- In [3]: #I recommend installing Anaconda for ease: #https://www.continuum.io/downloads #certain packages i.e. seaborn, do not come with the standard installation. A simple search for #how to 'install seaborn with anaconda' will easily find the installation instructions. #i.e. in this case: #http://seaborn.pydata.org/installing.html #conda install seaborn
- In [ ]: #With regards to the data, and the study during which this data was collected, rights belong to: #Friedmann, Peter. Criminal Justice Drug Abuse Treatment Studies (CJ-DATS): Step 'N Out, 2002-2006 [Unit ed States]. #ICPSR30221-v1. Ann Arbor, MI: Inter-university Consortium for Political and Social Research [distributo r], 2011-07-27. #They made the entire dataset openly available at: #http://doi.org/10.3886/ICPSR30221.v1
- In [4]: #indicate the directory where your data folder, in this case the file 'stepnout.csv', is located #in this case, it sits in a folder on the dropbox cloud. cd Dropbox/Data visualization/Course 5/ 32ec92f8b8c6d83a374ae0989bec1447 StepNOut

/Users/RI/Dropbox/Data visualization/Course 5/ 32ec92f8b8c6d83a374ae0989bec1447 StepNOut

21/03/2017 11:40

```
In [252]: #import the packages we will use
          import pandas as pd
          import numpy as np
          import seaborn as sns
          import matplotlib.pyplot as plt
          import statsmodels.formula.api as smf
          import statsmodels.stats.multicomp as multi
          import statsmodels.api as sm
          import scipy.stats
          from sklearn.cross validation import train test split
          from sklearn.tree import DecisionTreeClassifier
          from sklearn.metrics import classification report
          import sklearn.metrics
          from sklearn import datasets
          from sklearn.ensemble import ExtraTreesClassifier
          from sklearn.ensemble import RandomForestClassifier
          from sklearn import preprocessing
          from sklearn.linear model import LassoLarsCV
          from sklearn.cluster import KMeans
          %matplotlib inline
          sns.set(style="whitegrid", color codes=True)
```

```
In [192]: #load the full dataset
full_dataset = pd.read_csv('stepnout.csv')
```

In [231]: #show the first ten rows of the dataset to get an idea of the variables and their values full\_dataset.head(10)

Out[231]:

	cid	AGE	CHISP	CHETHN	cblack	casian	cnative	cwhite	cpacisl	cothrac	 arst24tlfb_9	arst25tlfb_9	alctlfb_9	alc5tlfb_
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1	2181	26.0	1.0	7.0	NaN	NaN	NaN	NaN	NaN	1.0	 0.0	0.0	1.0	1.0
2	2183	46.0	1.0	5.0	NaN	NaN	NaN	NaN	NaN	1.0	 0.0	0.0	0.0	0.0
3	2185	26.0	1.0	5.0	NaN	NaN	NaN	NaN	NaN	1.0	 0.0	0.0	1.0	1.0
4	2187	37.0	1.0	7.0	NaN	NaN	NaN	NaN	NaN	1.0	 0.0	0.0	0.0	0.0
5	2188	41.0	0.0	NaN	1.0	NaN	NaN	NaN	NaN	NaN	 0.0	0.0	1.0	0.0
6	2189	45.0	0.0	NaN	NaN	NaN	NaN	1.0	NaN	NaN	 0.0	0.0	0.0	0.0
7	2191	30.0	0.0	NaN	1.0	NaN	NaN	NaN	NaN	NaN	 0.0	0.0	1.0	0.0
8	2192	29.0	1.0	7.0	NaN	NaN	NaN	NaN	NaN	1.0	 0.0	0.0	1.0	0.0
9	2193	35.0	1.0	5.0	NaN	NaN	NaN	NaN	NaN	1.0	 0.0	0.0	0.0	0.0

10 rows × 691 columns

- In [8]: #rename all columns to small caps
  full\_dataset.rename(columns = lambda x: x.lower(), inplace=True)
- In [9]: #same result, different way of doing it
  #full\_dataset.columns = map(str.lower, full\_dataset.columns)
- In [10]: #same result, different way of doing it
  #df = df.rename(columns=lambda x: x.replace('\$', ''))

```
In [11]: #try to get all columns, one sees there are 691
full_dataset.columns
```

ahian

In [12]: #print all columns/variables
 print(full\_dataset.describe().to\_string())

/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/numpy/lib/function\_base.py:3834: RuntimeWarning: Invalid value encountered in percentile
RuntimeWarning)

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lcsf002	lcsf003	lcsf004	lcsf005	lcsf006	lcsf007	lcsf008	lcsf009	lcsf010
lcsf011	lcsf012	lcsf013	lcsfsc1	lcsfsc2	lcsfsc3	lcsf014	lcsf015	rctr
csex	dy30free d	ly180free	clive	lngliv	homeless	suprliv	livsp	jobsp
numch	minorch	livch	supch	fostch	depend	cmstat	lngmst	grade
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p13 fmrel8 prrel7 sbrel6 frchr5	majsup fmrel9 prrel8 sbrel7 frchr6	fmrel1 fmrel10 prrel9 sbrel8 frchr7	fmrel2 prrel1 prrel10 sbrel9 frchr8	fmrel3 prrel2 sbrel1 sbrel10 frchr9	fmrel4 prrel3 sbrel2 frchr1 frchr10	fmrel5 prrel4 sbrel3 frchr2 frchr11	fmrel6 prrel5 sbrel4 frchr3 gangev	fmrel7 prrel6 sbrel5 frchr4 gangc

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c10cmlf	c10cm6m	c10cm30	c11cmlf	c11cm6m	c11cm30	c12cmlf	c12cm6m	c12cm30
c13cmlf	c13cm6m	c13cm30	c14cmlf	c14cm6m	c14cm30	c14cmin	c14cmdt	c15cmlf
c15cm6m	c15cm30	c15cmin	c15cmdt	c16cmlf	c16cm6m	c16cm30	c16cmin	c16cmdt
c17cmlf	c17cm6m	c17cm30	c17cmin	c17cmdt	c18cmlf	c18cm6m	c18cm30	c18cmin
c18cmdt	c19cmlf	c19cm6m	c19cm30	c19cmin	c19cmdt	c20cmlf	c20cm6m	c20cm30
c20cmin	c20cmdt	c21cmlf	c21cm6m	c21cm30	c22cmlf	c22cm6m	c22cm30	c23cmlf
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health	painlf	dprslf	anxlf	hallulf	conclf	viollf	sidelf	sattlf
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hluc30d	agelcrk	crck6m	crck30d	age1coc	coc6m	coc30d	ag1coin	coinj30
age1hco	hcoc6m	hcoc30d	ag1hcin	hcinj30	age1hmo	hmth6m	hmth30d	ag1hmin
hminj30	age1her	her6m	her30d	ag1hein	heinj30	age1mth	mthd6m	mthd30d
ag1mtin	mtinj30	agelopi	opiat6m	opiat30d	aglopin	opinj30	age1met	meta6m
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		30.000000						
		18.000000						0000 602.00000
								1.000000 11.
		9999.000000						
		0000 9000.00		000000 9999				13.000000 150
00.000000							00000 4.0	
		000000 4.0		.000000 4				4.000000 4
$\Lambda$ $\Lambda$ $\Lambda$ $\Lambda$ $\Lambda$								
	4.000000							
4.000000	4.000000	4.000000	4.00000	0 4.00000	4.000	000 4.00	00000 4.00	$000   4.000000 \\ 0000   4.00000 \\ 4.000000   4.$

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000000 4.000000 4.000000 50.000000 30.000000 11.000000 46.000000 15.000000 2.000000 811.000000 4.000000 45.000000 66,000000 4.000000 9.000000 11,000000 60.000 59,000000 50.000000 52,000000 42,000000 10,000000 60.000000 42,000000 11,000000 80.00000 .000000 60.000000 3.000000 9.000000 7.000000 1.000000 59,000000 5.000000 999.000000 8.000000 1.000000 1,000000 8.000000 8.000000 20.000000 999.000000 105.000000 210.00000 0 280.000000 60.000000 90.000000 540.000000 4.000000 4.000000 4.000000 180.000000 180. 000000 25.000000 999.000000 999.000000 500.000000 999.000000 999.000000 999.000000 999,000000 999.000000 999.000000 999.000000 999.000000 10.000000 10.000000 1.000000 5.000000 5.000 5.000000 5.000000 5.000000 5.000000 5.000000 5,000000 5.000000 5,000000 000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5,000000 5,000000 5,00000 5.000000 5.000000 5,000000 5.000000 5,000000 5.000000 5,000000 5.000000 5.000000 000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.00000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5. 000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.00000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5,000000 5. 000000 5.000000 5.000000 5,000000 5.000000 5.000000 5.000000 5,000000 5,000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 5.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 7.000000 5.000000 276.000000 243.000000 269.000000 269.000000 1.000000 1.000000 1.000000 1.00000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 0.0 1.000000 1.000000 1.000000 0.0 1.000000 0.0 0.0 1.000000 0.0 1.000000 1.000000 1.000000 1.000000 1.000000 1,000000 1.000000 1.000000 1.0 00000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 0.0 0.0 0.0 1.000000 1.00 0000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 0.0 0.0 1.000000 1.000000 1.000000 0.0 1.000000 0.0 1.000000 0.0 0.0 0.0 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 9.000000 1.000000 9.000000 1.000000 13.000000 1.000000

```
In [13]: #initial exploratory data analysis
#frequency distributions for categorical variables
#graphical representations
#calculations of center and spread for quantitative variables
```

In [15]: #we take a look at the first 30 rows
reduced\_dataset.head(30)

Out[15]:

		cond	csex	age	clive	majsup	allarrests	anyarrest	alldrugs	anydrugs	allcrimes	anycrime	arrest_9mo	reincarc_9mo	num
(	)	1	1	30.0	5	11.0	2.0	1.0	2.0	1.0	2.0	1.0	1	1	2
-		1	1	26.0	5	11.0	0.0	0.0	1.0	1.0	1.0	1.0	1	1	1
2	2	1	1	46.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
3	3	0	1	26.0	4	9.0	0.0	0.0	1.0	1.0	2.0	1.0	0	0	0
4	ŀ	0	1	37.0	4	6.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
į	5	1	1	41.0	4	9.0	0.0	0.0	1.0	1.0	1.0	1.0	0	0	0
6	6	1	1	45.0	5	11.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
7	7	0	1	30.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
8	3	0	1	29.0	5	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
Ś	,	0	1	35.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0

10	0	1	35.0	4	9.0	NaN	NaN	NaN	NaN	NaN	NaN	0	0	0
11	0	1	37.0	4	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
12	0	1	24.0	4	9.0	0.0	0.0	1.0	1.0	0.0	0.0	0	0	0
13	1	1	24.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
14	1	1	32.0	4	9.0	0.0	0.0	1.0	1.0	0.0	0.0	0	0	0
15	0	1	54.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
16	1	1	24.0	4	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
17	1	1	25.0	5	9.0	4.0	1.0	1.0	1.0	4.0	1.0	1	1	1
18	1	1	29.0	4	9.0	0.0	0.0	0.0	0.0	1.0	1.0	1	1	1
19	1	1	41.0	4	1.0	0.0	0.0	1.0	1.0	1.0	1.0	0	0	0
20	1	1	30.0	5	5.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
21	0	1	21.0	5	9.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
22	1	1	37.0	4	1.0	0.0	0.0	3.0	1.0	1.0	1.0	1	1	2
23	1	1	39.0	5	1.0	0.0	0.0	1.0	1.0	0.0	0.0	0	0	0
24	0	1	28.0	4	9.0	0.0	0.0	1.0	1.0	0.0	0.0	0	0	0
25	0	1	34.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
26	1	1	19.0	4	9.0	0.0	0.0	0.0	0.0	1.0	1.0	0	0	0
27	1	1	41.0	4	1.0	0.0	0.0	9.0	1.0	0.0	0.0	0	0	0
28	1	1	24.0	5	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0	0	0
29	0	1	35.0	4	1.0	NaN	NaN	NaN	NaN	NaN	NaN	0	0	0

```
In [16]: #we check the length of the dataset
         print(len(reduced dataset))
         476
In [17]: #we confirm the number of columns
         print(len(reduced dataset.columns))
         17
         #we rename some of the columns to give them more meaningful names
In [18]:
         reduced dataset.rename(columns = {'csex': 'sex', 'clive': 'living situation', 'majsup': 'support'}, inpl
         ace=True)
         /Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/pandas/core/frame.py:2754: SettingWithCopyWarni
         A value is trying to be set on a copy of a slice from a DataFrame
         See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
         g-view-versus-copy
           **kwargs)
In [19]: #convert all variables to numeric ones
```

