Pandas

Today we are going to really start using pandas. Lets review the libraries or packages and what they have done so far.

- 1. We always start byt typing %matploltlib inline. This is a built in magic command that enables us to plot the data right into our ipython notebook https://ipython.org/ipython-doc/3/interactive/magics.html
- 2. we import matplotlib.pyplot as plt. This turns on all the graphing capabilities and then uses the shorthand plt. for when we call functions from matplotlib. This used to be called pylab but was updated to pyplot. Then we say fig,ax=plt.subplots() and all the plot functions go into fig and ax
- 3. we import numpy as np. This turns on math functions and we use the shorthand np.
- 4. from scipy we import stats. scipy gives us a lot of analysis functions and we use linear regression from stats.
- 5. Now we are also going to use pandas. Pandas is database management. It lets us take complicated datasets and anlyze them. You can think of it like a supercharged excel where you combine the organization of excel with the power of a programming language. It can do amazing things and I am still learning every day. So lets get started!
- 6. What is pandas? http://pandas.pydata.org/index.html and here is the documentation http://pandas.pydata.org/pandas-docs/stable/
- 7. import pandas as pd!!!!!!
- 8. On a final note you can see I made a numbered list in markdown. To do that you type a number a period and then two spaces.
- 9. Also in terms of line numnbers. I turn my line numbers on so it is easier to debug. Do this under view

```
In [2]: %matplotlib inline
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from scipy import stats
```

GDP_Lifespan_all_data.csv

Libby_Thesis_Data.csv

Importing files

We are going to start by finding our csv file and reading it in

I only want to list the csv files so I can see what I can read in. so I will do Is *.csv the star is a wildcard that means everything and then .csv is only ones that end in .csv. Today we are going to look at data from well water chemistry in Bangladesh. Specifically arsenic concentrations and if people drink the water. We will also look at the rest of the chemistry. We are looking at well water arsenic because drinking water with arsenic has negative long term health impacts. The US standard for arsenic is 10 ppb or 10 ug/L. The bangladesh standard is 50 ppb. Lets see what we can learn! We are going to try and learn about how many people drink water with 10 or 50 ppb arsenic. (to show the star I had to type *)

The data is on the edblogs siteh ttps://edblogs.columbia.edu/eescx3050-001-2015-3/category/classes/class-10-start-pandas/ or on the github site https://github.com/bmaillou/BigDataPython/blob/master/well_data.csv.

In [4]:	pwd	
Out[4]:	'/Users/bmaillou/Documents/w	ork-teaching/python/fall21/BigDataPython'
In [5]:	ls *.csv	
	Brian.csv CoreEM09GC01-extra-line.csv CoreEM09GC01.csv GDP-Lifespan - Copy.csv GDP-Lifespan.csv	gdp2 - Copy.csv gdp2.csv gdp2015.csv gdp_and_lifespan.csv gdp_only.csv

gdp_only_download.csv

mystery.csv

Well-As.csv	twoD1.csv
central_park.csv	weekly_mlo.csv
fldav_ljo.csv	well_data.csv
fldav_ljo_Yasna.csv	well_sites.csv
gdp.csv	

now we read in a well_data.csv. But I want to use pandas and not numpy.

But we are going to read in some data and try to analyze it. open the well_data.csv. It is for wells from Bangladesh. every well has an id#, a latitude and longitude, Depth, if people drink it and then some concentration data. lets use readcsv to get read in. In Pandas you are trying to get your data into a dataframe which is like an excel sheet. It will have column titles and an index for rows. It is all about the dataframes. When using pandas people name things 'df' a lot. That is shorthand for dataframe. I am not a good namer.

I am going to just name it df today.

In [3]: df=pd.read_csv('well_data.csv')

The data is now magically in the computers memory even if we can't see it we can access it!

This is important. Your output may not look like my output. It changes between computers depending on default settings when you installed. Don't worry. If you see data of descriptions you are fine.

just typing well_data will give us some descriptions of what we got! It used the first row for column names!

In [4]:	df												
Out[4]:		Well_ID	Lat	Lon	Depth	Drink	Si	Р	S	Ca	Fe	•••	1
	0	2	23.74	90.31	45	Y	NaN	NaN	NaN	NaN	NaN		Na
	1	14	23.62	90.60	60	Y	NaN	NaN	NaN	NaN	NaN		Na
	2	23	23.94	91.46	60	Y	NaN	NaN	NaN	NaN	NaN		N٤
	3	83	23.80	91.33	50	Y	48084.33842	0.936358	2085.570979	54666.48199	1.260031		76.20744
	4	84	23.98	90.81	150	Y	NaN	NaN	NaN	NaN	NaN		Na
	754	12516	24.71	90.41	160	Y	32379.64000	0.197380	3669.430000	39790.24000	0.341200		26.2992{
	755	12654	24.36	91.27	60	Y	25561.12000	0.090570	13771.370000	57630.63000	1.498350		19.72336
	756	72641	24.38	90.90	45	Ν	31319.48000	1.162550	38.300000	60905.16000	22.417560		14.1149
	757	76175	23.90	90.65	60	Ν	30605.53000	1.556120	4168.520000	66756.16000	12.793100		33.1803(
	758	141499	23.60	91.34	50	Ν	NaN	NaN	NaN	NaN	NaN		N٤

759 rows × 21 columns

In []:

Since we didn't set an index it just numbers each row and calls that the index. But that doesn't help us. I think we could set the well_id to the index. When you look at your data above. see how the numbers on the left have no title but are a little offset. That is the index. But what is an index. I am not sure. It is sort of like a master column that helps us organize the data. It will make more sense when we get to timeseries analysis. That is where pandas shines even more. But lets set and index and use well_id as that is the most important factor.

In [5]: df=df.set_index('Well_ID')

