2nd_dose_allocations': '98820'},
{'jurisdiction': 'Chicago',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '40770',
 '_2nd_dose_allocations': '40770'},
{'jurisdiction': 'Illinois',
 'week_of_allocations': '2021-06-21T00:00:00.000',
 '_1st_dose_allocations': '149490',
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  '_1st_dose_allocations': '0',
  '_2nd_dose_allocations': '0'},
{'jurisdiction': 'Guam',
  'week_of_allocations': '2021-03-08T00:00:00.000',
  '_1st_dose_allocations': '0',
  '_2nd_dose_allocations': '0'},
{'jurisdiction': 'Marshall Islands',
  'week_of_allocations': '2021-03-08T00:00:00.000',
  '_1st_dose_allocations': '0',
  '_2nd_dose_allocations': '0'}]
```

```
[29]: # Put the content into a DataFrame
      # Display the DataFrame
```

```
vaccines = pd.DataFrame(jsonhold)
vaccines
```

```
[29]:
```

	jurisdiction	week_of_allocations	_1st_dose_allocations	\
0	Connecticut	2021-06-21T00:00:00.000		54360
1	Maine	2021-06-21T00:00:00.000		21420
2	Massachusetts	2021-06-21T00:00:00.000		104580
3	New Hampshire	2021-06-21T00:00:00.000		21420
4	Rhode Island	2021-06-21T00:00:00.000		17280
..	
995	Hawaii	2021-03-08T00:00:00.000		18720

996	Nevada	2021-03-08T00:00:00.000	36270
997	American Samoa	2021-03-08T00:00:00.000	0
998	Guam	2021-03-08T00:00:00.000	0
999	Marshall Islands	2021-03-08T00:00:00.000	0

	_2nd_dose_allocations
0	54360
1	21420
2	104580
3	21420
4	17280
..	...
995	18720
996	36270
997	0
998	0
999	0

[1000 rows x 4 columns]

```
[ ]: # Same process
      # Just set the url as a variable
      # Combined a few other steps

      #response = requests.get(url)
      #vermont = pd.DataFrame(response.json())
      #vermont
```

1.1.6 A local example running on my machine

```
[ ]: #import requests
      url = 'http://localhost:8080/rest/applicants'
      response = requests.get(url)
      json1 = response.json()
      json1
      # Notice that all the data is wrapped in 'content'
```

```
[ ]: # We need to get the results inside of 'content'.
      #df5 = pd.DataFrame(json1['content'])
      #df5
```

1.2 Exercise - 15 minutes

1.2.1 Part 1

- Bring the csv file diabetes_inspect into a DataFrame. Name the dataframe df.

- What is its shape?

1.2.2 Part 2

- Read the outbreaks2.csv file into a pdf. Name the df something you will remember.
- The csv file does not have a header row.
- This data is a listing of food born illness outbreaks.
- Do not bring in the 6th or 8th columns. The df should have 10 columns.
- The remaining columns contain the following data. You can decide how to name the columns
 - year
 - month
 - state
 - location
 - food
 - species
 - status
 - illnesses
 - hospitalizations
 - fatalities
- What is the shape of this DataFrame?

A	B	C	D	E	G	I	J	K	L
Year	Month	State	Location	Food	Species	Status	Illnesses	Hospitalizati	Fatalities
1998	January	California	Restaurant				20	0	0
1998	January	California		Custard			112	0	0
1998	January	California	Restaurant				35	0	0
1998	January	California	Restaurant	Fish, Ahi	Scombroid toxin	Confirmed	4	0	0
1998	January	California	Private Home/Residence	Lasagna, Unspecified; Eggs, Other	Salmonella enterica	Confirmed	26	3	0
1998	January	California	Restaurant		Shigella boydii	Confirmed	25	3	0
1998	January	California	Restaurant				8	0	0
1998	January	California	Restaurant	Stuffing, Unspecified; Sandwich, Turkey	Salmonella enterica	Confirmed	4	3	0