In [193]: #mark all variables as numeric data, and signify, for the relevant ones, that they are categorical #rather than quantitative variables #errors='coerce' tells pandas to return invalid values as NaN rather than as the input values themselves reduced dataset['cond'] = pd.to numeric(reduced dataset['cond'], errors='coerce').astype('category') reduced dataset['sex'] = pd.to numeric(reduced dataset['sex'], errors='coerce').astype('category') reduced dataset['age'] = pd.to numeric(reduced dataset['age'], errors='coerce') reduced dataset['living situation'] = pd.to numeric(reduced dataset['living situation'], errors='coerce' ).astype('category') reduced dataset['support'] = pd.to numeric(reduced dataset['support'], errors='coerce').astype('category ') reduced dataset['allarrests'] = pd.to numeric(reduced dataset['allarrests'], errors='coerce') reduced dataset['anyarrest'] = pd.to numeric(reduced dataset['anyarrest'], errors='coerce').astype('cate gory') reduced dataset['alldrugs'] = pd.to numeric(reduced dataset['alldrugs'], errors='coerce') reduced dataset['anydrugs'] = pd.to numeric(reduced dataset['anydrugs'], errors='coerce').astype('catego') ry') reduced dataset['allcrimes'] = pd.to numeric(reduced dataset['allcrimes'], errors='coerce') reduced dataset['anycrime'] = pd.to numeric(reduced dataset['anycrime'], errors='coerce').astvpe('catego') ry') reduced dataset['arrest 9mo'] = pd.to numeric(reduced dataset['arrest 9mo'], errors='coerce').astype('ca tegory') reduced dataset['reincarc 9mo'] = pd.to numeric(reduced dataset['reincarc 9mo'], errors='coerce').astype ('category') reduced dataset['num arrest'] = pd.to numeric(reduced dataset['num arrest'], errors='coerce') reduced dataset['num reincarc'] = pd.to numeric(reduced dataset['num reincarc'], errors='coerce') reduced dataset['violent charge'] = pd.to numeric(reduced dataset['violent charge'], errors='coerce').as type('category')

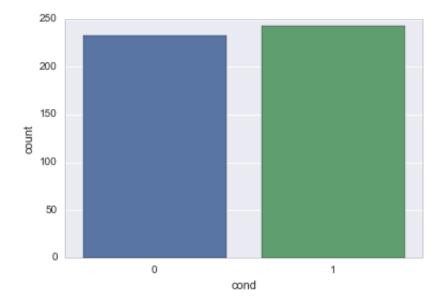
reduced dataset['property charge'] = pd.to numeric(reduced dataset['property charge'], errors='coerce').

astype('category')

```
In [21]: #now let's look at the counts and frequency distributions for these variables
          #for this, we will aim to get an idea of the shape, center, and spread of these variables
          #we will analyze the shape visually by checking for modality and skewness
          #we will check for measure of center such as mean, median and mode
          #we will check the spread through the standard deviation
In [194]: reduced dataset['cond'].value counts(sort=False, normalize = True)
Out[194]: 0
               0.489496
               0.510504
          Name: cond, dtype: float64
In [148]: reduced dataset['cond'].describe()
Out[148]: count
                    476
                      2
          unique
                      1
          top
          freq
                    243
          Name: cond, dtype: int64
 In [24]: #rules for visualizing data:
          #for visualizing a variable:
          #if it is categorical we use a bar chart i.e. sns's countplot function
          #if it is quantitative, we can combine a kernel density estimate and a histogram with sns's
          #distplot function
          #for visualizing two variables:
          # C-C: bivariate bar graph with sns factorplot
          # C-O: bivariate bar graph with sns factorplot
          # Q-Q: scatterplot with sns regplot
```

Step+N+Out

- In [25]: #given that the study condition is a categorical variable, we use a count plot to visualize it.
  sns.countplot(x='cond', data=reduced\_dataset)
- Out[25]: <matplotlib.axes.\_subplots.AxesSubplot at 0x1182c91d0>



- In [26]: #we check the counts per each value of the variable. sort=False tells pandas not to sort the results #by values. normalize = True tells it to return the relative frequencies rather than the absolute counts reduced\_dataset['sex'].value\_counts(sort=False, normalize=True)
- Out[26]: 1 0.829832 2 0.170168

Name: sex, dtype: float64

In [27]: #print all age values as a list to look through them
lis = [x for x in reduced\_dataset['age']]
print(lis)

[30.0, 26.0, 46.0, 26.0, 37.0, 41.0, 45.0, 30.0, 29.0, 35.0, 35.0, 37.0, 24.0, 24.0, 32.0, 54.0, 24.0,25.0, 29.0, 41.0, 30.0, 21.0, 37.0, 39.0, 28.0, 34.0, 19.0, 41.0, 24.0, 35.0, 27.0, 27.0, 40.0, 35.0, 2 3.0, 36.0, 36.0, 40.0, 24.0, 29.0, 22.0, 34.0, 23.0, 35.0, 36.0, 34.0, 40.0, 23.0, 44.0, 23.0, 28.0, 29 .0, 45.0, 42.0, 41.0, 39.0, 35.0, 30.0, 31.0, 24.0, 26.0, 32.0, 24.0, 25.0, 34.0, 28.0, 39.0, 27.0, 31. 0, 33.0, 27.0, 36.0, 33.0, 31.0, 24.0, 22.0, 33.0, 22.0, 35.0, 33.0, 29.0, 31.0, 45.0, 24.0, 39.0, 38.0 , 44.0, 29.0, 36.0, 33.0, 46.0, 27.0, 22.0, 25.0, 31.0, 22.0, 38.0, 36.0, 35.0, nan, 22.0, nan, 36.0, 3 8.0, 43.0, 36.0, 38.0, 42.0, 44.0, 35.0, 44.0, 24.0, 27.0, 23.0, 41.0, 35.0, 25.0, 34.0, 45.0, 34.0, 24 .0, 35.0, 59.0, 36.0, 30.0, 40.0, 20.0, 35.0, 32.0, 26.0, 27.0, 25.0, 28.0, 27.0, 30.0, 38.0, 41.0, 25. 0, 40.0, 42.0, 34.0, 27.0, 36.0, 29.0, 27.0, 50.0, 43.0, 29.0, 44.0, 42.0, 25.0, 28.0, 41.0, 22.0, 59.0 , 36.0, 32.0, 40.0, 25.0, 21.0, 38.0, 34.0, 40.0, 28.0, 25.0, 37.0, 41.0, 40.0, 24.0, 30.0, 37.0, 29.0, 29.0, 49.0, 37.0, 21.0, 26.0, 40.0, 28.0, 37.0, 38.0, 40.0, 28.0, 42.0, 40.0, 39.0, 24.0, 21.0, 46.0, 3 2.0, 26.0, 24.0, 31.0, 34.0, 26.0, 27.0, 26.0, 26.0, 28.0, 31.0, 50.0, 29.0, 35.0, 21.0, 22.0, 38.0, 29 .0, 42.0, 29.0, 45.0, 28.0, 42.0, 38.0, 23.0, 29.0, 41.0, 39.0, 37.0, 26.0, 22.0, 29.0, 30.0, 26.0, 47. 0, 47.0, 25.0, 32.0, 28.0, 38.0, 29.0, 24.0, 42.0, 22.0, 26.0, 46.0, 34.0, 44.0, 30.0, 23.0, 23.0, 30.0 , 23.0, 20.0, 46.0, 30.0, 24.0, 44.0, 34.0, 40.0, 24.0, 37.0, 37.0, 43.0, 48.0, 44.0, 29.0, 41.0, 22.0, 52.0, 42.0, 28.0, 51.0, 26.0, 43.0, 22.0, 46.0, 47.0, 43.0, 46.0, 23.0, 36.0, 24.0, 25.0, 47.0, 39.0, 3 1.0, 33.0, 35.0, 25.0, 38.0, 36.0, 42.0, 26.0, 44.0, 50.0, 36.0, 32.0, 47.0, 36.0, 29.0, 33.0, 44.0, 27 .0, 29.0, 46.0, 22.0, 37.0, 25.0, 39.0, 47.0, 30.0, 26.0, 40.0, 25.0, 27.0, 45.0, 22.0, 23.0, 28.0, 28. 0, 46.0, 44.0, 35.0, 41.0, 27.0, 43.0, 38.0, 38.0, 30.0, 40.0, 35.0, 33.0, 51.0, 32.0, 41.0, 28.0, 28.0 , 26.0, 31.0, 35.0, 33.0, 33.0, 31.0, 31.0, 43.0, 44.0, 29.0, 35.0, 36.0, 28.0, 56.0, 40.0, 25.0, 39.0, 29.0, 49.0, 22.0, 34.0, 20.0, 46.0, 42.0, 23.0, 24.0, 23.0, 22.0, 36.0, 50.0, 32.0, 24.0, 24.0, 30.0, 2 7.0, 34.0, 30.0, 28.0, 34.0, 24.0, 42.0, 36.0, 52.0, 20.0, 40.0, 37.0, 25.0, 24.0, 40.0, 33.0, 26.0, 25 .0, 27.0, 40.0, 47.0, 27.0, 35.0, 31.0, 51.0, 28.0, 36.0, 28.0, 28.0, 39.0, 40.0, 37.0, 36.0, 38.0, 48. 0, 18.0, 25.0, 32.0, 27.0, 43.0, 44.0, 27.0, 44.0, 19.0, 43.0, 27.0, 34.0, 34.0, 40.0, 34.0, 48.0, 49.0 , 44.0, 43.0, 33.0, 26.0, 32.0, 25.0, 39.0, 43.0, 42.0, 40.0, 44.0, 23.0, 31.0, 42.0, 34.0, 45.0, 57.0, 18.0, 23.0, 21.0, 21.0, 25.0, 29.0, 43.0, 59.0, 25.0, 22.0, 38.0, 41.0, 21.0, 49.0, 34.0, 48.0, 48.0, 4 6.0, 55.0, 50.0, 54.0, 54.0, 61.0, 50.0, 44.0, 21.0, 45.0, 45.0, 43.0, 51.0, 45.0, 19.0, 51.0, 24.0, 37 .0, 18.0, 33.0, 42.0, 42.0, 42.0, 24.0, 37.0, 31.0, 41.0, 39.0, 42.0]

In [28]: #the describe request gives us the count, mean, std, min, max, as well as the quartiles for the
#respective value distribution
reduced\_dataset['age'].dropna().describe()

Out[28]: count 474.000000 34.118143 mean std 8.836234 min 18.000000 25% 27.000000 50% 34.000000 41.000000 75% 61.000000 max

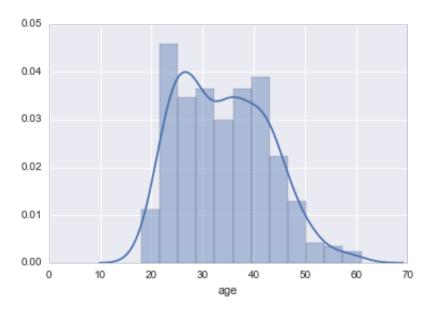
Name: age, dtype: float64

In [29]: #given that age is a quantitative variable, we use a distplot to visualize it.
sns.distplot(reduced dataset['age'].dropna())

/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/statsmodels/nonparametric/kdetools.py:20: Visib leDeprecationWarning: using a non-integer number instead of an integer will result in an error in the future

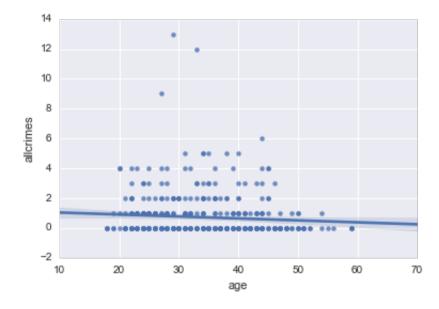
$$y = X[:m/2+1] + np.r_[0,X[m/2+1:],0]*1j$$

Out[29]: <matplotlib.axes.\_subplots.AxesSubplot at 0x1182729e8>



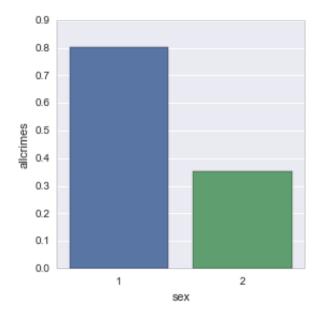
In [32]: #We use a regplot to plot two quantitative variables, age and allcrimes, while also having a regression
line suggesting
#any association present
sns.regplot(x='age', y='allcrimes', data=reduced\_dataset)