In [6]: df

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Pandas_Well_Data

24, 4:40 PM								Pandas_W	ell_Data				
Out[6]:		Lat	Lon	Depth	Drink		Si	Ρ	S	Ca	Fe		Ва
	Well_ID												
	2	23.74	90.31	45	Y		NaN	NaN	NaN	NaN	NaN	Ν	laN
	14	23.62	90.60	60	Y		NaN	NaN	NaN	NaN	NaN	Ν	laN
	23	23.94	91.46	60	Y		NaN	NaN	NaN	NaN	NaN	N	laN
	83	23.80	91.33	50	Y	48084.33	3842 0.93	6358 2	085.570979	54666.48199	1.260031	96.1595	587 76.20
	84	23.98	90.81	150	Y		NaN	NaN	NaN	NaN	NaN	N	laN
											· ···		
	12516	24.71	90.41	160	Y	32379.64	000 0.19	7380 36	69.430000	39790.24000	0.341200	77.7700	000 26.29
	12654	24.36	91.27	60	Y	25561.12	2000 0.09	0570 13	771.370000	57630.63000	1.498350	73.2300	00 19.72
	72641	24.38	90.90	45	Ν	31319.48	3000 1.16	2550	38.300000	60905.16000	22.417560	176.0200	000 14.1
	76175	23.90	90.65	60	Ν	30605.53	3000 1.55	6120 4 ⁻	168.520000	66756.16000	12.793100	178.8700	00 33.18
	141499	23.60	91.34	50	Ν		NaN	NaN	NaN	NaN	NaN	Ν	laN
In []: In [7]:	we can u df=df		the inde index										
In [8]:	df #s.	ince w	ve have	an in					on its ow				
Out[8]:		ell_ID	Lat		Depth		Si		P	S	Са	Fe	1
	0			90.31	45	Y	NaN	Na		NaN	NaN	NaN	Na
	1		23.62		60	Y	NaN	Na		NaN	NaN	NaN	Na
	2			91.46	60	Y	NaN	Na		NaN	NaN	NaN	Na
	3			91.33	50		084.33842	0.93635				30031	
	4			90.81	150	Y	NaN	Na		NaN	NaN	NaN	Na
		12516		90.41	160		379.64000	0.19738					26.29928
	755 ´	12654	24.36	91.27	60	Y 25	561.12000	0.09057	0 13771.37	0000 57630.	63000 1.49	8350	19.72336

758 141499 23.60 91.34

72641 24.38 90.90

76175 23.90 90.65

756

757

759 rows × 21 columns

Or we could just read in the data with the index set.

In [9]: df=pd.read_csv('well_data.csv',index_col='Well_ID')

If you don't know the column name you can use the column number!

45

60

50

Ν

Ν

Ν

31319.48000

30605.53000

NaN

1.162550

1.556120

NaN

38.300000

NaN

4168.520000

60905.16000 22.417560 ...

66756.16000 12.793100

NaN

In [10]: df=pd.read_csv('well_data.csv',index_col=0)

The first great trick of pandas!

The describe function. It gives you amazing summary statistics lickety-split!

14.1149

33.18036

Na

...

...

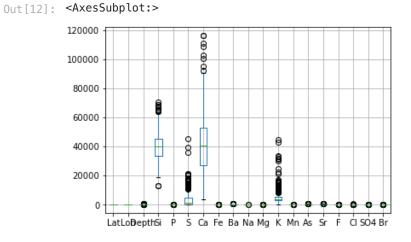
NaN

[11]:	df.de	escribe()								
[11]:		Lat	Lon	Depth	Si	Р	S	Ca	Fe	
	count 759.000000		759.000000	759.000000	407.000000	407.000000	407.000000	407.000000	407.000000	4
	mean	23.789249	90.641199	65.554677	40101.151444	0.809323	3407.292389	41129.291921	5.556200	
	std 0.578493		0.578800	42.186161	10117.680290	0.902860	5364.247733	20161.130827	5.153779	
	min	22.780000	89.610000	0.000000	12605.576700	0.008210	-41.390000	3577.160000	-0.003680	
	25%	23.285000	90.155000	45.000000	33200.310900	0.151957	149.635000	26996.273955	1.706806	
	50%	23.790000	90.650000	50.000000	40021.490000	0.507850	1220.877945	40166.830000	3.931310	
	75%	24.300000	91.130000	70.000000	45369.825000	1.189271	4341.695000	52976.458285	8.531585	
	max	24.770000	91.650000	523.000000	70304.057950	5.477616	45035.460000	116040.620000	30.192230	29

A hint of what is to come! But we just got all of our summary statistics.

In [12]:

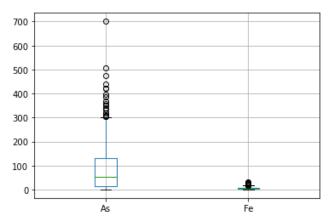
df.boxplot()



That boxplot was hard to see. What if we just look at As and Fe?

```
In [13]: fig,ax=plt.subplots()
    df.boxplot(column=['As', 'Fe'],ax=ax)
```

```
Out[13]: <AxesSubplot:>
```



this plotting is a little diffrent then how we have been plotting. Pandas has some built in plotting so you can make some really nice and quick plots. But these plots are a little harder to customize. So we will be doing both types of plotting depending on the goal. The goal could be a quick view versus a profesional looking plot.

We can also just get a list of our columns.

In [14]:	df.columns
Out[14]:	Index(['Lat', 'Lon', 'Depth', 'Drink', 'Si', 'P', 'S', 'Ca', 'Fe', 'Ba', 'Na', 'Mg', 'K', 'Mn', 'As', 'Sr', 'F', 'Cl', 'S04', 'Br'], dtype='object')
	Why did the columns not have parantheses? I am learning this. But each dataframe has attributes and methods.
	Methods uses paranthesis. Think of it as having to do something. An attribute just tells you about the dataframe and
	doesn't need parantheses. Methods can take extra arguments.
	Remember NaN is not a number. We are going to use this to our advantage!
	shape still gives us the shape. We can call it two different ways
In [15]:	df.shape
Out[15]:	(759, 20)
T. [10]	nn chono(df)
In [16]:	np.shape(df)
Out[16]:	(759, 20)

Stop and think for a second. What does this shape mean?

It means we are starting to analyze a lot of data. It is a dataset with 759 rows or wells and 20 columns or different parameters. This will already get hard to deal with in excel!

We have to slow down and learn some Pandas basics. this is a critical section. Take your time

Now how do we get at our data. How do we slice it. There are many ways. lets go through them all.

.ix

.loc

.iloc

[]

We are going to do a lot of practice and then I tried to make a cheat sheet/table. Take lots of notes.

[] works like normal except you can only use integers on rows and names on columns. you can't use integers on both rows and columns.

I am putting .head() on the print statements to save paper. You don't need them. It just shows the first 5 rows

[17]:	df[:].	<pre>df[:].head() #I am including head to shorten my printouts</pre>										
17]:		Lat	Lon	Depth	Drink	Si	Р	S	Ca	Fe	Ва	N
	Well_ID											
	2	23.74	90.31	45	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
	14	23.62	90.60	60	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
	23	23.94	91.46	60	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
	83	23.80	91.33	50	Y	48084.33842	0.936358	2085.570979	54666.48199	1.260031	96.159587	76.20744
	84	23.98	90.81	150	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na

In [18]: print(df['As'].head())

