lab08

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1 lab 08 - Distances and PCA

Name: Robb Alexander and Ryan Bailis Class: CSCI349 Semester: 2021SP Instructor: Brian King

```
[1]: # Setting things up
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import scipy.stats as stats
from scipy.stats import zscore
from sklearn.metrics import pairwise_distances
from sklearn.decomposition import PCA
```

1) Set up a pandas data frame with the following 8 observations and 3 variables:

```
2 test3 8 non-null int8 dtypes: category(2), int8(1) memory usage: 424.0+ bytes test1 test2 test3
```

```
[2]:
     ΑO
             Α
                 excellent
                                 25
     Α1
             C
                       fair
                                 32
     A2
             С
                                 60
                       good
     АЗ
             В
                                 53
                       fair
                                 23
     A4
             Α
                       poor
     A5
             В
                                 37
                 excellent
     A6
             C
                       good
                                 45
     A7
             В
                                 49
                       good
```

2) Show the output of df.describe(include='all'). What does the include='all' parameter do?

```
[3]:

"""

include='all' essentially shows everything

even if a column contains Nan values that

would normally be excluded from the basic

describe() function.

"""

df.describe(include='all')
```

```
[3]:
             test1 test2
                                 test3
     count
                  8
                         8
                              8.000000
     unique
                  3
                         4
                                   NaN
     top
                  В
                                   NaN
                     good
                  3
     freq
                         3
                                   NaN
                NaN
                            40.500000
     mean
                      {\tt NaN}
                NaN
                       {\tt NaN}
                            13.416408
     std
     min
                {\tt NaN}
                      {\tt NaN}
                            23.000000
     25%
               NaN
                      NaN 30.250000
     50%
               NaN
                      NaN 41.000000
     75%
                NaN
                      NaN 50.000000
                NaN
                      NaN
                            60.000000
     max
```

3) Show the output of df.test1.cat.categories, and df.test2.cat.categories. What is this showing? Does it work for df.test3.cat.categories? (If not, then comment this line out.)

```
[4]: """

The .cat variable is getting the categorical
Accessor of the variable and then it shows
the indexes with the .categories caller.
This is the categories labels
```

```
11 11 11
     df.test1.cat.categories
[4]: Index(['A', 'B', 'C'], dtype='object')
[5]: df.test2.cat.categories
[5]: Index(['poor', 'fair', 'good', 'excellent'], dtype='object')
[6]: # df.test3.cat.categories
    4) Show the output of df.test1.cat.codes, and df.test2.cat.codes. What is this showing?
[7]: """
     This is showing the numbers that each one
      of the observations have according to the
      category numbers. This is the converted
      strings into numerics.
      This is the categories' int mappings
     11 11 11
     df.test1.cat.codes
[7]: AO
           0
     A1
           2
     A2
           2
     ΑЗ
     A4
     Α5
           1
     A6
           2
     Α7
           1
     dtype: int8
[8]: df.test2.cat.codes
[8]: AO
    A1
           1
     A2
           2
     АЗ
           1
     Α4
           0
           3
     A5
           2
     A6
     Α7
     dtype: int8
```

5) Report the counts of each level of the categorical variables.

```
[9]: df.test1.cat.codes.value_counts()
 [9]: 1
           3
      2
           3
      0
           2
      dtype: int64
[10]: df.test2.cat.codes.value_counts()
[10]: 2
           3
      1
           2
      3
           2
      0
            1
      dtype: int64
```

6) Report a cross tabulation (i.e. contingency table) between test1 and test2. Include the margins (i.e. the sum of the rows and the columns) in your reported table (HINT: Look up pandas crosstab() function)

```
[11]: pd.crosstab(df.test1, df.test2, margins=True)
```

```
[11]: test2 poor fair good excellent
      test1
                        0
                                                 2
      Α
                 1
                                           1
                 0
                                                 3
      В
                        1
                               1
                                           1
      C
                 0
                        1
                               2
                                           0
                                                 3
      All
                 1
                        2
                               3
                                           2
                                                 8
```

7) From the previous table, store the contingency table without the margins in a variable called observed

```
[12]: observed = pd.crosstab(df.test1, df.test2)
```