Out[32]: <matplotlib.axes.\_subplots.AxesSubplot at 0x1187067f0>



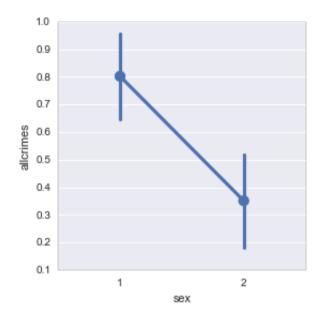
In [33]: #categorical explanatory variable 'sex' and quantitative response variable 'allcrimes'
sns.factorplot(x='sex', y='allcrimes', data=reduced\_dataset, kind='bar', ci=None)

Out[33]: <seaborn.axisgrid.FacetGrid at 0x11bd14d68>



In [38]: sns.factorplot(x='sex', y='allcrimes', kde='bar', data=reduced\_dataset)

Out[38]: <seaborn.axisgrid.FacetGrid at 0x11c365978>



In [39]: #before starting to manipulate the dataset itself, we make a copy, and will work on the copy
#rather than the original reduced dataset
rdc = reduced\_dataset.copy()

In [40]: ct1 = rdc.groupby('anyarrest').size()\*100/len(rdc['anyarrest'])
 print(ct1)

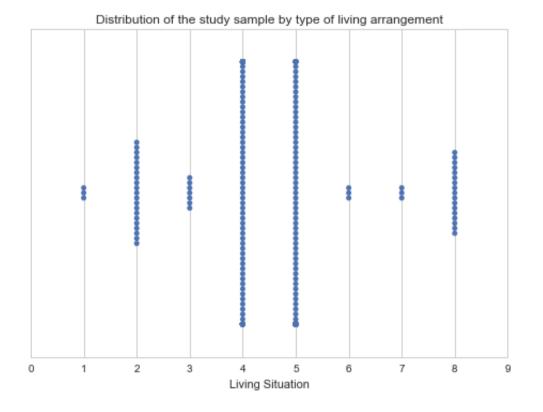
## anyarrest

0.0 69.327731 1.0 23.319328 dtype: float64

```
In [41]: rdc['anyarrest'].value counts(sort=False, dropna=False, normalize=True)
 Out[41]:
           0.0
                  0.693277
                  0.233193
           1.0
          NaN
                  0.073529
          Name: anyarrest, dtype: float64
 In [42]: rdc['cond'].isnull().value counts()
Out[42]: False
                   476
          Name: cond, dtype: int64
In [310]: rdc['anyarrest'].isnull().sum()
Out[310]: 35
 In [46]: rdc['anyarrest'] = rdc['anyarrest'].replace(11, np.nan)
 In [47]: rdc['anyarrest'].value counts()
Out[47]: 0.0
                 330
          1.0
                 111
          Name: anyarrest, dtype: int64
 In [48]: living dic = \{1: 1, 2:2, 3:1, 4:1, 5:1, 6:1, 7:1, 8:2\}
 In [49]: rdc['living situation'] = rdc['living situation'].map(living dic)
```

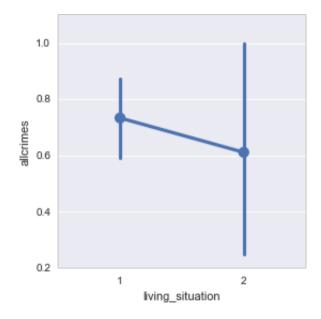
```
In [152]: sns.swarmplot('living_situation', data=reduced_dataset)
    plt.xlabel('Living Situation')
    plt.title('Distribution of the study sample by type of living arrangement')
```

Out[152]: <matplotlib.text.Text at 0x11dc74ba8>



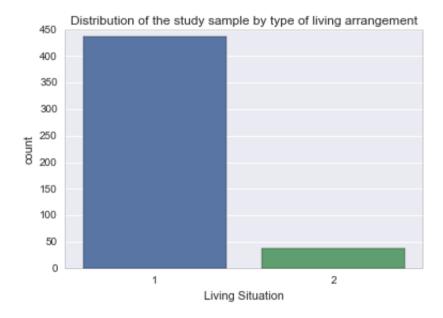
In [142]: sns.factorplot(x='living\_situation', y='allcrimes', data=rdc)

Out[142]: <seaborn.axisgrid.FacetGrid at 0x11d796c50>



```
In [149]: sns.countplot('living_situation', data=rdc)
    plt.xlabel('Living Situation')
    plt.title('Distribution of the study sample by type of living arrangement')
```

#### Out[149]: <matplotlib.text.Text at 0x11dac9080>



In [140]: rdc['age'].describe()

Out[140]: count 473.0 unique 4.0 top 20.0 freq 178.0

Name: age, dtype: float64

```
In [51]: def age group(x):
              if x < 30:
                  return 20
              elif x < 40:
                  return 30
              elif x < 50:
                  return 40
              elif x < 61:
                  return 50
In [52]: rdc['age'] = rdc['age'].map(age group)
In [53]: rdc['age']
Out[53]: 0
                 30.0
         1
                 20.0
         2
                 40.0
          3
                 20.0
                 30.0
                 40.0
         5
                 40.0
                 30.0
         8
                 20.0
                 30.0
         9
         10
                 30.0
         11
                 30.0
         12
                20.0
         13
                 20.0
         14
                30.0
         15
                 50.0
         16
                20.0
         17
                20.0
         18
                 20.0
         19
                 40.0
```

20	30.0
21	20.0
22	30.0
23	30.0
24	20.0
25	30.0
26	20.0
27	40.0
28	20.0
29	30.0
	• • •
446	40.0
447	40.0
448	50.0
449	50.0
450	50.0
451	50.0
452	NaN
453	50.0
454	40.0
455	20.0
456	40.0
457	40.0
458	40.0
459	50.0
460	40.0
461	20.0
462	50.0
463	20.0
464	30.0
465	20.0
466	30.0
467	40.0
468	40.0
469	40.0

> 20.0 470 471 30.0 472 30.0 473 40.0 474 30.0 475 40.0 Name: age, dtype: float64

In [141]: #describing the dataset/the variables of the dataset, after we group the values in the dataset #by age category rdc.groupby('age').describe()

/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/numpy/lib/function base.py:3834: RuntimeWarning

: Invalid value encountered in percentile RuntimeWarning)

## Out[141]:

		age_binned	age_unprocessed	allarrests	allcrimes	alldrugs	living_situation	num_arrest	num_reincarc
age									
	count	0.0	178.000000	168.000000	168.000000	168.000000	178.000000	178.000000	178.000000
	mean	NaN	25.044944	0.494048	0.767857	0.708333	1.073034	0.786517	0.573034
	std	NaN	2.787827	1.110629	1.555262	0.864151	0.260926	0.962247	0.749955
20.0	min	NaN	18.000000	0.000000	0.000000	0.000000	1.000000	0.000000	0.000000
20.0	25%	NaN	23.000000	NaN	NaN	NaN	1.000000	0.000000	0.000000
	50%	NaN	25.000000	NaN	NaN	NaN	1.000000	1.000000	0.000000
	75%	NaN	27.000000	NaN	NaN	NaN	1.000000	1.000000	1.000000
	max	NaN	29.000000	9.000000	13.000000	4.000000	2.000000	4.000000	4.000000
	count	0.0	153.000000	142.000000	142.000000	142.000000	153.000000	153.000000	153.000000

	mean	NaN	34.568627	0.429577	0.866197	0.767606	1.052288	0.699346	0.483660
	std	NaN	2.747642	0.886446	1.612139	1.089378	0.223337	1.026523	0.708079
30.0	min	NaN	30.000000	0.000000	0.000000	0.000000	1.000000	0.000000	0.000000
	25%	NaN	32.000000	NaN	NaN	NaN	1.000000	0.000000	0.000000
	50%	NaN	35.000000	NaN	NaN	NaN	1.000000	0.000000	0.000000
	75%	NaN	37.000000	NaN	NaN	NaN	1.000000	1.000000	1.000000
	max	NaN	39.000000	5.000000	12.000000	4.000000	2.000000	7.000000	3.000000
	count	0.0	120.000000	110.000000	110.000000	110.000000	120.000000	120.000000	120.000000
	mean	NaN	43.316667	0.245455	0.581818	0.700000	1.125000	0.400000	0.308333
	std	NaN	2.573241	0.706281	1.183850	1.324089	0.332106	0.737928	0.645551
40.0	min	NaN	40.000000	0.000000	0.000000	0.000000	1.000000	0.000000	0.000000
40.0	25%	NaN	41.000000	NaN	NaN	NaN	1.000000	0.000000	0.000000
	50%	NaN	43.000000	NaN	NaN	NaN	1.000000	0.000000	0.000000
	75%	NaN	45.000000	NaN	NaN	NaN	1.000000	1.000000	1.000000
	max	NaN	49.000000	6.000000	6.000000	9.000000	2.000000	5.000000	5.000000
	count	0.0	22.000000	19.000000	19.000000	19.000000	22.000000	22.000000	22.000000
	mean	NaN	53.000000	0.263158	0.157895	0.473684	1.090909	0.181818	0.181818
	std	NaN	3.207135	0.561951	0.374634	0.841191	0.294245	0.394771	0.394771
50.0	min	NaN	50.000000	0.000000	0.000000	0.000000	1.000000	0.000000	0.000000
30.0	25%	NaN	50.250000	NaN	NaN	NaN	1.000000	0.000000	0.000000
	50%	NaN	51.500000	NaN	NaN	NaN	1.000000	0.000000	0.000000

	75%	NaN	54.750000	NaN	NaN	NaN	1.000000	0.000000	0.000000
	max	NaN	59.000000	2.000000	1.000000	3.000000	2.000000	1.000000	1.000000

```
In [55]: #quartile split
         rdc['age unprocessed'] = reduced dataset['age']
         print('Age - 4 categories - guartiles')
         age binned = rdc['age binned'] = pd.qcut(rdc['age unprocessed'], 4, labels = ["1=0%tile", "2=25%tile",
                                                           "3=50%tile", "4=75%tile"]).value counts(sort=False, dro
         pna=True)
         print(age binned)
         Age - 4 categories - quartiles
         1=0%tile
                      135
         2=25%tile
                      115
         3=50%tile
                      116
         4=75%tile
                      108
         Name: age unprocessed, dtype: int64
         /Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/pandas/indexes/category.py:118: RuntimeWarning:
         Values and categories have different dtypes. Did you mean to use
         'Categorical.from codes(codes, categories)'?
           data = Categorical(data, categories=categories, ordered=ordered)
         /Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/pandas/indexes/category.py:118: RuntimeWarning:
         None of the categories were found in values. Did you mean to use
         'Categorical.from codes(codes, categories)'?
           data = Categorical(data, categories=categories, ordered=ordered)
```

```
In [56]: #Hypothesis testing:
         #1. specify the null hypothesis and the alternate hypothesis
         #2. choose a sample
         #3. assess the evidence
         #4. draw conclusions
         #Definition: assessing evidence provided by the data in favor or against each hypothesis
         #about the population
         #a result is statistically significant if it is unlikely to have occurred by chance
         #p value is also the type 1 error rate: the number of times we would be wrong in rejecting the
         #null hypothesis when it is true
         #p=0.03: if we reject the null hypothesis, we would be correct 97/100 times.
         #Bivariate statistical tools:
         #ANOVA; chi-square; correlation coefficient
         #ANOVA F Test: are the differences among the sample means due to true differences among the
         #population means, or merely due to sampling variability
         #F is the variation among samples means divided by the variation within groups
         #for explanatory variables with multiples levels, F test and p value do not tell us why the
         #group means are not equal, or how. there are many ways in which this can be the case.
         #before performing these analyses, one needs to use the .dropna() function to include only
         #valid data
```

# In [57]: #we will test the null hypothesis that the study condition (subject or control) and number of crimes are not related. test1\_data = rdc[['cond', 'allcrimes']].dropna() len(test1\_data)

Out[57]: 441

```
In [58]: test1 = smf.ols(formula='allcrimes ~ C(cond)', data=test1_data).fit()
    print(test1.summary())
```