8/24, 4:40 PM	Pandas_well_Data
	Well_ID 2 NaN 14 NaN 23 NaN 83 78.97747 84 NaN Name: As, dtype: float64
In [19]:	<pre>print (df[:]['As'].head()) #This is the same as the one above showing the rows</pre>
	Well_ID 2 NaN 14 NaN 23 NaN 83 78.97747 84 NaN Name: As, dtype: float64
In [21]:	<pre>print (df[30:50]['As']) #This prints rows 30-50.</pre>
	#Don't get confused as well_ID is our index and the name of the row
	Well_ID 330 10.233204 333 NaN 342 NaN 356 NaN 374 18.365596 389 50.285003 397 115.834040 398 NaN 402 17.755544 403 81.859568 410 NaN 414 NaN 415 87.102492 417 NaN 418 386.827954 420 79.798479 421 142.409968 434 NaN 475 270.785974 478 56.883257 Name: As, dtype: float64
In [22]:	<pre>print (df[30:50:2]['As']) #we skipped by twos!</pre>
	Well_ID 330 10.233204 342 NaN 374 18.365596 397 115.834040 402 17.755544 410 NaN 415 87.102492 418 386.827954 421 142.409968 475 270.785974 Name: As, dtype: float64 But you can pass a list to the columns you want! SEE the double brackets??? It is a list in the brackets!
In [23]:	df[30:50:2][['As','Depth']]
Out[23]:	As Depth
	Well_ID 330 10.233204 45
	330 10.233204 45

Well_ID		
330	10.233204	45
342	NaN	30
374	18.365596	45
397	115.834040	45
402	17.755544	30

As Depth

Well_ID		
410	NaN	60
415	87.102492	60
418	386.827954	65
421	142.409968	150
475	270.785974	55

And the order doesn't matter. So somehow it is smart about rows and columns

Pandas_Well_Data

In I	15
	20

df[['As','Depth']][30:50:2]

As Depth

\cap		нI	Г	2	5	1	
U	u	L I	L	2	2	1	=

Well_ID		
330	10.233204	45
342	NaN	30
374	18.365596	45
397	115.834040	45
402	17.755544	30
410	NaN	60
415	87.102492	60
418	386.827954	65
421	142.409968	150
475	270.785974	55

In [26]: df[['Depth', 'As']][30:50:2]

	Depth	As
Well_ID		
330	45	10.233204
342	30	NaN
374	45	18.365596
397	45	115.834040
402	30	17.755544
410	60	NaN
415	60	87.102492
418	65	386.827954
421	150	142.409968
475	55	270.785974

In [156...

Out[26]:

What I am teaching you is easy and hard at the same time. Take your time. It is a lot. I am showing you how to get at data. I just showed you brackets and now I am going to show you .loc. Also remember I just add .head to shorten the printouts. you can remove it.

.loc only uses names of the index and the columns.

THIS IS DIFFERENT. It is saying if my index matches this name then print it. This is a little confusing our the wells have numbers for names

Sometimes I put print sometimes not. It doesn't always matter and sometimes one looks nicer than the other.

In [28]:	df.loc	:[:].he	ead()	#gives	s us a	ll rows with	n all ind	exes				
Out[28]:		Lat	Lon	Depth	Drink	Si	Р	S	Са	Fe	Ва	٨
	Well_ID											
	2	23.74	90.31	45	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
	14	23.62	90.60	60	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
	23	23.94	91.46	60	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
	83	23.80	91.33	50	Y	48084.33842	0.936358	2085.570979	54666.48199	1.260031	96.159587 7	6.20744
	84	23.98	90.81	150	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
In [29]:	<pre>[29]: df.loc[101:156] #gives us all rows with all indexes but the numbers have t</pre>				rs have to m	natch an ind	ex. The n	umbers had	to ma			
Out[29]:		Lat		Depth		Si	Р	S	Са	Fe	Ва	
Out[23]:	Well_ID	Lut	Lon	Deptii	DIIIK	01	·	0	<u> </u>	10	Bu	
	101	24.40	90.26	60	Y	34311.71514	0.117534	2618.717799	42646.99574	1.843156	58.666191	28.28
	107	24.02	89.67	45	Ν	NaN	NaN	NaN	NaN	NaN	NaN	
	110	23.39	91.35	45	Y	47417.95635	1.095644	113.180915	46848.09017	11.740445	131.582974	23.58
	112	24.61	91.18	60	Y	37289.99489	2.448648	13.335397	65129.07627	8.923465	134.435231	15.87
	116	22.96	89.77	60	Y	NaN	NaN	NaN	NaN	NaN	NaN	
	130	22.94	89.97	60	Ν	44023.88418	1.172086	1023.167741	80183.25742	6.349396	159.171636	32.82
	153	24.17	90.81	45	Y	40523.43773	0.091676	2848.048146	40703.88184	1.869486	69.657716	29.68
	156	22.84	91.56	60	Ν	48375.82211	0.979053	1420.255478	52694.25919	13.020352	129.253387	23.80
In [30]:	df.loc	[101]	# j	ust ca	ll one	index. The	is is wel	l 101				
Out[30]:	Lat Lon Depth Drink Si P S Ca Fe Ba Na Mg K Mn As Sr F Cl SO4 Br Name: 3	34: 0.1 26 1. 58 22 22 1. 28 12 0 38	24.4 90.26 60 Y 311.7 17534 18.72 42647 84316 .6662 8.281 .5784 NaN 19269 .0709 3.043 .1994 .1123 7.518 .0552	object								
In [32]:	df.loc			-	index	doesn't exis	st you ge	t an error				
In []:												
	But we can use column names											

In [33]:	df.loc	[:]['As'].h	ead()		
Out[33]:	2 14 23 83 7 84 Name: A	NaN NaN NaN 78.97747 NaN NaN		mas as well as multiple brackets	
In [34]:	df.loc	[:,'As'].he	ad ()		
Out[34]:	84 Name: A	NaN NaN NaN 78.97747 NaN NaN		ows if you just get one column then it turns from a dataframe to a series	
In [35]:		(type(df.lo (type(df.lo			
	<class< td=""><td>'pandas.co</td><td>re.seri</td><td>es.Series'> .es.Series'></td><td></td></class<>	'pandas.co	re.seri	es.Series'> .es.Series'>	
		•		netimes be important. A series is more like a set of values.	
In [36]:	print	(df.loc[330	:500:2]]['As'])	
	402 410 415 418 421 475 481 488 500 Name: A We can a	10.233204 NaN 18.365596 115.834040 17.755544 NaN 87.102492 386.827954 142.409968 270.785974 NaN NaN NaN NaN S, dtype:	of name	es	
In [37]:	df.loc	[330:500:2]		,'Depth']]	
Out[37]:	Well_ID	As	Depth		
	330	10.233204	45		
	342	NaN	30		
	374	18.365596	45		
	397	115.834040	45		
	402	17.755544	30		
	410	NaN	60		
	415	87.102492	60		
	440	200 027054	C F		

 $local host: 8888/nbconvert/html/Documents/work-teaching/python/spring\ 24/BigDataPython/Pandas_Well_Data.ipynb?download=falsered between the second second$

418 386.827954

421 142.409968

475 270.785974

65

150

55

	As	Depth
Well_ID		
481	NaN	60
488	NaN	45
500	NaN	60

iloc

iloc only uses integers. So now this is row numbers. NOT the index. look the Well_ID compared to the iloc numberss

In [38]:	df.ilc	df.iloc[:].head()										
Out[38]:		Lat	Lon	Depth	Drink	Si	Р	S	Ca	Fe	Ва	٨
	Well_ID											
	2	23.74	90.31	45	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
	14	23.62	90.60	60	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
	23	23.94	91.46	60	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
	83	23.80	91.33	50	Y	48084.33842	0.936358	2085.570979	54666.48199	1.260031	96.159587	76.20744
	84	23.98	90.81	150	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na

In [39]: df.iloc[101:110:2] #this is row numbers now. so the index is not matching.