8) Run a chi-squared test to determine whether test1 and test2 are dependent. Use the contingency table from the previous step. Clearly report the chi2 statistic, the p value, and the degrees of freedom, and then use the p-value to clearly state whether test1 and test2 are independent (assume p=0.05 threshold to test for independence)

```
[13]: c2, p, dof, expected = stats.chi2_contingency(observed)

print("Chi Squared Value:", c2)

print("Degrees of Freedom", dof)

print("p-value:", p, "\nTherefore we can conclude that 0.37 > 0.05,\nrejecting

→the null hypothesis that the two variables\nare independent; so they are

→thus, dependent")
```

Degrees of Freedom 6

p-value: 0.37528525266160834

```
Therefore we can conclude that 0.37 > 0.05, rejecting the null hypothesis that the two variables are independent; so they are thus, dependent
```

9) Create a new data frame called df_num, that represents a numeric version of the above. Do NOT do any rescaling of your variables yet! NOTE: If you do this from a dataframe that has the categorical variables set up properly, then this step is simple to do. The two choices I generally follow are either: 1) use the cat member of your categorical data, which stores a CategoricalAccessor object (look it up), or use one of the encoders in the sklearn.preprocessing module. The first option is easier, and yet another reason why it's so important to take the time to preprocess your data as correctly and error-free as possible.

Data columns (total 3 columns):

Column Non-Null Count Dtype
--- ----
0 test1 8 non-null int64
1 test2 8 non-null int64
2 test3 8 non-null int8
dtypes: int64(2), int8(1)
memory usage: 500.0+ bytes

[14]:		test1	test2	test3
	AO	0	3	25
	A1	2	1	32
	A2	2	2	60
	AЗ	1	1	53
	A4	0	0	23
	A5	1	3	37
	A6	2	2	45
	A7	1	2	49

10) As you learned in lecture, you must rescale your data to fall on a similar scale. There are different approaches to doing so. A standardized z-score is among the most common, but not necessarily always the best approach, especially when you are dealing with numeric representations of true categorical data. Rescaling your data to all fall

between 0 and 1 is also a common approach, particularly when you have categorical data.

```
[15]: scaler = pre.MinMaxScaler()

df_num_zeroone = df_num.copy()

df_num_zeroone[:] = scaler.fit_transform(df_num)

df_num_zeroone
```

```
[15]:
          test1
                    test2
                               test3
      ΑO
                 1.000000
                           0.054054
            0.0
      Α1
            1.0
                 0.333333
                           0.243243
      A2
                 0.666667
                            1.000000
            1.0
      АЗ
            0.5 0.333333
                           0.810811
      Α4
            0.0 0.000000
                           0.000000
      A5
            0.5
                 1.000000
                           0.378378
            1.0
                0.666667
      A6
                           0.594595
      A7
            0.5
                 0.666667
                           0.702703
```

11) Notice the value of test3. Quite often, when we have solid knowledge of what we expect our range to be, then we can rescale our data using that knowledge. In the case of test3, you learned that the data must fall between 0 and 100. Therefore, reassign test3 so that the min and max before rescaling are assuming to be between 0 and 100, respectively. (i.e. simply divide the original variable by 100)

```
[16]: df_num_zeroone.loc[:,'test3'] = df_num.test3 / 100
df_num_zeroone
```

```
[16]:
          test1
                     test2
                            test3
      ΑO
            0.0
                 1.000000
                             0.25
      Α1
            1.0
                 0.333333
                             0.32
                 0.666667
                             0.60
      A2
            1.0
      AЗ
            0.5
                 0.333333
                             0.53
      Α4
            0.0
                 0.000000
                             0.23
      A5
            0.5
                 1.000000
                             0.37
      A6
            1.0
                 0.666667
                             0.45
      Α7
            0.5
                             0.49
                 0.666667
```