===========	=======	==========	=========	:========	=========	====
Dep. Variable:		allcrimes	R-square	ed:	0	.005
Model:		OLS	Adj. R-s	squared:	0.002	
Method:	L	east Squares	F-statis	stic:	2	.019
Date:	Thu,	09 Mar 2017	Prob (F-	-statistic):	0	.156
Time: 07:11:26			Log-Like	elihood:	<b>-</b> 79	1.21
No. Observations: 441			AIC:		1	586.
Df Residuals: 439			BIC:		1	595.
Df Model:		1				
Covariance Type	:	nonrobust				
=======================================		========	=======			=====
	coef	std err	t	P> t	[95.0% Conf.	<pre>Int.]</pre>
Intercept	0.8241	0.099	8.304	0.000	0.629	1.019
C(cond)[T.1]	-0.1974	0.139	-1.421	0.156	-0.470	0.076
Omnibus:		382.656	 Durbin-W	======================================	 1	•=== •930
<pre>Prob(Omnibus):</pre>		0.000	Jarque-E	Jarque-Bera (JB):		.014
Skew:		3.695	Prob(JB):		0.00	
Kurtosis:		23.831	Cond. No.		2.64	
=======================================	=======	========	=======			====

OLS Regression Results

#### Warnings:

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```
In [59]: #now we examine the means and stds
         grouped1 mean = test1 data.groupby('cond').mean()
         print(grouped1_mean)
               allcrimes
         cond
         0
                0.824074
                0.626667
In [60]: grouped1 std = test1 data.groupby('cond').std()
         print(grouped1 std)
               allcrimes
         cond
         0
                1.710861
                1.166190
In [61]: #we repeat the same analysis with arrests
         test2 data = rdc[['cond', 'allarrests']].dropna()
         test2 = smf.ols(formula = 'allarrests ~ C(cond)', data=test2 data).fit()
         print(test2.summary())
```

Step+N+Out

#### OLS Regression Results

===========	=======	=========	=======			=====	
Dep. Variable:		allarrests	R-square	ed:	0.001		
Model:		OLS	Adj. R-s		-0.002		
Method:	L	east Squares	F-statis	tic:	0	.2401	
Date:	Thu,	09 Mar 2017	Prob (F-	statistic):	0.624		
Time:		07:11:26	Log-Like	elihood:	-593.82		
No. Observations	No. Observations:		AIC:			1192.	
Df Residuals:		439	BIC:			1200.	
Df Model:							
Covariance Type:		nonrobust					
=======================================	coef	std err	t	P> t	======== [95.0% Conf	. Int.]	
Intercept	0.4213	0.063	6.642	0.000	0.297	0.546	
C(cond)[T.1]		0.089	-0.490		-0.218	0.131	
Omnibus:		408.823	====== Durbin-W	======================================	=======	==== 1.861	
Prob(Omnibus):		0.000	Jarque-E	sera (JB):	1111	7.974	
Skew:		4.036	Prob(JB)	:	0.00		
Kurtosis:		26.236	Cond. No	) <b>.</b>		2.64	

#### Warnings:

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified.

```
In [62]: grouped2_mean = test2_data.groupby('cond').mean()
    print(grouped2_mean)
```

allarrests cond 0 0.421296 1 0.377778

#### OLS Regression Results

		OLS Regles:	sion Results	· -=======		
Dep. Variable: Model: Method: Date: Time: No. Observations: Df Residuals: Df Model: Covariance Type:	Thu, 09 Mar 2017 07:11:26 439 435 3 nonrobust		R-squared: Adj. R-squ F-statisti Prob (F-st Log-Likeli AIC: BIC:	ared: .c: :atistic):	0.0 0.0 1.8 0.1 -786. 158	005 302 .46 67
	coef			P> t	======================================	Int.]
Intercept C(age)[T.30.0] C(age)[T.40.0] C(age)[T.50.0]	0.0983 -0.1860	0.166 0.179	0.591 -1.040	0.555 0.299	-0.538	0.989 0.425 0.166 0.084
Omnibus: Prob(Omnibus): Skew: Kurtosis:		382.848 0.000 3.709 24.196	Durbin-Wat Jarque-Ber Prob(JB): Cond. No.	a (JB):		

# Warnings:

```
In [65]: #given that we have an explanatory categorical variable with multiple levels, we use the
    #tuckey hsd test
    tuckey1 = multi.MultiComparison(test3_data['allcrimes'], test3_data['age'])
    res1 = tuckey1.tukeyhsd()
    print(res1.summary())
```

Multiple Comparison of Means - Tukey HSD, FWER=0.05

\_\_\_\_\_

```
group1 group2 meandiff lower upper reject

20.0 30.0 0.0983 -0.3305 0.5272 False
20.0 40.0 -0.186 -0.6475 0.2754 False
20.0 50.0 -0.61 -1.5206 0.3007 False
30.0 40.0 -0.2844 -0.7622 0.1935 False
30.0 50.0 -0.7083 -1.6274 0.2108 False
40.0 50.0 -0.4239 -1.3586 0.5108 False
```

- In [66]: #We will now test two other hypotheses:
  #Hypothesis(0)(a): the study condition (0 or 1) and the committing of a crime are independent
  #i.e. there is no relationship between them
  #Hypothesis(0)(b): there is no relationship between age and being arrested during the study
  #period
- In [67]: #contingency table of observed counts
  #when creating contingency tables, we put the response variable first (therefore vertical in
  #table), and the explanatory variable second, therefore horizontal at the top of the table.
  ct1 = pd.crosstab(rdc['anycrime'], rdc['cond'])
  print(ct1)

```
cond     0     1
anycrime
0.0      139     154
1.0      77     71
```

```
In [68]: #column percentages
         colsum = ct1.sum(axis=0)
         colpct = ct1/colsum
         print(colpct)
         cond
                          0
                                    1
         anvcrime
         0.0
                   0.643519 0.684444
         1.0
                   0.356481 0.315556
In [69]: #chi square test
         print('chi-square value, p value, expected counts')
         cs1 = scipy.stats.chi2 contingency(ct1)
         print(cs1)
         chi-square value, p value, expected counts
         (0.65446000714878627, 0.41852260938554675, 1, array([[ 143.51020408, 149.48979592],
                [ 72.48979592, 75.51020408]]))
In [70]: #now the second hypothesis test
         rdc['age'] = rdc['age'].astype('category')
         rdc['anyarrest'] = rdc['anyarrest'].astype('category')
In [71]: #contingency table of observed counts
         ct2 = pd.crosstab(rdc['anyarrest'], rdc['age'])
         print(ct2)
         age
                    20.0 30.0 40.0 50.0
         anyarrest
         0.0
                           105
                                        15
                     118
                                  90
                      50
                                  20
         1.0
                            37
```

```
In [72]: #column percentages
         colsum2 = ct2.sum(axis=0)
         colpct2 = ct2/colsum2
         print(colpct2)
                        20.0
                                  30.0
                                            40.0
                                                      50.0
         age
         anvarrest
         0.0
                    0.702381 0.739437 0.818182 0.789474
         1.0
                    0.297619 0.260563 0.181818 0.210526
In [73]: #chi square test
         print('chi-square value, p value, expected counts')
         cs2 = scipy.stats.chi2 contingency(ct2)
         print(cs2)
         chi-square value, p value, expected counts
         (4.9451035874289762, 0.17586134722188973, 3, array([[ 125.52164009, 106.09567198,
                                                                                             82.18678815,
                                                                                                            14.
         195899771,
                [ 42.47835991, 35.90432802, 27.81321185, 4.80410023]]))
In [74]: #We therefore cannot reject the null hypothesis, but given that the explanatory variable has several lev
         els,
         #we cannot know why the null hypothesis was not rejected
         #we therefore perform a 2 by 2 comparison
In [75]: sub1 = rdc.copy()
         recode1 = \{20:20, 30:30\}
         sub1['comp1v'] = sub1['age'].map(recode1)
```

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```
In [153]: ct3 = pd.crosstab(sub1['cond'], sub1['comp1v'])
          print(ct3)
          comp1v 20.0 30.0
          cond
          0
                    95
                          71
          1
                    83
                          82
  In [ ]: #column percentages
          colsum3 = ct3.sum(axis=0)
          colpct3 = ct3 / colsum3
          print(colpct3)
  In [ ]: #chi square test
          print('chi square value, p value, expected values')
          cs3 = scipy.stats.chi2 contingency(ct3)
          print(cs3)
  In [ ]: recode2 = {20:20, 40:40}
          sub1['comp2v'] = sub1['age'].map(recode2)
  In [ ]: ct4 = pd.crosstab(sub1['anyarrest'], sub1['comp2v'])
          print(ct4)
  In [ ]: colsum4 = ct4.sum(axis=0)
          colpct4 = ct4/colsum4
          print(colpct4)
  In [ ]: print('chi square value, p value, expected values')
          cs4 = scipy.stats.chi2 contingency(ct4)
          print(cs4)
```

Step+N+Out

```
In [ ]: recode3 = {20:20, 50:50}
        sub1['compv3'] = sub1['age'].map(recode3)
In [ ]: ct5 = pd.crosstab(sub1['anyarrest'], sub1['compv3'])
        print(ct5)
In [ ]: colsum5 = ct5.sum(axis=0)
        colpct5 = ct5/colsum5
        print(colpct5)
In [ ]: print('chi square value, p value, expected values')
        cs5 = scipy.stats.chi2_contingency(ct5)
        print(cs5)
In [ ]: recode4 = \{30:30, 40:40\}
        sub1['compv4'] = sub1['age'].map(recode4)
In [ ]: ct6 = pd.crosstab(sub1['anyarrest'], sub1['compv4'])
        print(ct6)
In [ ]: colsum6 = ct6.sum(axis=0)
        colpct6 = ct6 / colsum6
        print(colpct6)
In [ ]: print('chi square value, p value, expected values')
        cs6 = scipy.stats.chi2 contingency(ct6)
        print(cs6)
In [ ]: recode6 = {30:30, 50:50}
        sub1['compv6'] = sub1['age'].map(recode6)
```

```
In [ ]: ct7 = pd.crosstab(sub1['anyarrest'], sub1['compv6'])
        print(ct7)
In [ ]: | colsum7 = ct7.sum(axis=0)
        colpct7 = ct7/colsum7
        print(colpct7)
In [ ]: print('chi square value, p value, expected values')
        cs7 = scipy.stats.chi2 contingency(ct7)
        print(cs7)
In [ ]: recode7 = {40:40, 50:50}
        sub1['compv7'] = sub1['age'].map(recode7)
In [ ]: ct8 = pd.crosstab(sub1['anyarrest'], sub1['compv7'])
        print(ct8)
In [ ]: colsum8 = ct8.sum(axis=0)
        colpct8 = ct8 / colsum8
        print(colpct8)
In [ ]: print('chi square value, p value, expected values')
        cs8 = scipy.stats.chi2 contingency(ct8)
        print(cs8)
```

In [ ]: #now we will test whether there is a relationship between two quantitative variables, age\_unprocessed an d allcrimes

#for this we use the pearson correlation test

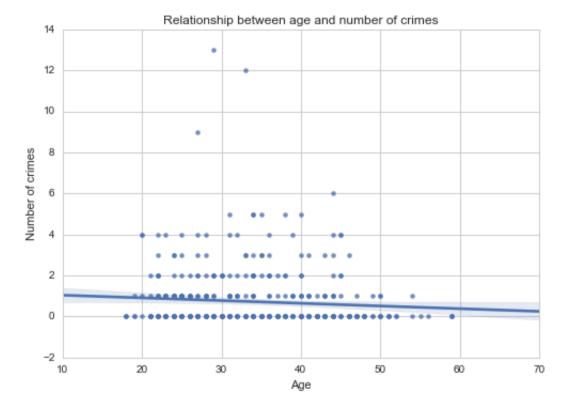
#r, going from -1 to 1 only tells us whether the two variables are linearly related. they may be related in nonlinear ways

#therefore it's always important to look at r in parallel with a scatterplot of the two variables
#r squared is a measure of how much variability in one variable can be explained by the other variable
#to calculate the pearson coefficient we need to remove all missing values

In [ ]: rdc.columns

```
In [154]: scat1 = sns.regplot(x='age_unprocessed', y='allcrimes', fit_reg=True, data=rdc)
    plt.xlabel('Age')
    plt.ylabel('Number of crimes')
    plt.title('Relationship between age and number of crimes')
    scat1
```

Out[154]: <matplotlib.axes.\_subplots.AxesSubplot at 0x11dc61860>



In [ ]: data\_pearson\_test = rdc[['age\_unprocessed', 'allcrimes']].dropna()

```
In [ ]: print('association between age and number of crimes')
        print(scipy.stats.pearsonr(data pearson test['age unprocessed'],
                                                    data pearson test['allcrimes']))
In [ ]: #a moderator is a third variable that affects the direction and/or strength between your explanatory and
        response variables
        #the question is, is our response variable associated with our explanatory variable, for each level of o
        ur third variable?
In [ ]: #let's see if gender is a moderator variable for test group -> allcrimes
In [ ]: sub11 = rdc[['cond', 'sex', 'allcrimes']].dropna()
In [ ]: rdc['sex'].value counts()
In [ ]: sub12 = sub11[sub11['sex']==1]
        sub13 = sub11[sub11['sex']==2]
In [ ]: model 12 = smf.ols(formula='allcrimes ~ C(cond)', data=sub12).fit()
        print(model 12.summary())
        model 13 = smf.ols(formula='allcrimes ~ C(cond)', data=sub13).fit()
        print(model 13.summary())
In [ ]: sub12.groupby('cond').mean()
In [ ]: sub12.groupby('cond').std()
In [ ]: sub13.groupby('cond').mean()
```

- In [ ]: sub13.groupby('cond').std()
  In [ ]: sns.factorplot(x='cond', y='allcrimes', kind='bar', data=sub12)
  In [ ]: sns.factorplot(x='cond', y='allcrimes', kind='bar', data=sub13)
- In []: #we would test for moderator variables with the chi square test the same way
  #divide the population into the sublevels of the third variables
  #conduct a chi square test for each to see if the relationship is statistically significant for each lev
  el
  #we would visualize it with a linegraph factorplot(kind='point')
- In [ ]: #to know if your data is obersevational or experimental, you ask if the explanatory variable was manipul ated or not #data reporting tells you what is happening, but data analysis tells you why it is happening

In [ ]: #randomization works best as your sample size approaches infinity. for small sizes, imbalances in the groups

#can occur. if you check randomized studies, one of first steps is to check for imbalances between group s

#on covariates. this is also why we can conclude that variables are associated, but hardly that one caus es the other

#statistical control: include unbalanced covariates as additional explanatory variables in the study.