Out[39]:		Lat	Lon	Depth	Drink	Si	Р	S	Ca	Fe	Ва	Na
	Well_ID											
	3058	23.92	90.47	60	Y	NaN	NaN	NaN	NaN	NaN	NaN	Nal
	3060	23.62	91.56	60	Y	37199.96208	0.949837	13.694978	51862.73844	10.061950	119.569875	11.641918
	3103	23.33	90.12	130	Y	NaN	NaN	NaN	NaN	NaN	NaN	Nal
	3112	24.43	91.05	45	Y	41513.82677	1.697027	14.933618	47308.53575	14.961449	133.853170	22.606914
	3179	23.18	90.78	50	Y	NaN	NaN	NaN	NaN	NaN	NaN	Nal

and column number. But we use a column seperator......

64

In [40]: df.iloc[101:110:2,5]

Out[40]: Well_ID	
3058 NaN	
3060 0.949837	
3103 NaN	
3112 1.697027	
3179 NaN	
Name: P, dtype: f	loat

In [41]: df.iloc[101:110:2,2:5]

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Pandas_Well_Data

		E 4			
	114	1 /1	1		-
J	ut	14	-	1	-

Well_ID			
3058	60	Y	NaN
3060	60	Y	37199.96208
3103	130	Y	NaN
3112	45	Υ	41513.82677
3179	50	Y	NaN

Depth Drink

Just to boggles your bind a little.....

Depth

In [42]: df.iloc[101:110:2,[2,5,8]] #I just had it show columns 2,5,8

Fe

Si

Out[42]:

Well_ID			
3058	60	NaN	NaN
3060	60	0.949837	10.061950
3103	130	NaN	NaN
3112	45	1.697027	14.961449
3179	50	NaN	NaN

Ρ

ix was phased out.

```
In [43]: print (df.ix[101:110:2,[2,5,8]])
```

```
AttributeError Traceback (most recent call last)
<ipython-input-43-6c367ed50da5> in <module>
----> 1 print (df.ix[101:110:2,[2,5,8]])
~/anaconda3/lib/python3.8/site-packages/pandas/core/generic.py in __getattr__(self, name)
5137 if self._info_axis._can_hold_identifiers_and_holds_name(name):
5138 return self[name]
-> 5139 return object.__getattribute__(self, name)
5140
5141 def __setattr__(self, name: str, value) -> None:
AttributeError: 'DataFrame' object has no attribute 'ix'
```

In []:

Dot notation

I am not sure if that is the official name but here is how it works

What I didn't show you is a dot notation.

df.As.head() In [44]: Well_ID Out[44]: 2 NaN 14 NaN 23 NaN 83 78.97747 84 NaN Name: As, dtype: float64 Dot notation only works if you name your columns well. No minus signs or spaces.

In [45]: df.As[20:30]

Out[45]:	Well_	ID		
	233		NaN	
	237		NaN	
	275		NaN	
	279		NaN	
	280	5	.364619	
	283		NaN	
	287		NaN	
	290		NaN	
	292	53	.097829	
	295		NaN	
	Name:	As,	dtype:	float64

All examples in one place so maybe we can make sense of them?

Name	Description
[]	
well_data[:]	all data
well_data[:]['As']	all arsenic data
well_data.loc[:,'As']	basically the same as above
well_data['As']	all arsenic data.
well_data[1:10]['As']	arsenic data from rows 1-10 excluding 10
well_data[1:10:2]['As']	same but skipping by two
well_data[1:10:2][['As','Depth']]	for As and depth. note the double brackets.
well_data[['As','Depth']][1:10:2]	order doesn't matter
well_data[['Depth','As']][1:10:2]	order doesn't matter.
you can't use column numbers	

Description	loc
gives us all rows with all columns	well_data.loc[:]
needs to be an index Gives us by index number not row number.	well_data.loc[101:156]
and we can skip	well_data.loc[101:156:2]
and we can do column names	well_data.loc[101:156:2]['As']
	well_data.loc[:]['As']
is the same as above. I have bugs where one works but other doesn't	well_data.loc[:,'As']
we can do multiple columns	well_data.loc[101:156:2][['As','Depth']]

iloc	
well_data.iloc[:]	gives it all.
well_data.iloc[101:110:2]	does row numbers.
well_data.iloc[101:110:2,5]	row number by column number
well_data.iloc[101:110:2,2:5]	mulitple row multiple number
well_data.iloc[101:110:2,[2,5,8]]	select columns

ix phased out

Dot notation.	This can be very nice.
well_data.As	gives all arsenic data
well_data.As[1:5]	gives rows 1-5

You can use boolean choices to get the data you want. For example I gave the description if people drink or don't drink from their well. Lets count that.

value_counts is a great first function. It just counts for you. Simple but very helpful.

I am going to do the same thing many different ways! value_counts is a function that counts each

In [46]:	df['Drink'].value_counts()
Out[46]:	Y 614 N 144 Name: Drink, dtype: int64 is the same as
In [48]:	df.Drink.value_counts()
Out[48]:	Y 614 N 144 Name: Drink, dtype: int64 Is the same as (I am trying to teach you pandas)
In [49]:	<pre>df.iloc[:,3].value_counts()</pre>
Out[49]:	Y 614 N 144 Name: Drink, dtype: int64 Is the same as (I am trying to teach you pandas)
In [50]:	<pre>df.loc[:,'Drink'].value_counts()</pre>
Out[50]:	Y 614 N 144 Name: Drink, dtype: int64 Now you should be able to access your data. I always forget the semantics. Look online or back at your cheat sheets. That is why I made the cheat sheet above.
	Now we can sub-select data very easily.
	We can return a boolean based on results.
In [51]:	df['Drink']=='Y'
Out[51]:	Well_ID 2 True 14 True 23 True 83 True 84 True

True

12516	Ti	rue			
12654	Ti	rue			
72641	Fa	lse			
76175	Fa	lse			
141499	9 Fa	lse			
Name:	Drink,	Length:	759 ,	dtype:	bool
Also do	it with t	he dot not	ation		

In [53]: df.Drink=='Y'

Out[53]:	Well_ID 2 14 23 83 84	True True True True True
		True True False False False ink, Length: 759, dtype: bool only want data from wells people drink from? we can ask for that. Remember I just added the .head() to
In [59]:	df[df.D	rink=='Y'].head()

Out[59]:

	Lat	Lon	Depth	Drink	Si	Р	S	Ca	Fe	Ва	٨
Well_ID											
2	23.74	90.31	45	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
14	23.62	90.60	60	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
23	23.94	91.46	60	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na
83	23.80	91.33	50	Y	48084.33842	0.936358	2085.570979	54666.48199	1.260031	96.159587	76.20744
84	23.98	90.81	150	Y	NaN	NaN	NaN	NaN	NaN	NaN	Na

What if we only wanted arsenic concentrations where people drink the water?