12) Compute a single distance matrix called distmat_zeroone. Use a standard Euclidean distance measure. Your reported result should be an 8x8 matrix with appropriately labeled rows and columns. (HINT – study the output of the distance matrix functions! They do not output a square matrix. As discussed in lecture you can use either pairwise_distances from scikit-learn, or the pdist and squareform functions in scipy.spatial.distance). Be sure to label the rows and columns to be the index from the input dataframe. Round it to 3 significant digits.

```
[17]: names = ['A0','A1','A2','A3','A4','A5','A6','A7']
```

```
[17]:
           ΑO
                        A2
                              A3
                                    A4
                                          Α5
                                                A6
                                                       A7
                 A1
     AO 0.000 1.204 1.111 0.879 1.000 0.514 1.073
                                                   0.647
        1.204 0.000
                     0.435 0.542
     A1
                                 1.058 0.835 0.358
                                                    0.625
        1.111 0.435
                     0.000 0.605 1.258 0.643 0.150 0.512
     A2
     A3 0.879 0.542 0.605 0.000 0.672 0.686 0.606 0.336
     A4 1.000 1.058
                     1.258 0.672 0.000 1.127 1.222 0.873
     A5 0.514 0.835 0.643 0.686 1.127 0.000 0.606 0.354
     A6 1.073 0.358 0.150 0.606 1.222 0.606 0.000 0.502
     A7 0.647 0.625 0.512 0.336 0.873 0.354 0.502 0.000
```

13) Output the top three closest pairs of observations. You MUST write Python code to report these results! Do not simply print out your distance matrix and tell me your answers! Consider that this may have been thousands of observations! Always generate reported answers in code whenever you can! For each pair, output the pair of observations from the original dataframe, and the distance between them. (HINT: I found this easier to do with the output of pdist.)

```
[18]: [(1.204, ('A0', 'A1')), (1.111, ('A0', 'A2')), (0.879, ('A0', 'A3')), (1.0, ('A0', 'A4')), (0.514, ('A0', 'A5'))]
```

```
[19]: small = heapq.nsmallest(3, output)
for i, (dist, (row, col)) in enumerate(small):
    print(f'Closest: #{i}: [\'{row}\\', \'{col}\\'] dist={dist}\')
    print(pd.concat([df.loc[row], df.loc[col]], axis=1).T, "\n")
```

```
Closest: #0: ['A2', 'A6'] dist=0.15
  test1 test2 test3
```

```
A2
       C good
                  60
       C good
A6
                  45
Closest: #1: ['A3', 'A7'] dist=0.336
  test1 test2 test3
АЗ
       В
          fair
Α7
       В
         good
                  49
Closest: #2: ['A5', 'A7'] dist=0.354
              test2 test3
   test1
                       37
A5
       B excellent
Α7
       В
                       49
               good
```

A1

C

fair

32

14) Now, output the three most distant (least similar) pairs of observations. Again, for each pair, output the two observations, and the distance between them

```
[20]: largest = heapq.nlargest(3, output)
      for i, (dist, (row, col)) in enumerate(largest):
          print(f'Farthest: #{i}: [\'{row}\', \'{col}\'] dist={dist}')
          print(pd.concat([df.loc[row], df.loc[col]], axis=1).T, "\n")
     Farthest: #0: ['A2', 'A4'] dist=1.258
        test1 test2 test3
            C good
     A2
            A poor
                       23
     A4
     Farthest: #1: ['A4', 'A6'] dist=1.222
        test1 test2 test3
     Α4
            A poor
                       23
            С
              good
     A6
                       45
     Farthest: #2: ['A0', 'A1'] dist=1.204
        test1
                   test2 test3
               excellent
     ΑO
            Α
                            25
```

15) Create a new data frame, df_num_binarized, that stores the a binarized version for test1 and test2.

```
[21]: df_num_binarized = pd.get_dummies(data=df, prefix=['test1', 'test2'])

cols = list(df_num_binarized.columns)
df_num_binarized = df_num_binarized[cols[1:] + cols[0:1]]

df_num_binarized.loc[:,'test3'] = df_num_binarized.test3 / 100
df_num_binarized
```

```
[21]:
           test1_A test1_B test1_C test2_poor test2_fair test2_good \
      ΑO
                 1
                           0
                                      0
      Α1
                 0
                           0
                                                   0
                                                                 1
                                                                              0
                                      1
      A2
                 0
                           0
                                      1
                                                   0
                                                                 0
                                                                              1
      АЗ
                 0
                            1
                                      0
                                                   0
                                                                 1
                                                                              0
      Α4
                  1
                           0
                                      0
                                                   1
                                                                 0
                                                                              0
      Α5
                 0
                           1
                                      0
                                                   0
                                                                 0
                                                                              0
      A6
                 0
                            0
                                      1
                                                   0
                                                                 0
                                                                              1
                            1
                                      0
                                                   0
                                                                 0
      Α7
                                                                              1
           test2_excellent
                             test3
      ΑO
                               0.25
                          1
                          0
                               0.32
      Α1
                               0.60
                          0
      A2
                          0
                               0.53
      ΑЗ
      Α4
                               0.23
      Α5
                          1
                               0.37
      A6
                          0
                               0.45
      A7
                          0
                               0.49
```