#In a true experiment, 3 conditions:

#1. only one variable is manipulated

#2. we have a control group

#3. random assignment

#In theory, in this case one can determine causality

**#Quasi** experiment:

#1. only one variable is manipulated

#2. control group

#3. no random assignment; groups pre selected. i.e. drug users study.

#To improve a quasi experimental design: add confounding variables; have a control group; use a pre-test /post-test design

#confounder=control variable=covariate=third variable=lurking variable

In [ ]: #identifying a confounding variable does not allow to establish causation, just to get closer to a causa 1 connection.

#due to infinite number of possible lurking variables, observational studies cannot rly establish causat ion

#a lurking of confounding variable is a third variable that is associated with both the explanatory and response

#variables.

#i.e. x=firefighters; y=damage caused by a fire. plot would suggest more firefighters cuases more fire d amage.

#in reality there is a third lurking variable that influences both, seriousness of the fire.

#In a study we want to demonstrate that our statistical relationship is valid even after controlling for confounders.

# In [ ]: #Linear regression:

#multivariate linear regression for quantitative response variable

#logistic regression for binary categorical response variable

#Assumptions:

#Normality: residuals from our linear regression model are normally distributed. if they are not, #our model may be misspecified.

#Linearity: association between explanatory and response variable is linear

#Homoscedasticity (or assumption of constant variance): variability in the response variable is the same at all levels

#of the explanatory variable. i.e. if spread of residuals values increases as you move along x axis, ass umption is

#false.

#Independence: observations are not correlated with each other. Longitudinal data can violate this assum ption, as well

#as hierarchical nesting/clustering data i.e. looking at students by classes. this assumption is the most t serious

#to be violated, and also cannot be fixed by modifying the variables. the data structure itself is the p roblem.

#We have to contend with:

#Multicollinearity: explanatory variables are highly correlated with each other. this can mess up your p arameter estimates

#or make them highly unstable. Signs: 1. highly associated variable not significant. 2. negative regress ion coefficient

#that should be positive. 3. taking out an explanatory variable drastically changes the results. #Outliers: can affect your regression line, meaning it will not fit the data as well as it should.

```
In [ ]: #multiple regression model allows us to find the relationship between one explanatory variable and the
        #reponse variable, while controlling (holding constant at 0) all the other variables.
        #categorical sex (1 and 2); age restricted to 18-25 group: each variable needs to include a meaningful
        #value of 0, so as to make it easier to interpret the coefficients
        #for a categorical variable, we can just recode one of the values to be 0
        #for a quantitative variable, we have to center it. Centering = subtracting the mean of a variable
        #from the value of the variable. We are therefore recoding it so that its mean=0.
        #Note: do not center the response variable.
In [ ]: #We will create a multiple regression model, investigating the relationship between the study group 'con
        d', 'aqe', 'sex',
        #and the quantitative response variable 'allcrimes'. We will then do the same for 'allarrests'.
```

#we will first center the explanatory variables. for categorical variables, one of the categories needs to be 0, for #quantitative variables, we need to subtract the mean from each value.

```
In [81]: rdc linear = rdc[['cond', 'age unprocessed', 'sex', 'allcrimes', 'allarrests']].dropna()
         len(rdc linear)
```

Out[81]: 439

```
In [83]: rdc linear['age unprocessed c'] = rdc['age unprocessed']-rdc['age unprocessed'].mean()
         print(rdc linear['age unprocessed c'].mean())
```

-0.24115029362859228

```
In [84]: rdc linear['sex'].value counts()
```

Out[84]: 1 362 77

Name: sex, dtype: int64

```
In [85]: recode20 = {1:1, 2:0}
         rdc linear['sex c'] = rdc linear['sex'].map(recode20)
```

In [92]: model20 = smf.ols(formula = 'allcrimes ~ C(cond)', data=rdc linear).fit() print(model20 summary())

print(mode120.	summary())						
		OLS Regres	sion Resul	.ts			
Dep. Variable:		allcrimes	R-square	ed:	0.004		
Model:		OLS	Adj. R-s	quared:	0	.002	
Method: Le		east Squares	F-statis	stic:	1	.890	
Date: Thu,		09 Mar 2017	Prob (F-	·statistic):	0	.170	
Time:		07:20:13	Log-Like	elihood:	-788.43		
No. Observation	439	AIC:		1581.			
Df Residuals:		437	BIC:		1589.		
Df Model:		1					
Covariance Type	e: 	nonrobust					
	coef	std err	t	P> t	[95.0% Conf.	Int.]	
Intercept	0.8241	0.099	8.288	0.000	0.629	1.019	
C(cond)[T.1]	-0.1918	0.140	-1.375	0.170	-0.466	0.082	
Omnibus:		380.446	 Durbin-W			.931	
<pre>Prob(Omnibus):</pre>		0.000	Jarque-E	Bera (JB):	8874.157		
Skow.		3 688	Prob(JB)	•		0 00	

===============					
Kurtosis:	23.754	Cond. No.		2	2.64
Skew:	3.688	Prob(JB):		(	0.00
Prob(Omnibus):	0.000	Jarque-Bera	(JB):	8874.	.157
Omnibus:	380.446	Durbin-Watso	on:	1.	.931
C(cond)[T.1] -0.	.1918	-1.3/5 0 =======	.170 :======	-0.466 =======	0.082 ====

# Warnings:

In [93]: model21 = smf.ols(formula = 'allcrimes ~ C(cond)+age\_unprocessed\_c', data=rdc\_linear).fit()
 print(model21.summary())

	0	LS Regress	ion Results			
=======================================		=======		========	========	
Dep. Variable:	a	llcrimes	R-squared:		0.010	
Model:		OLS	Adj. R-squar	ed:	0.006	
Method:	Least	Least Squares F			2.260	
Date:	Thu, 09 Mar 2017 P		Prob (F-stat	istic):	0.106	
Time:		07:20:16	Log-Likeliho	od:	-787.12	
No. Observations:		439	AIC:		1580.	
Df Residuals:		436	BIC:		1592.	
Df Model:		2				
Covariance Type:	n	onrobust				
=============				========		=====
	coef		t		[95.0% Conf.	<pre>Int.]</pre>
Intercept	0.8186				0.623	1.014
C(cond)[T.1]	-0.1871	0.139	-1.344	0.180	-0.461	0.087
age_unprocessed_c	-0.0129	0.008	-1.619		-0.029	0.003
Omnibus:		380.213	======== Durbin-Watso		1.935	
Prob(Omnibus):		0.000	Jarque-Bera	(JB):	8859.859	
Skew:			Prob(JB):	` ,	0.00	
Kurtosis:			Cond. No.		20.0	
=======================================	.=======					

In [94]: model22 = smf.ols(formula = 'allcrimes ~ C(cond)+age\_unprocessed\_c+C(sex)', data=rdc\_linear).fit()
 print(model22.summary())

OLS Regression Results							
Dep. Variable:	a	llcrimes	R-squared:		0.023		
Model:		OLS	Adj. R-squar	ed:	0.017		
Method:	Least	Least Squares F			3.471		
Date:	Thu, 09	Mar 2017	Prob (F-stat	istic):	0.0162		
Time:		07:20:38	Log-Likeliho	od:	-784.19		
No. Observations:		439	AIC:		1576.		
Df Residuals:		435	BIC:		1593.		
Df Model:		3					
Covariance Type:	n	onrobust					
=========	coef	std err	========= t	P> t	========= [95.0% Conf	. Int.]	
Intercept	0.8988	0.104	8.627	0.000	0.694	1.104	
C(cond)[T.1]	-0.1921	0.139	-1.387	0.166	-0.464	0.080	
C(sex)[T.2]	-0.4412	0.182	-2.417	0.016	-0.800	-0.082	
age_unprocessed_c	-0.0116	0.008	-1.459	0.145	-0.027	0.004	
Omnibus:	=======	377.881	======== Durbin-Watso	======== on:	1.975		
Prob(Omnibus):		0.000	Jarque-Bera	(JB):	8785.690		
Skew:		3.650	Prob(JB):		0.00		
Kurtosis:	=======	23.665	Cond. No.		23.7		

In [95]: model23 = smf.ols(formula = 'allcrimes ~ C(cond)+age\_unprocessed\_c + I(age\_unprocessed\_c\*\*2)+C(sex)', da
 ta=rdc\_linear).fit()
 print(model23.summary())

OLS Regression Results									
Dep. Variable: Model: Method: Date: Time: No. Observations: Df Residuals: Df Model: Covariance Type:	allcrimes OLS Least Squares Thu, 09 Mar 2017 07:21:30 439 434 nonrobust	Adj. R-sq F-statist Prob (F-s Log-Likel	quared: ic: :tatistic):	0	0.033 0.024 3.673 0.00590 782.07 1574. 1595.				
=======================================	coef	std err	t	P> t	[95.0% Cc	onf. Int.]			
C(sex)[T.2]	1.0387 -0.2112 -0.4521 -0.0072 ** 2) -0.0017	0.138 0.182 0.008	-1.527 -2.485 -0.876	0.128 0.013 0.382	-0.810 -0.023	0.061 -0.095 0.009			
Omnibus: Prob(Omnibus): Skew: Kurtosis:	374.130 0.000 3.602 23.351	Prob(JB):	era (JB):	85	1.982 525.279 0.00 320.				

```
In [97]: model24 = smf.ols(formula = 'allarrests ~ C(cond)', data=rdc_linear).fit()
    print(model24.summary())
```

		OLS Regres	sion Resul	ts			
Dep. Variable: allarrests			 R-squared:		0.000		
Model:	OLS		Adj. R-s	Adj. R-squared:		-0.002	
Method:	Least Squares		F-statistic:		0.2025		
Date:	Sat,	Sat, 11 Mar 2017		<pre>Prob (F-statistic):</pre>		0.653	
Time:		17:20:34	•		-591.96		
No. Observations:		439	AIC:		1188.		
Df Residuals:	s: 437		BIC:		1196.		
Df Model:		1					
Covariance Type	<b>:</b>	nonrobust					
=========	coef				[95.0% Conf	. Int.]	
Intercept	0.4213		6.629		0.296	0.546	
C(cond)[T.1]	-0.0401	0.089	-0.450	0.653	-0.215	0.135	
Omnibus:	=======	406.395	====== Durbin-W	======================================	=======	===== 1.862	
<pre>Prob(Omnibus):</pre>		0.000			10979.833		
Skew:		4.027	• ,		0.00		
Kurtosis:		26.138	•		2.64		
=======================================			=======			====	

In [99]: model25 = smf.ols(formula = 'allarrests ~ C(cond)+age\_unprocessed\_c', data=rdc\_linear).fit()
 print(model25.summary())