This is weird again.

You are saying only give me As.

```
In [60]: df[df.Drink=='Y']['As'].head()
```

Out[60]: Well_ID

2 NaN 14 NaN 23 NaN 83 78.97747 84 NaN Name: As, dtype: float64

In the crazy world of pandas where you put the Arsenic doesn't matter.

```
In [62]: df['As'][df.Drink=='Y'].head()
```

Out[62]: Well_ID

2 NaN 14 NaN 23 NaN 83 78.97747 84 NaN Name: As, dtype: float64

Say you wanted to do an intervention. You would want to go to the houses with the highest arsenic first. So we could ask what are the well id's for people who drink water and their arsenic is greater than 250 ppb. This would be poeple

with high exposure! We would need to use an and statement.

In pandas you do this two ways.

- np.logical_and().
- Else you can use the & but YOU NEED PARANTHESE. REMEMBER THIS!!!!! It will come back and help you.

We would try to convince these households to switch. The drinking water standard is 10 ppb. This is really crazy high exposure.

In [64]: df['As'][(df.Drink=='Y') & (df.As>250)]

Out	641	Well	ID

well_	LD
475	270.785974
2821	285.971884
2841	506.750799
2977	282.519542
4545	439.690000
4689	267.553524
4793	271.752307
4987	255.620635
5060	368.900000
5557	351.206317
5717	700.890000
5788	422.070000
6137	309.920000
6583	339.300000
7007	304.690000
8051	308.880000
8522	256.610000
9362	299.530000
Name:	As, dtype: float64

A second way to do boolean and in pandas. Remember. When you split a line at a comma in a fucntion you don't need to use the . I do this to make the packets print better. You don't need to do it. But also line breaks can just make things cleaner and easier to see

In [67]: df['As'][np.logical_and(df.Drink=='Y' , df.As>250)]

Out[67]:	475 2821 2841 2977 4545 4689 4793 4987 5060 5557 5717	270.785974 285.971884 506.750799 282.519542 439.690000 267.553524 271.752307 255.620635 368.900000 351.206317 700.890000
	4689	267-553524
	5557	351.206317
	5717	700.890000
	5788	422.070000
	6137	309.920000
	6583	339.300000
	7007	304.690000
	8051	308.880000
	8522	256.610000
	9362	299.530000
	Name:	As, dtype: float64

Can we look at who drinks from their wells and if they don't drink is it beacuse it has more arsenic?

Another way to word this.

What is the average arsenic in wells people drink from?

What is the averarge arsenic in wells people don't drink from.

Use decribe

In [68]: *#wells people drink from* print('Arsenic of wells where people drink') print (df['As'][df.Drink=='Y'].describe()) # wells people don't drinkfrom print ('\nArsenic of wells where people don\'t drink') #to put the ' in the line I added a \ befo print (df['As'][df.Drink=='N'].describe()) Arsenic of wells where people drink count 336.000000 72.484421 mean 91.571489 std min 0.000000 25% 9.962403 50% 39.975000 75% 99.294435 700.890000 max Name: As, dtype: float64 Arsenic of wells where people don't drink 71.000000 count 171.105792 mean std 107.308224 1.368709 min 25% 81.614239 50% 150.250000 250.245000 75% max 473.340000 Name: As, dtype: float64

What do the results above show?

I am going to come back to groupby here and there and we will do a whole packet on it. But when your brain can think through groupby it makes things simpler. So here we are going to groupby drink and then describe As. It should do what we just did in one line and make a nicer output.

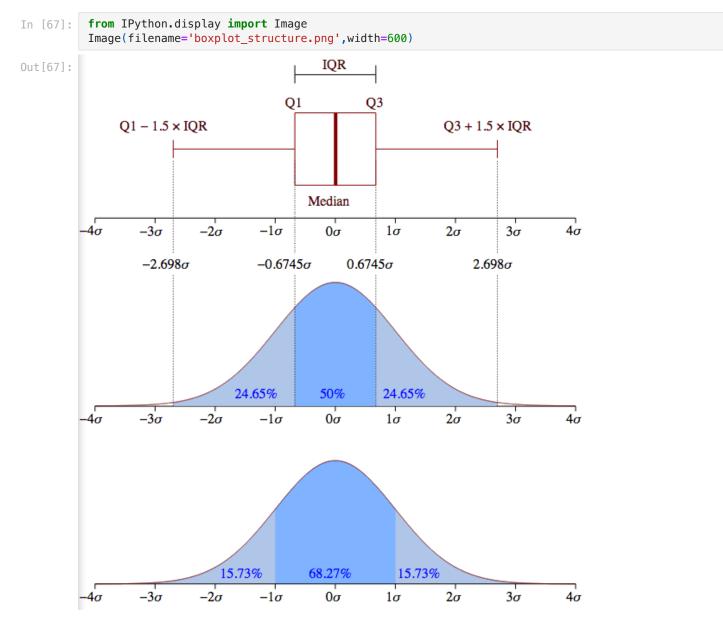
```
In [71]: df.groupby('Drink')['As'].describe()
```

	5		(2.2						
ut[71]:		count	mean	std	min	25%	50%	75%	max
	Drink								
	Ν	71.0	171.105792	107.308224	1.368709	81.614239	150.250	250.245000	473.34
	Y	336.0	72.484421	91.571489	0.000000	9.962403	39.975	99.294435	700.89
[]:									
(Could v	we disp	ly this data?)					
72]:	df.b	oxplot	(column='A	s')					
72]:	<axes< td=""><td>Subplo</td><td>t:></td><td></td><td></td><td></td><td></td><td></td><td></td></axes<>	Subplo	t:>						
	700			•					
	600								
	500			8					
	400			8					
	300 -								
	200								
	100 -								
	0								

As

Do you remember what a boxplot shows? I found this next picture on stackoverflow. No need to import. Just for your reference.

http://stackoverflow.com/questions/17725927/boxplots-in-matplotlib-markers-and-outliers



cool boxplots

But we really want two boxplots. One for people who drink and one for people who don't drink. I wasn't sure how to do it? So I googled pandas boxplot. Here are two of the links I got. See if you can figure it out! If you scroll down on the first link you should find the answer.... You will want your boxplots grouped. (only spend 2 minutes on this and I will come help you. Don't go down a rabbit hole on this. Answer at the ened)

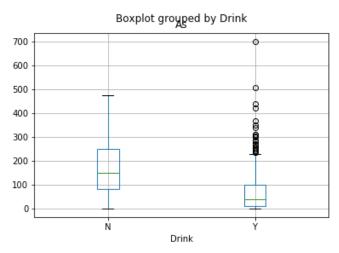
- click on the first link
- scroll down to where it says boxplots.
- Now scroll a little further to where you see the boxplots that say "grouped by x"
- look in the code.
- see if you can find a keyword argument in the parantheses that could hep you and figure out what column to pass

https://pandas.pydata.org/docs/reference/api/pandas.DataFrame.boxplot.html

http://stackoverflow.com/questions/23232989/boxplot-stratified-by-column-in-python-pandas

In [65]:

Out[65]: <matplotlib.axes._subplots.AxesSubplot at 0xc0542e8>



What difference do you notice about the arsenic concentrations of peeople drinking from their wells?