16) Now, compute distmat_binarized by computing the distance matrix for the df_binarized.

```
[22]: names = ['A0','A1','A2','A3','A4','A5','A6','A7']
distmat_binarized = pd.DataFrame(pairwise_distances(X=df_num_binarized).

round(3), index=names, columns=names)
```

17) Report the three closest pairs, and the three most distant pairs from distmat_binarized

```
diag_df = distmat_binarized.mask(np.tril(np.ones(distmat_binarized.shape,u
→dtype=bool)))

pairs = list(itertools.product(names, names))

output = []

for i, v in enumerate(pairs):
    value = diag_df.loc[v[0], v[1]]
    if not np.isnan(value):
        output.append((value, v))
```

```
[24]: small = heapq.nsmallest(3, output)
for i, (dist, (row, col)) in enumerate(small):
    print(f'Closest: #{i}: [\'{row}\', \'{col}\'] dist={dist}')
    print(pd.concat([df.loc[row], df.loc[col]], axis=1).T, "\n")
```

```
A6
            C good
                        45
     Closest: #1: ['A0', 'A4'] dist=1.414
        test1
                    test2 test3
     A0
             Α
                excellent
                             25
                             23
     A4
             Α
                     poor
     Closest: #2: ['A3', 'A7'] dist=1.415
        test1 test2 test3
     АЗ
            В
               fair
                        53
     A7
            В
               good
                        49
[25]: largest = heapq.nlargest(3, output)
      for i, (dist, (row, col)) in enumerate(largest):
          print(f'Farthest: #{i}: [\'{row}\', \'{col}\'] dist={dist}')
          print(pd.concat([df.loc[row], df.loc[col]], axis=1).T, "\n")
     Farthest: #0: ['A2', 'A4'] dist=2.034
        test1 test2 test3
     A2
             С
                good
            Α
     A4
               poor
                        23
     Farthest: #1: ['A0', 'A2'] dist=2.03
        test1
                    test2 test3
             Α
                excellent
                             25
     AO
     A2
             C
                             60
                     good
     Farthest: #2: ['A3', 'A4'] dist=2.022
        test1 test2 test3
     АЗ
            В
               fair
                        53
                        23
     Α4
             Α
               poor
```

18) Take a moment and compare and contrast your results. Which method do you think have the better results? Why? Which variable do you think was the distinguishing player in affecting the different outcomes between both of the above approaches to transforming your data to numeric results? Why? Summarize what would have been the best transformation to make for all three variables that would have given the most accurate results. The general difference between the two is that it seems that the binarization one judges distance in the amount of exact matches instead of the caring about the order as much. By looking at the closest distance for the binarized one shows that it thinks poor and excellent are close enough to be the lowest distance. The binarized one thus cares more for the numerical chances with less focus on the So the important variable we had to take care of was test3. In the first one (zeroone), it cared more for the test2 and less so for the values of test3, while the opposite for the second one. So the best choice for test1 is binarization, test2 is zeroone, and test3 is factor of 100

19) Load in your next dataset using the following: df_car_crashes = sns.load_dataset('car_crashes')

```
[26]: df_car_crashes = sns.load_dataset('car_crashes')
```