	0	LS Regress	ion Results			
Dep. Variable:	allarrests		R-squared:		0.016	
Model:			Adj. R-square	ed:	0.011	
Method:	Least Squares		F-statistic:		3.519	
Date:	Sat, 11 Mar 2017		Prob (F-stati	stic):	0.0305	
Time:	17:21:10		Log-Likelihoo	od:	-588.55	
No. Observations:		439	AIC:		1183.	
Df Residuals:		436	BIC:		1195.	
Df Model:		2				
Covariance Type:	n	onrobust				
	coef	std err	======== t 		======================================	 . Int.]
Intercept	0.4157	0.063			0.292	0.540
C(cond)[T.1]	-0.0354	0.089	-0.399	0.690	-0.209	0.139
	0.0001	0.003	0.000	0.000		
age_unprocessed_c	-0.0133	0.005		0.009	-0.023	-0.003
, ,	-0.0133	0.005		0.009		
age_unprocessed_c	-0.0133	0.005	-2.614 ========	0.009	-0.023	
<pre>age_unprocessed_c ====================================</pre>	-0.0133	0.005 ==================================	-2.614 ======= Durbin-Watson	0.009	-0.023 	
<pre>age_unprocessed_c ====================================</pre>	-0.0133	0.005 ==================================	-2.614 	0.009	-0.023  1.877 10991.230	

# Warnings:

In [103]: model26 = smf.ols(formula = 'allarrests ~ C(cond)+age\_unprocessed\_c+C(sex)', data=rdc\_linear).fit()
 print(model26.summary())

	0	LS Regress:	ion Results			
Dep. Variable:	allarrests		R-squared:		0.021	
Model:	OLS		Adj. R-squar	ed:	0.015	
Method:	Least Squares		F-statistic:		3.179	
Date:	Sat, 11 Mar 2017		Prob (F-stat	istic):	0.0239	
Time:	17:22:42		Log-Likelihood:		-587.30	
No. Observations:		439	AIC:		1183.	
Df Residuals:		435	BIC:		1199.	
Df Model:		3				
Covariance Type:		onrobust				
=======================================		std err			[95.0% Conf	. Int.]
Intercept	0.4490	0.067	6.749		0.318	0.580
C(cond)[T.1]	-0.0374	0.088	-0.423	0.673	-0.211	0.136
C(sex)[T.2]	-0.1834	0.117	-1.574	0.116	-0.412	0.046
age_unprocessed_c	-0.0127	0.005	-2.504			-0.003
Omnibus:		402.755	======== Durbin-Watso		1.884	
Prob(Omnibus):		0.000	Jarque-Bera (JB):		10846.200	
Skew:		3.970	Prob(JB):		0.00	
Kurtosis:		26.020	Cond. No.		23.7	

# Warnings:

•	2 ( ) ,					
	OLS Regres	sion Result	s			
Dep. Variable:	allarrests	R-squared	:		0.022	
Model:	OLS	Adj. R-sq	uared:	0.013 2.409 0.0487 -587.24 1184. 1205.		
Method:	Least Squares	F-statist	ic:			
Date:	Sat, 11 Mar 2017	Prob (F-s	tatistic):			
Time:	17:22:34	Log-Likel	ihood:			
No. Observations:	439	AIC:				
Df Residuals:	434	BIC:				
Df Model:	4					
Covariance Type:	nonrobust					
	coef		t 		[95.0% Con	if. Int.]
Intercept	0.4640				0.307	0.621
_	-0.0395	0.089	-0.445	0.657	-0.214	0.135
C(sex)[T.2]	-0.1846	0.117	-1.582	0.114	-0.414	0.045
age_unprocessed_c	-0.0123	0.005	-2.325	0.021	-0.023	-0.002
<pre>I(age_unprocessed_c</pre>	** 2) -0.0002	0.001	-0.342	0.733	-0.001	0.001
Omnibus:	402.301	======= Durbin-Wa	tson:	=======	1.880	
Prob(Omnibus):	0.000	Jarque-Be	ra (JB):	10785.618		
Skew:	Prob(JB): 0.00					
Kurtosis:	25.951	Cond. No.			320.	

# Warnings:

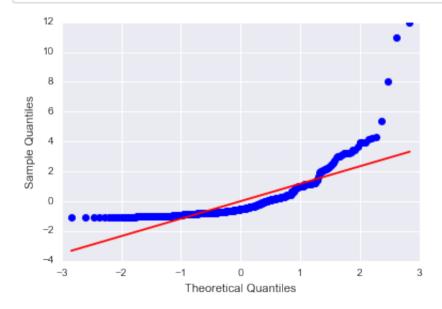
```
In [105]: #group means and sd
          print('Mean')
          ds1 = rdc linear.groupby('cond').mean()
          print(ds1)
          Mean
                age unprocessed allcrimes allarrests age unprocessed c
                                                                             sex c
          cond
          0
                      33.694444
                                  0.824074
                                              0.421296
                                                               -0.423699 0.819444
          1
                      34.053812
                                  0.632287
                                             0.381166
                                                               -0.064332 0.829596
In [106]: print('Standard Deviation')
          ds2 = rdc linear.groupby('cond').std()
          print(ds2)
          Standard Deviation
                age unprocessed allcrimes allarrests age unprocessed c
                                                                             sex c
          cond
          0
                       9.026533 1.710861
                                             1.013082
                                                                9.026533 0.385543
          1
                       8.438798 1.169907
                                             0.850545
                                                                8.438798 0.376833
  In [ ]: #For each response variable, allcrimes and allarrests, we choose the model that gave us the highest over
          a11
          #explanatory power, and run further tests to check for evidence of model misspecification.
          #If model is correctly specified, residuals are not correlated with explanatory variables.
          #If data fails to meet regression assumptions, or misses explanatory variables, we have model misspecifi
```

#Intercept = value of response variable when all explanatory variables are held constant at 0

cation.

In [110]: #Q-Q plot for normality
 fig4 = sm.qqplot(model23.resid, line='r')
 #red line represents residuals we would expect if model residuals were normally distributed
 #our residuals below deviate significantly from red line, especially at lower and higher quantiles, mean
 ing they do not
 #follow a normal distribution. This means curvilinear association we saw is not fully explained by our m
 odel. We could add

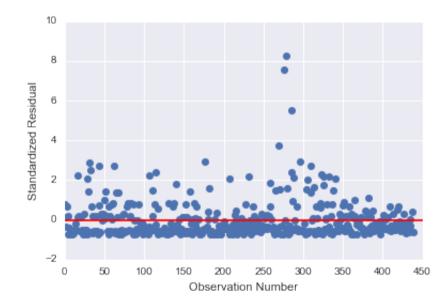
#more explanatory variables.



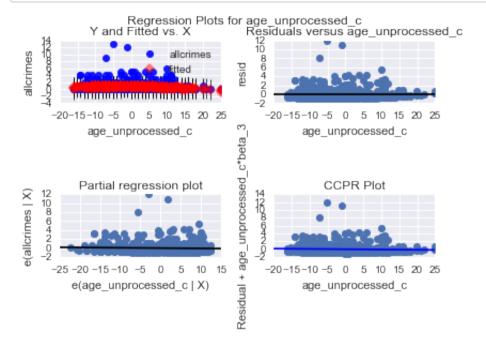
In [ ]: #normalizing or standardizing values makes them fit a standard normal distribution

# In [111]: #simple plot of residuals stdres = pd.DataFrame(model23.resid\_pearson) plt.plot(stdres, 'o', ls='None') l = plt.axhline(y=0, color='r') plt.ylabel('Standardized Residual') plt.xlabel('Observation Number') #resid\_pearson normalizes our model's residuals #ls='none' means points will not be connected #we expect most residuals to fall within 2sd of the mean. More than 2 are outliers, and more than 3 extr eme outliers. #if more than 1% of our observations have standardized residuals with an absolute value greater than 2.5 , or more than 5% #have one greater than or equal to 2, there is evidence that the fit of the model is poor. largest cause of this is ommission #of important explanatory variables in our model.

Out[111]: <matplotlib.text.Text at 0x11d0630b8>

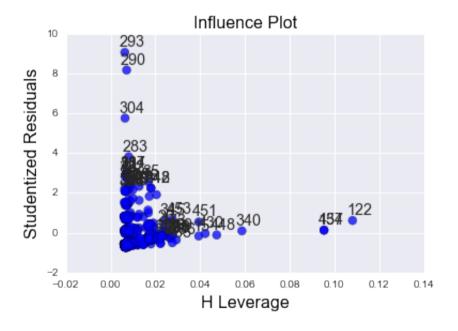


In [113]: #additional regression diagnostic plots
#fig1 = plt.figure(figsize(12,8))
fig1 = sm.graphics.plot\_regress\_exog(model23, 'age\_unprocessed\_c')



In [114]: #leverage plot
 fig3 = sm.graphics.influence\_plot(model23, size=8)
 print(fig3)
 #we see that we have extreme outliers, but they are low leverage, meaning they do not have an undue infl
 uence on our
 #estimation of the regression model.
 #we also have high leverage observations, but they are not outliers.
#we have no observations that are both high leverage and outliers.

# Figure(480x320)



In [ ]: #now we will apply logistic regression models to the binary response variables anycrime and anyarrest #the data management has already been performed

In [158]: rdc\_logistic['anycrime'] = pd.to\_numeric(rdc\_logistic['anycrime'], errors='coerce')
 rdc\_logistic['anyarrest'] = pd.to\_numeric(rdc\_logistic['anyarrest'], errors='coerce')

-0.557

0.303

Optimization terminated successfully.

-0.1268

Current function value: 0.563817

Iterations 5

Logit Regression Results

No. Observations: 441 Dep. Variable: anyarrest Logit Df Residuals: Model: 439 Method: MLE Df Model: Sun, 12 Mar 2017 Pseudo R-squ.: 0.0006709 Date: Time: 15:21:29 Log-Likelihood: -248.64 True LL-Null: converged: -248.81 LLR p-value: 0.5634 \_\_\_\_\_\_ coef std err [95.0% Conf. Int.] Intercept -1.02590.154 -6.645 0.000 -1.328-0.723

-0.578

0.563

0.220

C(cond)[T.1]

```
In [164]: #however, for logistic regression it makes much more sense to calculate the odds ratio
          #if OR=1, the model is not statistically significant
          #if OR<1, the response variable becomes less likely as the explanatory one increases
          #if OR>1, the response variable becomes more likely as the explanatory one increases
          print('Odds Ratios')
          print(np.exp(lreg1.params))
          #Study subjects in the treatment group (cond=1) are 0.88 times as likely to have had an arrest since rel
          ease on parole
          #as study subjects in the control group
          Odds Ratios
          Intercept
                          0.358491
          C(cond)[T.1]
                          0.880886
          dtype: float64
In [163]: # odd ratios with 95% confidence intervals
```

```
params = lreg1.params
conf = lreg1.conf_int()
conf['OR'] = params
conf.columns = ['Lower CI', 'Upper CI', 'OR']
print (np.exp(conf))
#we have 95% confidence that the population odds ratio will be between 0.57 and 1.35.
```

```
Lower CI Upper CI OR
Intercept 0.264892 0.485161 0.358491
C(cond)[T.1] 0.572854 1.354554 0.880886
```

# In [ ]: #Machine learning encompasses a wide range of statistical methods. These can be used for: #1. Describe associations #2. Search for patterns #3. Make predictions #We typically do not use ML with hypotheses in mind. instead we learn from the data #We learn from the test set. #Accuracy = test error rate. The rate at which it correctly classifies or estimates. #Goal is to minimize test error rate. #Linear regression: accuracy = mean squared error #Variance = change in parameter estimates across different data sets #Bias = how far off model estimated values are from true values #ideally we want low variance and low bias, but they are negatively associated. As one decreases, the ot her increases. #Generally, complexity of model leads to high variance and low bias #Simple models will have lower variance, but also be more biased. #Logistic regression: accuracy = how well the model classifies observations

# In [ ]: #Supervised Prediction includes: #Linear regression #Pattern recognition **#Discriminant analysis** #Multivariate function estimation #Supervised ML techniques #Decision trees #Like linear regression, decision trees are designed for supervised prediction problems. #Root node, and terminal nodes or leaves. #Growing the tree process: binary splits maximize correct classification; all cut points are tested; sub groups showing #similar outcomes are generated #Validating the tree: cross validation guards against overfit. A random subset is tested and only 'branc hes' that improve #the classification are retained #Selected sub tree is the lowest probability of misclassification #Trees allow the handling of many variables that cannot be done as efficiently in linear regression. The v can also uncover #constellations of variables that can predict high or low rates of the response variable.