```
[ ]: # Part 1
```

2 Renaming Preview

2.1 Rename using names in read_csv

```
[36]: imm = pd.read_excel('/Users/jamescody/Documents/python-data/immunotherapy.
↳xlsx', sheet_name = 'Immuno1')
imm.head()
```

```
[36]:   sex  age  Time  Number_of_Warts  Type  Area  induration_diameter  \
0    1   22   2.25                14    3    51                    50
1    1   15   3.00                 2    3   900                    70
2    1   16  10.50                 2    1   100                    25
3    1   27   4.50                 9    3    80                    30
4    1   20   8.00                 6    1    45                     8
```

	Result_of_Treatment
0	1
1	1
2	1
3	1
4	1

```
[37]: imm = pd.read_excel('/Users/jamescody/Documents/python-data/immunotherapy.xlsx',
                        names = ('a','b','c','d','e','f','g','h'),
                        sheet_name = 'Immuno1')
imm.head()
```

```
[37]:
```

	a	b	c	d	e	f	g	h
0	1	22	2.25	14	3	51	50	1
1	1	15	3.00	2	3	900	70	1
2	1	16	10.50	2	1	100	25	1
3	1	27	4.50	9	3	80	30	1
4	1	20	8.00	6	1	45	8	1

2.2 Rename all columns using rename()

```
[38]: imm.rename(columns = {'a':'a1','b':'b1','c':'c1','d':'d1','e':'e1','f':'f1','g':
    ↳ 'g1','h':'h1'}, inplace = True)
imm.head()

# An alternative - using the axis=1
# imm.rename({'a':'a1','b':'b1','c':'c1','d':'d1','e':'e1','f':'f1','g':
    ↳ 'g1','h':'h1'}, axis =1, inplace = True)
```

```
[38]:
```

	a1	b1	c1	d1	e1	f1	g1	h1
0	1	22	2.25	14	3	51	50	1
1	1	15	3.00	2	3	900	70	1
2	1	16	10.50	2	1	100	25	1
3	1	27	4.50	9	3	80	30	1
4	1	20	8.00	6	1	45	8	1

2.3 Rename some columns using rename()

```
[39]: imm.rename(columns = {'b1':'BB','e1':'EE'}, inplace = True)
imm.head()
```

```
[39]:
```

	a1	BB	c1	d1	EE	f1	g1	h1
0	1	22	2.25	14	3	51	50	1
1	1	15	3.00	2	3	900	70	1
2	1	16	10.50	2	1	100	25	1
3	1	27	4.50	9	3	80	30	1

```
4    1    20    8.00    6    1    45    8    1
```

2.4 Rename using setaxis()

```
[40]: imm.set_axis(['AA','BB','CC','DD','EE','FF','GG','HH'], axis = 'columns',  
    ↪inplace = True)  
imm.head()
```

```
[40]:
```

	AA	BB	CC	DD	EE	FF	GG	HH
0	1	22	2.25	14	3	51	50	1
1	1	15	3.00	2	3	900	70	1
2	1	16	10.50	2	1	100	25	1
3	1	27	4.50	9	3	80	30	1
4	1	20	8.00	6	1	45	8	1

2.5 Rename using .columns

```
[41]: imm.columns = ['AaA','BbB','CcC','DdD','EeE','FfF','GgG','HhH']  
imm.head()
```

```
[41]:
```

	AaA	BbB	CcC	DdD	EeE	FfF	GgG	HhH
0	1	22	2.25	14	3	51	50	1
1	1	15	3.00	2	3	900	70	1
2	1	16	10.50	2	1	100	25	1
3	1	27	4.50	9	3	80	30	1
4	1	20	8.00	6	1	45	8	1

2.6 Rename using str.replace (a string function)

```
[42]: imm.columns = imm.columns.str.replace('CcC','CCC')  
imm.head()
```

```
[42]:
```

	AaA	BbB	CCC	DdD	EeE	FfF	GgG	HhH
0	1	22	2.25	14	3	51	50	1
1	1	15	3.00	2	3	900	70	1
2	1	16	10.50	2	1	100	25	1
3	1	27	4.50	9	3	80	30	1
4	1	20	8.00	6	1	45	8	1

```
[ ]:
```