20) Preprocess your data. Minimally, you should move the state code to become the index for the dataframe, and then drop that column from your dataframe. Show the first five rows.

```
[27]: df_car_crashes.index = df_car_crashes.abbrev
    df_car_crashes.drop('abbrev', axis=1, inplace=True)
    df_car_crashes.head(5)
```

```
[27]:
                      speeding alcohol not_distracted no_previous ins_premium \
      abbrev
      ΑL
               18.8
                         7.332
                                  5.640
                                                  18.048
                                                                15.040
                                                                              784.55
      ΑK
               18.1
                         7.421
                                  4.525
                                                  16.290
                                                                17.014
                                                                             1053.48
      AZ
               18.6
                         6.510
                                   5.208
                                                  15.624
                                                                17.856
                                                                              899.47
      AR.
               22.4
                         4.032
                                  5.824
                                                  21.056
                                                                21.280
                                                                              827.34
               12.0
                         4.200
                                                  10.920
                                                                              878.41
      CA
                                  3.360
                                                                10.680
```

```
ins_losses
abbrev
AL 145.08
AK 133.93
AZ 110.35
AR 142.39
CA 165.63
```

21) Create a new dataframe called df_car_crashes_zscore that represents the zscore transformation for df_car_crashes. Again, show the first five rows.

```
[28]: df_car_crashes_zscore = df_car_crashes.apply(zscore)
df_car_crashes_zscore.head(5)
```

```
[28]:
                                   alcohol not_distracted no_previous \
                 total speeding
      abbrev
      ΑL
              0.737446
                        1.168148 0.439938
                                                   1.002301
                                                                0.277692
      ΑK
              0.565936
                        1.212695 -0.211311
                                                  0.608532
                                                                0.807258
      ΑZ
              0.688443 0.756709 0.187615
                                                  0.459357
                                                                1.033141
      AR.
              1.619498 -0.483614 0.547408
                                                   1.676052
                                                                1.951700
      CA
             -0.928653 -0.399524 -0.891763
                                                  -0.594276
                                                               -0.891968
              ins_premium ins_losses
      abbrev
      ΑL
                -0.580083
                             0.430514
      ΑK
                 0.943258
                            -0.022900
      ΑZ
                 0.070876
                            -0.981778
      AR
                -0.337701
                             0.321125
```

22) Create a distance matrix called distmat_cars based on the df_car_crashes_zscore. Display the entire distance matrix.

[29]: distmat_cars = pd.DataFrame(pairwise_distances(X=df_car_crashes_zscore), u

→index=df_car_crashes_zscore.index, columns=df_car_crashes_zscore.index)

distmat_cars

[29]:	abbrev abbrev	AL	AK	AZ	AR	CA	CO	\
	AL	0.000000	1.848559	1.875942	2.616265	3.450372	2.687193e+00	
	AK	1.848559	0.00000	1.461454	2.961700	3.503615	2.674699e+00	
	AZ	1.875942	1.461454	0.000000	2.592934	3.873068	2.690949e+00	
	AR	2.616265	2.961700	2.592934	0.000000	4.773783	4.082043e+00	
	CA	3.450372	3.503615	3.873068	4.773783	0.000000	1.382473e+00	
	CO	2.687193	2.674699	2.690949	4.082043	1.382473	2.107342e-08	
	CT	4.022977	3.719860	4.341409	5.511677	1.360401	2.171261e+00	
	DE	2.426745	1.305465	2.322674	3.375632	2.801377	2.348937e+00	
	DC	6.627070	5.923340	6.312294	7.648001	3.725440	4.294426e+00	
	FL	2.863217	2.014017	2.471886	2.722795	3.294284	2.884757e+00	
	GA	2.740439	2.609142	2.619278	3.132498	1.994627	1.592712e+00	
	HI	1.967575	2.309612	2.090333	3.887437	4.367894	3.364956e+00	
	ID	3.227303	3.462721	2.330031	4.160356	4.023485	2.745180e+00	
	IL	2.648799	2.872699	2.779181	4.036791	1.416124	5.753356e-01	
	IN	2.936806	3.210155	2.440773	3.699517	2.832514	1.789320e+00	
	IA	3.114833	3.587012	2.842219	3.547224	3.017301	2.214125e+00	
	KS	1.858608	2.173195	1.774256	2.804821	2.488592	1.447705e+00	
	KY	1.951130	2.154966	1.855779	1.793132	3.376552	2.635520e+00	
	LA	3.864433	3.260930	4.243512	4.014378	4.931876	4.790391e+00	
	ME	2.803998	3.213492	2.324265	4.160980	3.400465	2.191438e+00	
	MD	4.050022	3.809894	4.515303	5.093197	1.647736	2.550081e+00	
	MA	5.453420	5.160132	5.288766	6.535539	2.506906	3.019168e+00	
	MI	3.471272	2.994360	3.509289	4.352050	1.698430	2.010389e+00	
	MN	4.735183	4.795405	4.670325	5.819223	2.042084	2.280417e+00	
	MS	4.470256	4.270578	4.110102	4.692240	3.437689	3.078930e+00	
	MO	1.086939	2.000188	2.032453	3.294652	2.593112	1.781583e+00	
	MT	3.494754	3.874574	2.985649	3.963502	6.286038	5.169746e+00	
	NE	3.307722	3.668479	2.927505	3.634205	2.980809	2.250106e+00	
	NV	2.248853	1.565833	2.021377	3.406048	2.217568	1.521520e+00	
	NH	3.706577	3.863496	3.575319	5.069694	2.025549	1.486772e+00	
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	NY	4.241067	3.472039	4.115832	5.286360	2.134562	2.525314e+00	
	NC	1.240410	2.276668	1.781326	3.167884	3.005854	1.983954e+00	
	ND	3.890314	4.709646	3.884830	3.080659	6.754302	5.867464e+00	
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[51 rows x 51 columns]