# In [ ]: #Strengths of decision trees

#Can select from a large number of variables those and their interactions that are most important in det ermining the

#target or response variable to be explained

#They are easy to interpret and visualize, especially when the tree is small

#Can handle large data sets well and can predict both binary, categorical target variables and also quan titative ones

#Limitations: small changes in the data can lead to different splits and this can undermine the interpre tability of the model

#and decision trees are not very reproducible on future data

In [201]: reduced\_dataset\_clean = reduced\_dataset.dropna()
 len(reduced\_dataset\_clean)

Out[201]: 433

In [205]: predictors = reduced\_dataset\_clean.ix[:, reduced\_dataset\_clean.columns != 'allcrimes']

In [206]: predictors

Out[206]:

	cond	sex	age	living_situation	support	allarrests	anyarrest	alldrugs	anydrugs	anycrime	arrest_9mo	reincarc_9mo	nun
0	1	1	30.0	5	11.0	2.0	1.0	2.0	1.0	1.0	1	1	2
1	1	1	26.0	5	11.0	0.0	0.0	1.0	1.0	1.0	1	1	1
2	1	1	46.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
3	0	1	26.0	4	9.0	0.0	0.0	1.0	1.0	1.0	0	0	0
4	0	1	37.0	4	6.0	0.0	0.0	0.0	0.0	0.0	0	0	0
5	1	1	41.0	4	9.0	0.0	0.0	1.0	1.0	1.0	0	0	0
6	1	1	45.0	5	11.0	0.0	0.0	0.0	0.0	0.0	0	0	0
7	0	1	30.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
8	0	1	29.0	5	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0
9	0	1	35.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
11	0	1	37.0	4	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0
12	0	1	24.0	4	9.0	0.0	0.0	1.0	1.0	0.0	0	0	0
13	1	1	24.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
14	1	1	32.0	4	9.0	0.0	0.0	1.0	1.0	0.0	0	0	0

15	0	1	54.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
16	1	1	24.0	4	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0
17	1	1	25.0	5	9.0	4.0	1.0	1.0	1.0	1.0	1	1	1
18	1	1	29.0	4	9.0	0.0	0.0	0.0	0.0	1.0	1	1	1
19	1	1	41.0	4	1.0	0.0	0.0	1.0	1.0	1.0	0	0	0
20	1	1	30.0	5	5.0	0.0	0.0	0.0	0.0	0.0	0	0	0
21	0	1	21.0	5	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
22	1	1	37.0	4	1.0	0.0	0.0	3.0	1.0	1.0	1	1	2
23	1	1	39.0	5	1.0	0.0	0.0	1.0	1.0	0.0	0	0	0
24	0	1	28.0	4	9.0	0.0	0.0	1.0	1.0	0.0	0	0	0
25	0	1	34.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
26	1	1	19.0	4	9.0	0.0	0.0	0.0	0.0	1.0	0	0	0
27	1	1	41.0	4	1.0	0.0	0.0	9.0	1.0	0.0	0	0	0
28	1	1	24.0	5	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0
30	0	1	27.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
31	0	1	27.0	4	9.0	1.0	1.0	2.0	1.0	1.0	1	1	1
438	1	1	25.0	4	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0
439	1	1	22.0	4	9.0	0.0	0.0	0.0	0.0	0.0	1	1	1
440	0	1	38.0	5	9.0	1.0	1.0	1.0	1.0	1.0	1	1	1
441	0	2	41.0	5	9.0	0.0	0.0	1.0	1.0	0.0	1	1	1

442	0	1	21.0	5	8.0	0.0	0.0	1.0	1.0	0.0	0	0	0
443	0	1	49.0	8	3.0	0.0	0.0	0.0	0.0	0.0	0	0	0
445	0	1	48.0	5	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0
446	1	1	48.0	2	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0
447	0	1	46.0	7	4.0	0.0	0.0	0.0	0.0	0.0	0	0	0
448	0	1	55.0	4	7.0	0.0	0.0	3.0	1.0	0.0	0	0	0
449	0	1	50.0	5	1.0	0.0	0.0	1.0	1.0	0.0	0	0	0
451	0	1	54.0	5	11.0	1.0	1.0	0.0	0.0	1.0	0	0	0
453	0	2	50.0	4	2.0	1.0	1.0	0.0	0.0	1.0	0	0	0
454	1	2	44.0	5	7.0	0.0	0.0	0.0	0.0	0.0	0	0	0
455	0	1	21.0	5	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
457	0	1	45.0	5	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0
458	0	2	43.0	5	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0
459	0	1	51.0	4	7.0	0.0	0.0	0.0	0.0	0.0	0	0	0
460	1	1	45.0	4	1.0	1.0	1.0	0.0	0.0	1.0	0	0	0
461	0	1	19.0	5	4.0	0.0	0.0	0.0	0.0	0.0	0	0	0
463	0	1	24.0	5	9.0	0.0	0.0	1.0	1.0	0.0	0	0	0
464	0	2	37.0	5	7.0	0.0	0.0	0.0	0.0	0.0	0	0	0
465	0	1	18.0	5	6.0	0.0	0.0	0.0	0.0	0.0	0	0	0
467	1	1	42.0	4	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
468	0	1	42.0	5	1.0	0.0	0.0	0.0	0.0	0.0	0	0	0

469	0	2	42.0	2	3.0	0.0	0.0	0.0	0.0	0.0	0	0	0
470	0	1	24.0	5	4.0	1.0	1.0	1.0	1.0	1.0	0	0	0
471	1	1	37.0	5	9.0	0.0	0.0	0.0	0.0	0.0	0	0	0
472	1	2	31.0	4	2.0	1.0	1.0	0.0	0.0	1.0	0	0	0
473	0	1	41.0	4	1.0	0.0	0.0	1.0	1.0	0.0	0	0	0

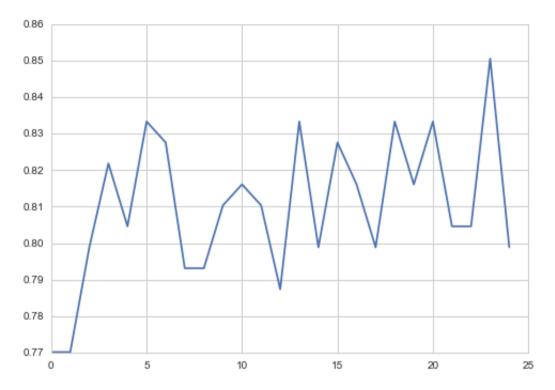
433 rows × 16 columns

```
In [215]: sklearn.metrics.confusion matrix(tar test, predictions)
Out[215]: array([[112,
                                                       01,
                       19,
                                             2,
                                                       01,
                                                       0],
                                                       0],
                        1, 0,
                                                      0],
                         0, 0, 0, 0, 0,
                   0,
                                                      01,
                                                       011)
In [216]: sklearn.metrics.accuracy score(tar test, predictions)
Out[216]: 0.78735632183908044
  In [ ]: #Random forests
          #Forests of trees
          #Splits on only ONE variable in each node. Variable with largest association with Target among
          #candidate variables. Only among variables randomly selected to be tested for that node.
          #First a subset of explanatory variables is selected at random
          #Next the node is split with the Best variable of the subset. After this node is split, a new list of su
          bset variables
          #is selected at random to split on the next node.
          #typical k fold values: 5 or 10
In [219]: classifier2 = RandomForestClassifier(n estimators=25)
          classifier2 = classifier.fit(pred train, tar train)
In [221]: predictions2 = classifier2.predict(pred test)
```

```
In [222]: sklearn.metrics.confusion matrix(tar test, predictions)
Out[222]: array([[112,
                                                  01,
                    0, 20,
                              7,
                                   2, 1,
                                             2,
                                                  01,
                        7,
                                             0, 1],
                                             0, 01,
                                 1, 2, 0, 0],
                   0, 1, 0, 0, 1, 0, 0],
                 , 0
                                                  011)
In [223]: sklearn.metrics.accuracy score(tar test, predictions)
Out[223]: 0.7931034482758621
In [224]: #fit an extra Trees model to the data
          model = ExtraTreesClassifier()
          model.fit(pred train, tar train)
          #display the relative importance of each attribute
          print(model.feature importances )
          \begin{bmatrix} 0.04472255 & 0.01493451 & 0.05962542 & 0.04225549 & 0.06900537 & 0.05969145 \end{bmatrix}
            0.03740676 0.04925347 0.04173209 0.46073854 0.00651952 0.02053235
            0.03376509 0.02278637 0.01254948 0.02448154]
In [227]: trees = range(25)
          accuracy = np.zeros(25)
In [228]: for idx in range(len(trees)):
              classifier = RandomForestClassifier(n estimators = idx + 1)
              classifier = classifier.fit(pred train, tar train)
              predictions = classifier.predict(pred test)
              accuracy[idx] = sklearn.metrics.accuracy score(tar test, predictions)
```

In [229]: plt.cla()
 plt.plot(trees, accuracy)

Out[229]: [<matplotlib.lines.Line2D at 0x11e9f2320>]



```
In [230]: #Lasso regression
#penalized regression method
#supervised learning method
#shrinkage and selection method
#shrinkage = constraints on parameters that shrinks coefficients to 0
#selection = identifies most imp. variables associated with response variable
#Can increase prediction accuracy and improve model interpretability vs standard OLS
#When lambda=0, it becomes OLS regression
#Bias increases and variance decreases as lambda increases
#in Lasso regression, penalty is not fair if variables are not on the same scale
#standardize all predictor variables to have means equal to 0 and sd = 1
#Lasso regression has several algorithms, among them LAR (least angle regression-)
#sklearn library refers to the penalty term as 'alpha'
```

```
In []: #Limitations of lasso regression
#1. Selection of variables is 100% statistically driven
#2. If predictors are correlated, lasso arbitrarily selects one
#3. Estimating p values is not straightforward
#4. Different selection methods or statistical softwares can provide different results
#5. No guarantee that selected model is not overfitted nor that it's the best model
#All regression models can produce meaningless models without human intervention
#Best approach is a combination of ML, human intervention, and independent application
```

In [233]: | predictors.columns

```
In [240]: predictors['cond'] = preprocessing.scale(predictors['cond'].astype('float64'))
          predictors['sex'] = preprocessing.scale(predictors['sex'].astype('float64'))
          predictors['age'] = preprocessing.scale(predictors['age'].astype('float64'))
          predictors['living situation'] = preprocessing.scale(predictors['living situation'].astype('float64'))
          predictors['support'] = preprocessing.scale(predictors['support'].astype('float64'))
          predictors['allarrests'] = preprocessing.scale(predictors['allarrests'].astype('float64'))
          predictors['anyarrest'] = preprocessing.scale(predictors['anyarrest'].astype('float64'))
          predictors['alldrugs'] = preprocessing.scale(predictors['alldrugs'].astype('float64'))
          predictors['anydrugs'] = preprocessing.scale(predictors['anydrugs'].astype('float64'))
          predictors['anycrime'] = preprocessing.scale(predictors['anycrime'].astype('float64'))
          predictors['arrest 9mo'] = preprocessing.scale(predictors['arrest 9mo'].astype('float64'))
          predictors['reincarc 9mo'] = preprocessing.scale(predictors['reincarc 9mo'].astype('float64'))
          predictors['num arrest'] = preprocessing.scale(predictors['num arrest'].astype('float64'))
          predictors['num reincarc'] = preprocessing.scale(predictors['num reincarc'].astype('float64'))
          predictors['violent charge'] = preprocessing.scale(predictors['violent charge'].astype('float64'))
          predictors['property charge'] = preprocessing.scale(predictors['property charge'].astype('float64'))
          /Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:1: SettingWithCopyWarning
          A value is trying to be set on a copy of a slice from a DataFrame.
          Try using .loc[row indexer,col indexer] = value instead
          See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
          q-view-versus-copy
            if name == ' main ':
          /Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:2: SettingWithCopyWarning
          A value is trying to be set on a copy of a slice from a DataFrame.
          Try using .loc[row indexer,col indexer] = value instead
          See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
          q-view-versus-copy
            from ipykernel import kernelapp as app
          /Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:3: SettingWithCopyWarning
```

```
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
  app.launch new instance()
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:4: SettingWithCopyWarning
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:5: SettingWithCopyWarning
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:6: SettingWithCopyWarning
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:7: SettingWithCopyWarning
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:8: SettingWithCopyWarning
```

```
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:9: SettingWithCopyWarning
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:10: SettingWithCopyWarnin
q:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:11: SettingWithCopyWarnin
q:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:12: SettingWithCopyWarnin
q:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:13: SettingWithCopyWarnin
```