Whenever you are comparing two populations a t-test should pop into your head!

t-test or student t-test

A t-test tells you if there is a significant difference between two means. Actually it tells you the probability that they are the same. Back to our friend the p-value! The first website seems to have a good explanation.

Whenever you are comparing two means you run a t-test.

I am going to repeat this. If you ever compare two populations with a mean you need to run a t-test to see if the differences are statistically significant.

You then need to choose if it is

- 1. Paired scipy.stats.ttest_rel
- 2. not paired scipy.stats.ttest_ind

By paired we mean you repeated the measure on the same thing. Can you track the same thing across two samples. For example The exam score of the same person before and after an intervention

By not paired we mean two different populations. Imagine we fed 100 people carrots and 100 people steak and we weighed them and wanted to know if their exam scores was differnt. That is not paired and also the worst experiment ever!

In terms of our arsenic example if we measured the same wells twice it would be paired. if we measured different sets of wells it is unpaired.

Finally, if you are doing unpaired you need to decide if the groups have the same or unequal variance. It is statiscally safer to choose unequal variance. But you can always look at your variance and decide.

Your results are a t statistic and a p-value. We want our p-value less than 0.05 or 0.01 again!

Back to our wells. We will run an unparied t-test with unequal variance.

so lets pass our arrays from aboce with Arsenic for Drink=Y and Drink=N

We are asking if the difference we see with our eyes in the boxplot is statistically significant for arsenic. You need the stats to verify!

THIS WILL FAIL!

```
Out[74]: Ttest_indResult(statistic=nan, pvalue=nan)
```

It failed b/c we have NaN's in our data (Not a Number). NaN's are nice as they keep track where we don't have data.

But scipy does not handle NaN's well.

So we need to get rid of them then do the math.

In pandas terms we need to drop the NaN's using the function dropna

```
In [75]: df['As'][df.Drink=='Y'].dropna()
```

```
Out[75]:
         Well ID
         83
                    78.977470
                    28.070949
          101
          110
                    96.885674
          112
                    80.627214
         153
                    39.249817
                    26,980000
          12363
                    21.740000
          12440
          12461
                   117.820000
         12516
                     0.130000
         12654
                    17.390000
         Name: As, Length: 336, dtype: float64
         So try again!
```

```
In [76]: stats.ttest_ind(df['As'][df.Drink=='Y'].dropna()
            ,df['As'][df.Drink=='N'].dropna()
            ,equal_var=False)
```

```
Out[76]: Ttest_indResult(statistic=-7.209206229150192, pvalue=1.4829579464861492e-10)
```

That is a small p-value!!!!!

So a signifcant difference!

You could say the mean arsenic concentration is lower in well where people drink then where they don't drink (p<0.01)

What wells do people drink from?

For our final exercise. Lets put it together and get data and then see if we can plot it. I want to know the number of people who are drinking from there wells based on the arsenic concentrations. Can we do the reverse. if the arsenic is <10,10-50, and >50 what is the value counts of drinking and not drinking. I chose these numbers because 10 ppb is the EPA and WHO drinking water limit. 50 ppb is the Bangladesh drinking water limit. We see negative health effects at 10ppb. Drinking water with 10 ppb arsenic is bad for you! It increases your risk of cardiovascular disease, cancers, and death!

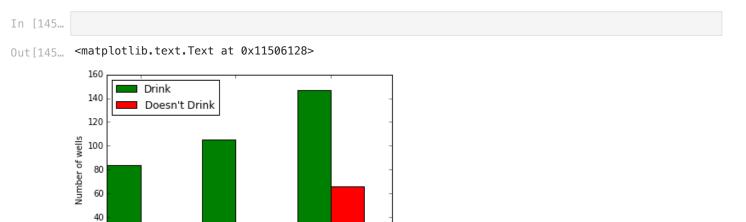
I would first just try and break the data into 3 groups and print out the results. So use your way of selecting data and select data based on the levels of arsenic. To do between 10 and 50 you will need to use an and statement and how to do those is different. you need to use a boolean function to choose two data sets! works by taking two arguments and then returning what happens the same way as if you did an and. but it works better. Remember we did this above.

Three Groups

20 0

- 1. <10 ppb arsenic
- 2. 10-50 ppb arsenic
- 3. >50 ppb arsenic
- 4. Print out the number of people drinking from wells with arsenic less than 10. you can use value_counts() and your selection method.

This is the graph we want to make



First start by counting who drinks less than 10.

10-50

Arsenic Concentration ppb

```
In [77]: print ('people drinking with arsenic <10')</pre>
```

df['Drink'][df.As<=10].value_counts()</pre>

people drinking with arsenic <10 Out[77]: Y 84 N 1

Name: Drink, dtype: int64

<10

1. Next use determine the people drinking from wells with arsenic more than 50.

>50

In [113...

```
people drinking with >50
Y 147
N 66
dtype: int64
```

1. Now use your logical_and() or & and parantheses to determine between 10 and 50.

In [115...

```
people drinking with 10-50
Y 105
N 4
dtype: int64
```

Looking at the data one by one is painful. Lets work on getting to our bar chart. This is a bad way of looking at the data. I would like to make bar plot. Here is my goal. Can we get there? Follow the next steps after the plot and see how it goes!

python does not make bar plots easy. But let's make one anyway

First lets look up bar plot. http://matplotlib.org/examples/api/barchart_demo.html This is the example on all the web pages. We can make sense of it. Lets do one step at a time. What plt.bar wants is (x,y,width). lets do it for As<10 first. Here is our data again. The x location and width become arbitrary to make it look pretty.

```
In [85]: df['Drink'][df.As<=10].value_counts()</pre>
```

```
Out[85]: Y
```

```
N 1
Name: Drink, dtype: int64
```

84

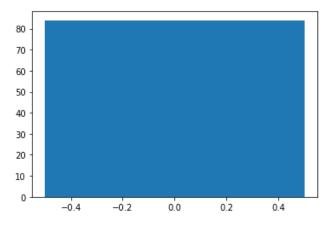
So we want to plot 84 Drink, 1 Doesn't drink. I will do it longhand first time.