23) An interesting way to suggest outliers is to take a distance matrix, aggregate the mean over each row or column, then sort the output in order. Why would this work? Because an observation that is an outlier should have a relatively high mean distance to all other observations! Do this, and output the entire ordered list in descending order. (HINT: DC should be your largest outlier.)

```
[30]: """
This works because the average distance is how
```

```
closely the data connects with other datas in
the list. If the average is high, then it means
the data is further and less similar, which is
the idea of an outlier.
"""

distmat_cars.mean().sort_values(ascending=False)

abbrev

DC = 5.265664
```

```
[30]: abbrev
     DC
            5.265664
            5.257310
     ND
      SC
            5.214264
     LA
            4.767480
     MT
            4.759816
     WV
            4.512766
     NJ
            4.310802
            4.197513
     MA
     MS
            4.048305
            3.897635
      AR
     MD
            3.749939
     MN
            3.719807
      OK
            3.618059
     HI
            3.591755
      SD
            3.585247
      TX
            3.572109
      NY
            3.531806
            3.514075
      CT
```

WI

ID

PA UT

RI WA

FL

VA

OR

CA

ME

NH

ΙA

AK

ΜI

NE

AL

DE

ΑZ

TN

3.468777

3.440899 3.440610

3.438146 3.434056

3.296993

3.205449

3.156358

3.130498

3.116879

3.105251

3.100905

3.098437

3.091304

3.072811

3.065206

3.033363

3.021971

2.999864

2.965254

17

```
ΚY
      2.964333
NM
      2.914953
IN
      2.841386
WY
      2.839487
VT
      2.820077
NC
      2.777656
OH
      2.746863
MO
      2.734449
GA
      2.726648
NV
      2.627495
CO
      2.617432
ΙL
      2.602264
KS
      2.573618
dtype: float64
```

24) From this analysis, which 4 states seem to be strongest outliers?

```
[31]: distmat_cars.mean().sort_values(ascending=False)[:4].index.values.tolist()
```

```
[31]: ['DC', 'ND', 'SC', 'LA']
```

The outliers are DC, ND, SC, LA.

25) OK. Let's explore the data visually. First, using the original, unscaled data frame df_car_crashes, create a scatter plot of insurance premiums vs. insurance losses, with total number of accidents as the size of the point. Create a label near to every point representing the two letter state code.

Insurance premium vs Insurance Losses across US states



26) Next, generate two interesting plots that show some relationships between variables in the data. Try to use as many variables as you can without creating chaos! Don't just throw in multiple variables for the sake of showing them, only include them if it makes sense to do so. Your aim is to derive meaning from your data. Good visualizations tell a story. Strive to use at least one additional variable as size, color, or shape in your data, so you can show more than just 2 variables on a single plot. Add titles, legends and label your axes as appropriate.

[33]:

11 11 1

The two plots I chose to show are the relationships between the variables in a scatter plot using total crashes as color. In the resulting matrix plot, it shows that this data is insufficient to draw a conclusion between alcohol, speeding, and not distracted. This is because they are not normalized with the total amount. The top lefts all have higher totals regardless of the type of crash.