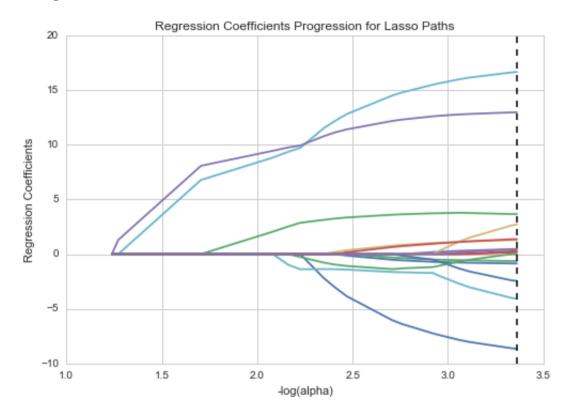
```
q:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:14: SettingWithCopyWarnin
q:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:15: SettingWithCopyWarnin
q:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
/Users/RI/Anaconda/anaconda/lib/python3.5/site-packages/ipykernel/ main .py:16: SettingWithCopyWarnin
q:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row indexer,col indexer] = value instead
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/indexing.html#indexin
q-view-versus-copy
```

# 

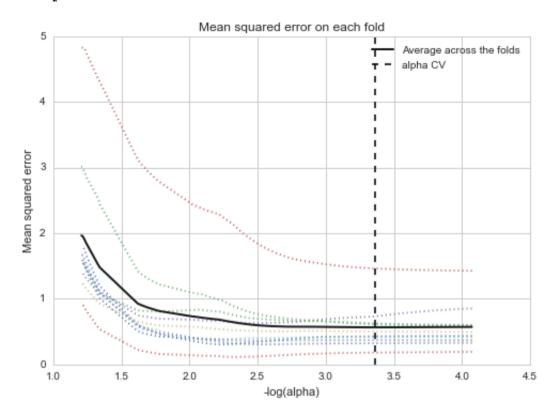
```
In [246]: # specify the lasso regression model
model=LassoLarsCV(cv=10, precompute=False).fit(pred_train,tar_train)
```

```
In [247]: # print variable names and regression coefficients
          dict(zip(predictors.columns, model.coef ))
Out[247]: {'age': 0.00922117951327419,
           'allarrests': 0.97816170401806635,
           'alldrugs': 0.23108994348020051,
            'anyarrest': -0.48346833180493959,
           'anycrime': 0.73524559191125771,
           'anydrugs': 0.01539580168267698,
           'arrest 9mo': 0.15463119466699699,
           'cond': -0.049336041215996017,
           'living situation': 0.027322802531668331,
           'num arrest': -0.13464215269435476,
           'num reincarc': 0.0,
           'property charge': 0.024425067866711093,
           'reincarc 9mo': -0.23122026644497329,
           'sex': -0.036216582632220544,
           'support': 0.074415953782179037,
           'violent charge': 0.072286686175727102}
In [248]: # plot coefficient progression
          m log alphas = -np.log10(model.alphas )
          ax = plt.qca()
          plt.plot(m log alphas, model.coef path .T)
          plt.axvline(-np.log10(model.alpha ), linestyle='--', color='k',
                      label='alpha CV')
          plt.ylabel('Regression Coefficients')
          plt.xlabel('-log(alpha)')
          plt.title('Regression Coefficients Progression for Lasso Paths')
```

Out[248]: <matplotlib.text.Text at 0x11ea7eac8>



Out[249]: <matplotlib.text.Text at 0x11f00aa90>



```
In [250]: # MSE from training and test data
          from sklearn.metrics import mean squared error
          train error = mean squared error(tar train, model.predict(pred train))
          test error = mean squared error(tar test, model.predict(pred test))
          print ('training data MSE')
          print(train error)
          print ('test data MSE')
          print(test error)
          training data MSE
          0.460347733345
          test data MSE
          0.54794067835
In [251]: # R-square from training and test data
          rsquared train=model.score(pred train,tar train)
          rsquared test=model.score(pred test,tar test)
          print ('training data R-square')
          print(rsquared train)
          print ('test data R-square')
          print(rsquared test)
          training data R-square
```

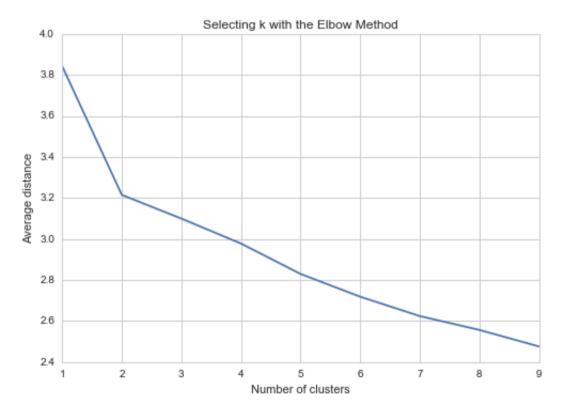
training data R-square 0.766331632034 test data R-square 0.789061561182

```
In [ ]: #Cluster analysis
          #Unsupervised learning method = no response variable included in the analysis
          #Goal: to have less variance within clusters, and more between clusters
          #Can also be used as a method of data reduction, to reduce number of variables
          #to the number of categorical variables equal to the clusters produced
          #Canonical discriminant analysis:
          #creates a smaller number of variables
          #linear combinations of clustering variables
          #canonical variables are ordered by proportion of variance accounted for
          #majority of variance is accounted for by first few canonical variables
In [255]: clustervar = predictors.copy()
In [257]: # split data into train and test sets
          clus train, clus test = train test split(clustervar, test size=.3, random state=123)
In [258]: # k-means cluster analysis for 1-9 clusters
          from scipy.spatial.distance import cdist
          clusters=range(1,10)
          meandist=[]
          for k in clusters:
              model=KMeans(n clusters=k)
              model.fit(clus train)
              clusassign=model.predict(clus train)
              meandist.append(sum(np.min(cdist(clus train, model.cluster centers , 'euclidean'), axis=1))
              / clus train.shape[0])
```

```
In [259]:
    """
    Plot average distance from observations from the cluster centroid
    to use the Elbow Method to identify number of clusters to choose
    """

    plt.plot(clusters, meandist)
    plt.xlabel('Number of clusters')
    plt.ylabel('Average distance')
    plt.title('Selecting k with the Elbow Method')
```

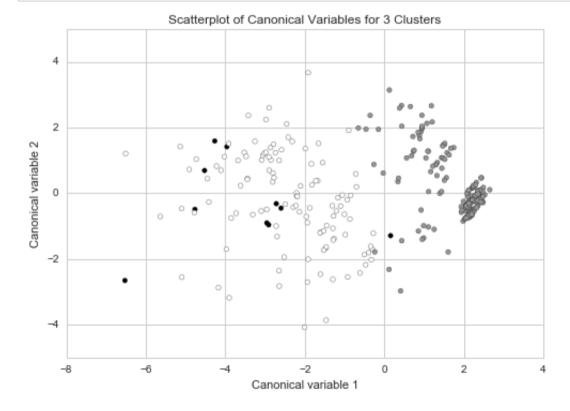
Out[259]: <matplotlib.text.Text at 0x11f661b70>



```
In [260]: # Interpret 3 cluster solution
    model3=KMeans(n_clusters=3)
    model3.fit(clus_train)
    clusassign=model3.predict(clus_train)
```

```
In [261]: # plot clusters

from sklearn.decomposition import PCA
pca_2 = PCA(2)
plot_columns = pca_2.fit_transform(clus_train)
plt.scatter(x=plot_columns[:,0], y=plot_columns[:,1], c=model3.labels_,)
plt.xlabel('Canonical variable 1')
plt.ylabel('Canonical variable 2')
plt.title('Scatterplot of Canonical Variables for 3 Clusters')
plt.show()
```



```
11 11 11
In [288]:
          BEGIN multiple steps to merge cluster assignment with clustering variables to examine
          cluster variable means by cluster
          # create a unique identifier variable from the index for the
          # cluster training data to merge with the cluster assignment variable
          clus train.reset index(level=0, drop=True)
          # create a list that has the new index variable
          cluslist=list(clus train['index'])
          # create a list of cluster assignments
          labels=list(model3.labels )
          # combine index variable list with cluster assignment list into a dictionary
          newlist=dict(zip(cluslist, labels))
          newlist
          # convert newlist dictionary to a dataframe
          newclus=pd.DataFrame.from dict(newlist, orient= 'index')
          newclus
          # rename the cluster assignment column
          newclus.columns = ['cluster']
          # now do the same for the cluster assignment variable
          # create a unique identifier variable from the index for the
          # cluster assignment dataframe
          # to merge with cluster training data
          newclus.reset index(level=0, drop=True)
          # merge the cluster assignment dataframe with the cluster training variable dataframe
          # by the index variable
          merged train=pd.merge(clus train, newclus, left index=True, right index=True)
          merged train.head(n=100)
          # cluster frequencies
          merged train.cluster.value counts()
```

```
Out[288]: 1 110
0 80
2 6
Name: cluster, dtype: int64

In [289]: """

END multiple steps to merge cluster assignment with clustering variables to examine cluster variable means by cluster
"""

# FINALLY calculate clustering variable means by cluster
clustergrp = merged_train.groupby('cluster').mean()
print ("Clustering variable means by cluster")
print(clustergrp)
```

Clustering variable means by cluster level 0 index cond age \ sex cluster 0 155.375000 218.900000 -0.125445 -0.132803 -0.115947 145.590909 224.527273 -0.061788 0.064352 -0.084813 1 2 61.666667 210.333333 -0.025412 -0.023272 -0.197488 living situation support allarrests anyarrest alldrugs \ cluster 0 0.126453 0.185858 0.048168 -0.085034 0.187033 -0.063659 -0.101742 0.012498 0.058328 -0.1218681 -0.078019 2 0.118736 -0.369018 0.176315 0.114828 anydrugs anycrime arrest 9mo reincarc 9mo num arrest \ cluster 0.264865 0.053554 -0.055090 -0.004977 -0.102726 0 -0.123996 0.094418 0.091929 0.035152 0.101847 1 2 -0.221593 -0.008142 0.280647 0.147061 0.036514 num reincarc violent charge property charge cluster -0.043510 0.059618 0.010329 0 0.012591 0.108077 0.091644 1 2 0.065391 -0.168830 -0.252749

```
In [307]: # validate clusters in training data by examining cluster differences in GPA using ANOVA
          # first have to merge GPA with clustering variables and cluster assignment data
          allarrests data['targets'] = pd.DataFrame(targets)
          # split GPA data into train and test sets
          arrests train, arrests test = train test split(allarrests data, test size=.3, random state=123)
          arrests train1=pd.DataFrame(arrests train)
          arrests train1.reset index(level=0, inplace=True)
          merged train all=pd.merge(arrests train1, merged train, left index=True, right index=True)
          sub1 = merged train all[['targets', 'cluster']].dropna()
          import statsmodels.formula.api as smf
          import statsmodels.stats.multicomp as multi
          allarrests mod = smf.ols(formula='targets ~ C(cluster)', data=sub1).fit()
          print (allarrests mod.summary())
          print ('means for all arrests by cluster')
          m1= sub1.groupby('cluster').mean()
          print (m1)
          print ('standard deviations for all arrests by cluster')
          m2= sub1.groupby('cluster').std()
          print (m2)
          mc1 = multi.MultiComparison(sub1['targets'], sub1['cluster'])
          res1 = mc1.tukeyhsd()
          print(res1.summary())
```

# OLS Regression Results

Dep. Variable:	targets	R-squared:	0.004
Model:	OLS	Adj. R-squared:	-0.007
Method:	Least Squares	F-statistic:	0.3610
Date:	Tue, 21 Mar 2017	Prob (F-statistic):	0.697
Time:	06:54:56	Log-Likelihood:	-358.59

No. Observations: 196 ATC: 723.2 Df Residuals: 193 BIC: 733.0 Df Model: Covariance Type: nonrobust coef std err P>|t| [95.0% Conf. Int.] Intercept 0.9000 0.170 5.298 0.000 C(cluster)[T.1] -0.1545 0.223 -0.692 0.490 0.565 1,235 -0.595 0.286 C(cluster)[T.2] -0.4000 0.643 -0.622 0.535 -1.668 0.868 Omnibus: 178.561 Durbin-Watson: 2.128 3045.652 Prob(Omnibus): 0.000 Jarque-Bera (JB): Skew: 0.00 3.497 Prob(JB): Kurtosis: 21.001 Cond. No. 7.01 \_\_\_\_\_\_

# Warnings:

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified. means for all arrests by cluster

targets

## cluster

- 0.900000
- 1 0.745455
- 2 0.500000

standard deviations for all arrests by cluster

targets

### cluster

- 0 1.879941
- 1 1.222393
- 2 0.836660

Multiple Comparison of Means - Tukey HSD, FWER=0.05

\_\_\_\_\_

group1 group2 meandiff lower upper reject

\_\_\_\_\_\_