remember it is ax.bar(x,y,width)

x=where to plot it and is a bit of a dumy variable y=the height of the bar width=how wide you want the bars

```
In [86]: # bar for people who drink
fig,ax=plt.subplots()
ax.bar(0,84,1)
```

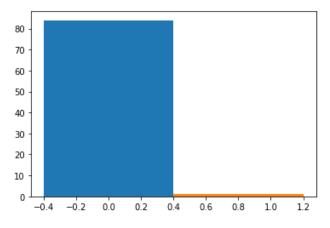
Out[86]: <BarContainer object of 1 artists>



The default width is 0.8 starting from 0. Now we need to add the doesn't drink.

```
In [88]: # bar for people who drink and don't drink
fig,ax=plt.subplots()
ax.bar(0,84,.8)
ax.bar(0.8,1,.8)
```

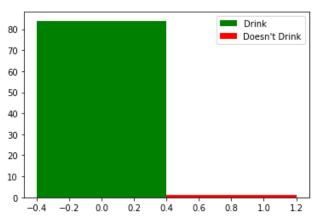
Out[88]: <BarContainer object of 1 artists>



Now we need to add colors and labels for a legend.

```
In [87]: fig,ax=plt.subplots()
ax.bar(0,84,0.8,color='g',label='Drink')
ax.bar(0.8,1,0.8,color='r',label="Doesn't Drink")
#I did double quotes so I could print the single quote
ax.legend(loc='best')
```

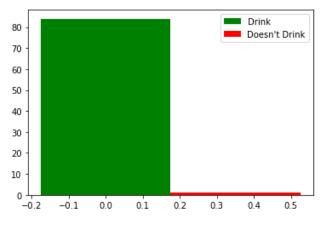




this is a disaster. We can't hard wire it all. We need to be better in our programming and be what people call pythonic. now instead of setting the x-axis to zero lets use np.arange. Then also lets set the width. We will also make the second bar start at one width

```
In [79]: fig,ax=plt.subplots()
width=0.35
xvalues=np.arange(1)
ax.bar(xvalues,84,width,color='g',label='Drink')
ax.bar(xvalues+width,1,width,color='r',label="Doesn't Drink")
#I did double quotes so I could print the single quote
ax.legend(loc='best')
```

Out[79]: <matplotlib.legend.Legend at 0x7f7fdf6d2880>



Now we are starting to make progress. But we need the other two sets of bars. We will need a set of yes and no values. so we need yes[0],yes[1],yes[2] representing our values. I would make a nump array of zeros and then fill it in. So to make a numpy array of zeros. then fill in the array. we know the length has to be three.

```
In [90]: yes=np.zeros(3)
print (yes)
```

[0. 0. 0.]

Now do the same for no. then set each one equal to the correct result that you have above where you printed out the results. don't print the results like you did above. set them to yes, no given the correct array spot. At the end you should now have yes and no set for the three levels.

In [85]:

[0. 0. 0.]

Now I will show you how to add the first yes and no

In [91]: yes=np.zeros(3)
no=np.zeros(3)

<pre>yes[0],no[0]=df['Drink'][df.As<=10].value_counts()</pre>	
print (' <mark>yes',</mark> yes)	
print ('no',no)	

yes [84. 0. 0.] no [1. 0. 0.]

Now can you do the other two?

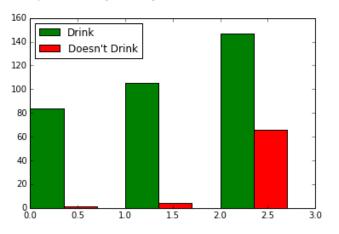
In [140...

[84. 105. 147.] [1. 4. 66.]

Now we can do a bar plot of yes and no. Go copy and past your barplot code from above. but now make the x-axis have an np.arange of 3 b/c we want 3 locations. And don't use the hardwired number put in your new yes and no arrays you just made.

In [141...

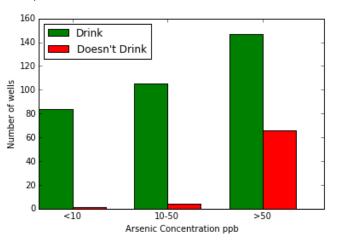
Out[141... <matplotlib.legend.Legend at 0x106ed908>



Now you are looking great with a wonderful graph. lets label everything. We just need an x-axis labeled correctly. Also, I would put all the code in one cell so it always works smoothly. If we go back to our webpage with the example we can use ax.set_xticks(xvalues+width/2) to get us the xticks we want. then we can add ax.set_xticklabels(('names','names','names')). We can also use ax.set_xlabel() and ax.set_ylabel()

```
In [144...
```

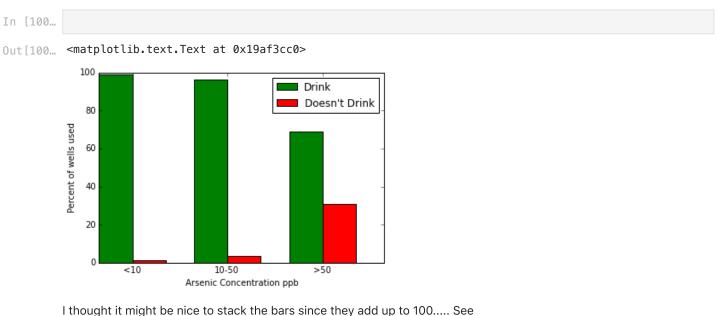
```
Out[144... <matplotlib.text.Text at 0x10f052e8>
```



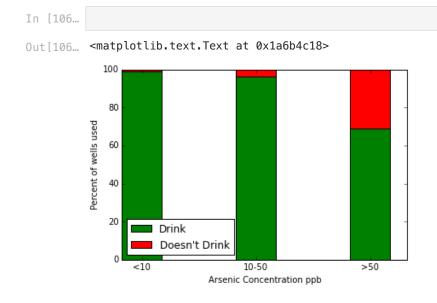
Now that is a great looking graph. You just need to add a figure caption. I would write something like

Number of wells categorized by if the respondents drink or don't drink from the well and stratified by arsenic concentration.

As a total bonus and if you have time you could change it from the number of wells to the proportion of wells in each category.



http://matplotlib.org/examples/pylab_examples/bar_stacked.html It is "easy" I used the bottom keyword. Then I removed the width offset and tweaked a few other things

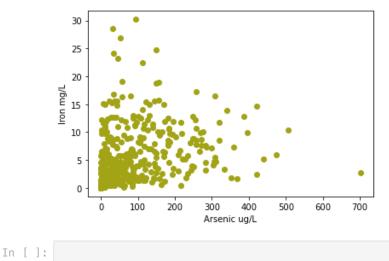


Some homework hints.

for the homework you will need to make scatter plots. They are easy to make in pandas. Here is one of Arsenic versus Iron. You can label your axes and change the color of your symbols.

```
In [92]: fig,ax=plt.subplots()
ax.scatter(df['As'],df['Fe'],c='xkcd:vomit')
ax.set_xlabel('Arsenic ug/L')
ax.set_ylabel('Iron mg/L')
```

Out[92]: Text(0, 0.5, 'Iron mg/L')



```
In []:
```

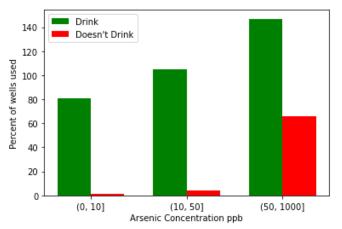
Below is a simpler method

We will learn more about this but I did advanced Python.