The matrix also shows that in general, total crashes tends to mean that the insurance will be higher. This should be expected, but these plots force us to use normalized data.

The second plot uses percentages of the total in relation with the alcohol us speeding. This is then sorted to show the percent of speeding normalized with the total. This shows that most states have more speeding than alcohol crashes, which should be the norm. The states with higher alcohol than speeding also seem to have 'notorious drivers' or are more rural states. An outlier is also seen in utah where there is very littler drinking in general due to their demographic.

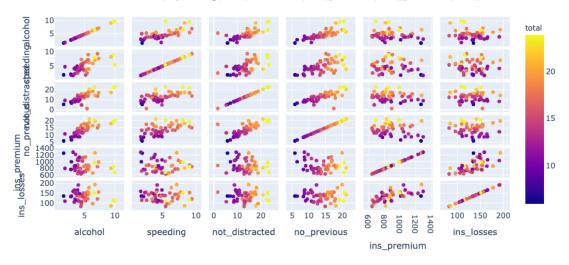
Another outlier is MT, which has the law that allows for the consumptiion of alcohol while driving if you're under a certain threshold.

```
fig = px.scatter_matrix(df_car_crashes, dimensions=["alcohol", "speeding", |
→"not_distracted", "no_previous", "ins_premium", "ins_losses"], color="total")
fig.update_layout(title_text='Scatter Matrix of alcohol, speeding, u
→not_distracted, no_previous, ins_premium, ins_losses')
fig.show()
df_car_crashes_percent = df_car_crashes.copy()
df_car_crashes_percent.alcohol /= df_car_crashes_percent.total
df_car_crashes_percent.speeding /= df_car_crashes_percent.total
df_car_crashes_percent.not_distracted /= df_car_crashes_percent.total
df_car_crashes_percent = df_car_crashes_percent.sort_values(by=['speeding'])
fig = px.line(df car crashes percent, x=df car crashes percent.

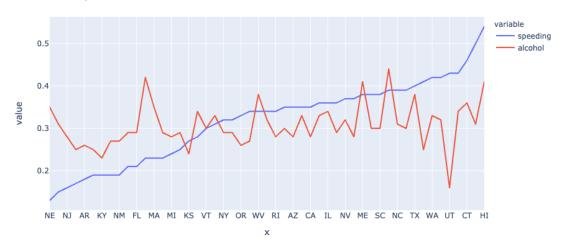
→sort_values(by=['speeding']).index, y=['speeding', 'alcohol'],

→render_mode='webgl')
fig.update layout(title text='Insurance premium vs Insurance Losses across US_
⇔states')
fig.show()
```

Scatter Matrix of alcohol, speeding, not_distracted, no_previous, ins_premium, ins_losses



Insurance premium vs Insurance Losses across US states



27) Run a full PCA on the z_score transformed data. Set n_components to be the same number of columns as the data. Be sure to fit the data to your PCA model, and then output the components, explained variance, and the explained variance ratio.

```
[34]: pca = PCA(n_components=len(df_car_crashes_zscore.columns))
    pca.fit(df_car_crashes_zscore)
    print("Components:\n", pca.components_, "\n")
    print("Explained Variance:\n", pca.explained_variance_, "\n")
    print("Explained Variance Ratio:\n", pca.explained_variance_ratio_)
```

Components:

components.					
[[0.47947078	0.37534719	0.45437635	0.4380328	0.45703414	-0.1308319
-0.06996048]					
[0.06893769	0.0765846	0.03345835	0.04237473	0.0961294	0.6852266
0.71252436]					
[-0.26908802	0.81826935	0.08293253	-0.12616845	-0.31798812	0.25614247
-0.26173503]					
[0.0389558 -	-0.36374887	0.14834351	0.1712655	0.03948141	0.65639617
-0.61839859]					
[0.14597659 -	-0.02282818	0.35479821	-0.85646854	0.33611019	0.04214531
-0.06327152]					
[-0.16970508 -	-0.22479246	0.7837677	0.08510479	-0.50401185	-0.11577348
0.17805184]					
[0.80082487	0.01784783	-0.15285774	-0.14247844	-0.55875371	0.04126619
-0.02804966]]					

Explained Variance:

```
[4.0942308 1.6095732 0.56161403 0.35753958 0.2863854 0.20263316 0.02802383]
```

```
Explained Variance Ratio:
[0.57342168 0.22543042 0.07865743 0.05007557 0.04011 0.02837999 0.00392491]
```

- 28) Use your intuition what do the weights of the first couple of components suggest where most of the variance in the data is coming from? It looks like the weights tend to be higher for the first two components. It is much lower for the latter few. This can also be seen in the fact that the first two comps have the highest variance ratio.
- 29) Create a plot of the cumulative sum of the explained variance. How many components will get you to 90% of the explained variance?

```
[35]: # 3 components

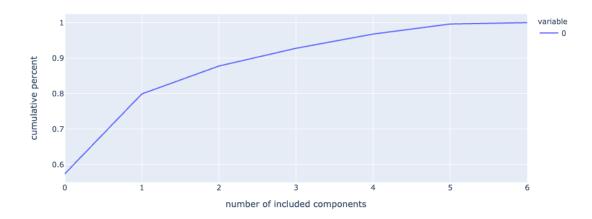
fig = px.line(pca.explained_variance_ratio_.cumsum(), labels={'index':'number_

→of included components','value':'cumulative percent'}, title='Number of_

→components needed for 90% cumulative variance', render_mode='webgl')

fig.show()
```

Number of components needed for 90% cumulative variance

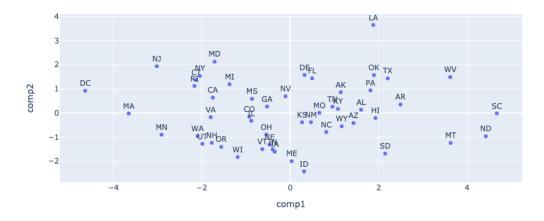


30) Transform the z_score transformed data using your PCA model (i.e. using the transform function of the pca object.) (NOTE: I often just store the transformed data temporarily as some arbitrary variable, X, to make it easier to manipulate the data for plotting.)

```
[36]: X = pca.transform(df_car_crashes_zscore)
    df_car_crashes_comps = df_car_crashes.copy()
    df_car_crashes_comps['comp1'] = X[:,0]
    df_car_crashes_comps['comp2'] = X[:,1]
    df_car_crashes_comps['comp3'] = X[:,2]
```

31) Generate a 2D plot using the first two principal components as your x and y coordinates. Be sure to label each point, and label your axes as component 1 and component 2, respectively.

Component 1 vs Component 2 of PCA



- 32) Compare the states you reported as potential outliers above to those that appear to be outliers from your plot. Do the same results seem to hold? The results still hold, we can tell that LA, DC, SC, and ND are still on the furthest edges of the plot.
- 33) Read how to generate a 3D scatterplot in seaborn or plotly, and use it to generate a scatterplot of the first 3 components.

```
[38]: fig = px.scatter_3d(df_car_crashes_comps, x='comp1', y='comp2', z='comp3', □

→text=df_car_crashes.index, title="Component 1 vs Component 2 vs Component 3 □

→of PCA")

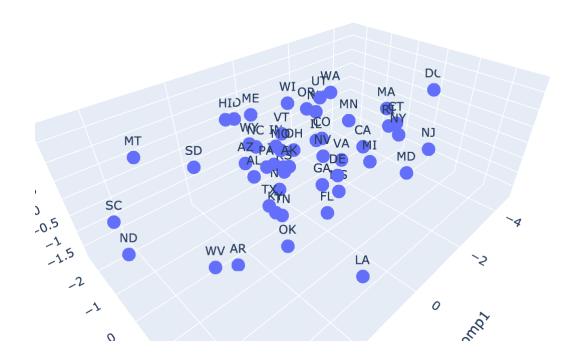
fig.update_layout(scene = dict(xaxis_title='comp1', yaxis_title='comp2', □

→zaxis_title='comp3'))

fig.show()
```

34) Do the same outliers still stand out? Yes, they four outliers are still the furthest away from the center cluster of points. But it also looks like MT and AR are also quite far away and could be seen as outliers too.

Component 1 vs Component 2 vs Component 3 of PCA



34) Do the same outliers still stand out?

Yes, they four outliers are still the furthest away from the center cluster of points. But it also looks like MT and AR are also quite far away and could be seen as outliers too.