- 1. I defined bins
- 2. I used groubpy to group the data by if people drink
- 3. I added a cut value to also group by the bins I cut and set by.
- 4. I then filled the group by the counts.
- 5. I then unstack and transpose and flip the matrix.
- 6. Then I can plot that new data.
- 7. The fun part is I can change the bins and it automtacillay updates!

```
In [109...
          bins=[0,10,50,1000]
          df_No_Yes=df.groupby(['Drink',pd.cut(df['As'],bins)])\
                                   .As.count().unstack().transpose()
          fig,ax=plt.subplots(1,1)
          width=0.35
          xvalues=np.arange(df_No_Yes.shape[0])
          ax.bar(xvalues,df_No_Yes.Y,width,color='g',label='Drink')
          ax.bar(xvalues+width,df_No_Yes.N,width,color='r',label="Doesn't Drink")
          #I did double quotes so I could print the single quote
          ax.legend(loc='best')
          #You can try numbers 1-8 for location. see http://matplotlib.org/1.3.1/users/legend_guide.html
          ax.set_xticks(xvalues+width/2)
          ax.set_xticklabels(df_No_Yes.index.values)#('<10','10-50','>50'))
          ax.set_xlabel('Arsenic Concentration ppb')
          ax.set_ylabel('Percent of wells used')
```

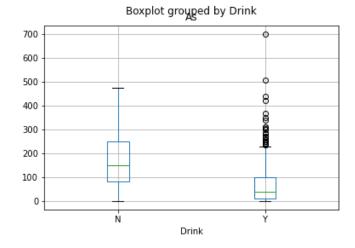
Out[109... Text(0, 0.5, 'Percent of wells used')



Answers



```
Out[73]: <AxesSubplot:title={'center':'As'}, xlabel='Drink'>
```



print ('people drinking with <10')</pre> In [79]:

df['Drink'][df.As<=10].value_counts()</pre>

people drinking with <10

```
84
Out[79]: Y
         Ν
                1
         Name: Drink, dtype: int64
```

In [80]: print ('\npeople drinking with >50 ')

df['Drink'][df.As>=50].value_counts()

people drinking with >50 Υ 147 Out[80]:

Ν 66 Name: Drink, dtype: int64

In [84]: print ('people drinking with 10-50')

df['Drink'][(df.As<=50)&(df.As>=10)].value_counts()

using np.logical_and #df['Drink'][np.logical_and(df.As<=50,df.As>=10)].value_counts()

people drinking with 10-50

4

Out[84]: Υ 105 Ν

Pandas_Well_Data

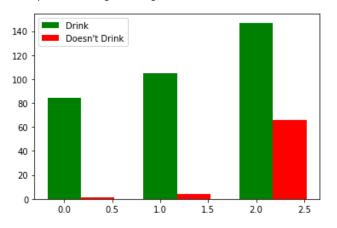
Name: Drink, dtype: int64

Pandas_Well_Data

```
In [94]: yes=np.zeros(3)
no=np.zeros(3)
yes[0],no[0]=df['Drink'][df.As<=10].value_counts()
yes[1],no[1]=df['Drink'][(df.As<=50)&(df.As>=10)].value_counts()
yes[2],no[2]=df['Drink'][df.As>=50].value_counts()
print (yes,no)
[ 84. 105. 147.] [ 1. 4. 66.]
```

In [95]: fig,ax=plt.subplots(1,1)
width=0.35
xvalues=np.arange(3)
ax.bar(xvalues,yes,width,color='g',label='Drink')
ax.bar(xvalues+width,no,width,color='r',label="Doesn't Drink") #I did double quotes so I could pr
ax.legend(loc='best')

Out[95]: <matplotlib.legend.Legend at 0x7f8862f94070>

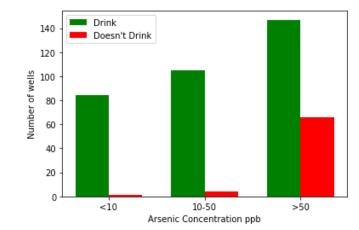


In [96]: yes=np.zeros(3)
no=np.zeros(3)
ves[0].no[0]=df

```
yes[0],no[0]=df['Drink'][df.As<=10].value_counts()
yes[1],no[1]=df['Drink'][(df.As<=50)&(df.As>=10)].value_counts()
yes[2],no[2]=df['Drink'][df.As>=50].value_counts()
fig,ax=plt.subplots()
```

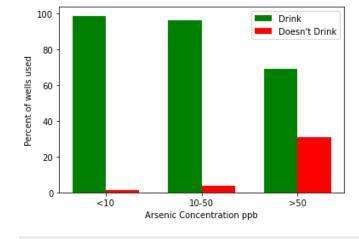
```
width=0.35
xvalues=np.arange(3)
ax.bar(xvalues,yes,width,color='g',label='Drink')
ax.bar(xvalues+width,no,width,color='r',label="Doesn't Drink")
#I did double quotes so I could print the single quote
ax.legend(loc='best')
ax.set_xticks(xvalues+width/2)
ax.set_xticklabels(('<10','10-50','>50'))
ax.set_xlabel('Arsenic Concentration ppb')
ax.set_ylabel('Number of wells')
```

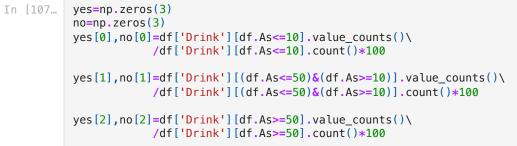
```
Out[96]: Text(0, 0.5, 'Number of wells')
```





Out[105... Text(0, 0.5, 'Percent of wells used')





```
fig,ax=plt.subplots(1,1)
width=0.35
xvalues=np.arange(3)
ax.bar(xvalues,yes,width,color='g',label='Drink')
ax.bar(xvalues,no,width,color='r',bottom=yes,label="Doesn't Drink")
#I did double quotes so I could print the single quote
ax.legend(loc=3) #You can try numbers 1-8 for location. see http://matplotlib.org/1.3.1/users/le
ax.set_xticks(xvalues+width/2)
ax.set_xticklabels(('<10','10-50','>50'))
ax.set_xlabel('Arsenic Concentration ppb')
ax.set_ylabel('Percent of wells used')
```

Out[107... Text(0, 0.5, 'Percent of wells used')

