Assignment 6: Text Analytics: String Distances

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In [4]: # loading the necessary libraries
          import pandas as pd
          import numpy as np
          import re
          import matplotlib.pyplot as plt
          import jellyfish
         input name = "Mukund" # initializing my name
 In [5]:
          babynames = pd.read csv("babyNamesUSYOB-full.csv") # Loading the babynames dataset
 In [6]:
         babynames.head()
 In [7]:
            YearOfBirth
 Out[7]:
                          Name Sex Number
          0
                  1880
                           Mary
                                        7065
                  1880
                          Anna
                                        2604
          2
                  1880
                          Emma
                                        2003
          3
                  1880
                       Elizabeth
                                        1939
                                  F
          4
                  1880
                         Minnie
                                        1746
 In [8]:
         babynames.shape
         (1858689, 4)
 Out[8]:
         # Preprocessing the babynames data
 In [9]:
          names = babynames['Name'].apply(lambda x: re.sub('[^a-zA-Z]', '', x.lower())).drop_dup
         # defining the distance metrics
In [10]:
          def hamming_distance(a, b):
              """Computes the Hamming distance between two strings."""
              if len(a) != len(b):
                  raise ValueError("Strings must be of equal length.")
              return sum(1 for x, y in zip(a, b) if x != y)
          def longest_common_substring_distance(a, b):
              """Computes the distance between two strings as the length of their longest common
              lengths = np.zeros((len(a) + 1, len(b) + 1))
              for i, x in enumerate(a):
                  for j, y in enumerate(b):
                      if x == y:
```

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lengths[i + 1][j + 1] = lengths[i][j] + 1
   return len(a) + len(b) - 2 * lengths.max()
def levenshtein distance(a, b):
   """Computes the Levenshtein distance between two strings."""
   return jellyfish.levenshtein distance(a, b)
def damerau levenshtein distance(a, b):
    """Computes the full Damerau-Levenshtein distance between two strings."""
   return jellyfish.damerau levenshtein distance(a, b)
def restricted damerau levenshtein distance(s1, s2):
   # Compute the standard Damerau-Levenshtein distance
   d = jellyfish.damerau_levenshtein_distance(s1, s2)
   # Count the number of adjacent transpositions
   transpositions = 0
   for i in range(min(len(s1), len(s2)) - 1):
       if s1[i] != s2[i] and s1[i+1] == s2[i] and s1[i] == s2[i+1]:
           transpositions += 1
   # Subtract the number of adjacent transpositions
   d_restricted = d - transpositions
   return d restricted
def qgram_distance(a, b, q):
    """Computes the q-gram distance between two strings."""
   a qgrams = set(a[i:i+q] for i in range(len(a) - q + 1))
   b_q arms = set(b[i:i+q] for i in range(len(b) - q + 1))
   return 1 - len(a_qgrams & b_qgrams) / len(a_qgrams | b_qgrams)
def jaro_distance(a, b):
    """Computes the Jaro distance between two strings."""
   return jellyfish.jaro_distance(a, b)
def jaro_winkler_distance(a, b, weight=0.1):
    """Computes the Jaro-Winkler distance between two strings."""
   return jellyfish.jaro_winkler(a, b)
def jaccard_distance(a, b, q):
    """Computes the Jaccard distance between two strings based on their q-gram profile
   a_q = set(a[i:i + q]  for i in  range(len(a) - q + 1))
   b_q rams = set(b[i:i+q] for i in range(len(b) - q + 1))
   return 1 - len(a_qgrams & b_qgrams) / len(a_qgrams | b_qgrams)
def cosine_distance(a, b, q):
   """Computes the cosine distance between two strings based on their q-gram profiles
   a_q = set(a[i:i+q]  for i  in range(len(a) - q + 1))
   b_qgrams = set(b[i:i+q] for i in range(len(b) - q + 1))
   intersection = len(a_qgrams & b_qgrams)
   norm_a = np.sqrt(len(a_qgrams))
```

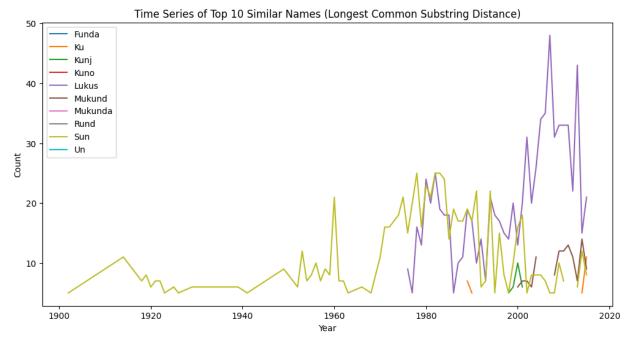
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norm b = np.sqrt(len(b qgrams))
              return 1 - intersection / (norm a * norm b)
          def soundex distance(a, b):
              """Computes the distance between two strings based on their Soundex encoding."""
             soundex a = jellyfish.soundex(a)
              soundex_b = jellyfish.soundex(b)
              return 0 if soundex_a == soundex_b else 1
In [11]: def name distance(name, distance metric):
              """Computes the distance between the input name and another name based on a given
             return distance metric(input name, name)
In [12]: # Computing the distance between the input name and each name in the dataset based on
          # babynames['hamming_distance'] = names.apply(lambda x: name_distance(x, hamming_dista
          babynames['longest_common_substring_distance'] = names.apply(lambda x: name_distance()
          babynames['levenshtein_distance'] = names.apply(lambda x: name_distance(x, levenshteir
          babynames['damerau_levenshtein_distance'] = names.apply(lambda x: name_distance(x, dam
          babynames['restricted_damerau_levenshtein_distance'] = names.apply(lambda x: name_dist
          babynames['qgram distance'] = names.apply(lambda x: name distance(x, lambda a, b: qgra
          babynames['jaro distance'] = names.apply(lambda x: name distance(x, jaro distance))
          babynames['jaro_winkler_distance'] = names.apply(lambda x: name_distance(x, jaro_wink]
          babynames['jaccard distance'] = names.apply(lambda x: name distance(x, lambda a, b: ja
          babynames['cosine_distance'] = names.apply(lambda x: name_distance(x, lambda a, b: cos
          babynames['soundex distance'] = names.apply(lambda x: name distance(x, soundex distance
In [14]:
         # Sorting the dataframes by each distance metric and selecting the top ten names
         # top10_hamming = babynames.sort_values('hamming_distance').head(10)['Name']
         top10 lcs = babynames.sort values('longest common substring distance').head(10)['Name'
          top10 levenshtein = babynames.sort values('levenshtein distance').head(10)['Name']
          top10 dl = babynames.sort values('damerau levenshtein distance').head(10)['Name']
          top10_restricted_dl = babynames.sort_values('restricted_damerau_levenshtein_distance')
          top10 qgram = babynames.sort values('qgram distance').head(10)['Name']
          top10_jaro = babynames.sort_values('jaro_distance', ascending=False).head(10)['Name']
          top10_jaro_winkler = babynames.sort_values('jaro_winkler_distance', ascending=False).
          top10_jaccard = babynames.sort_values('jaccard_distance').head(10)['Name']
          top10_cosine = babynames.sort_values('cosine_distance').head(10)['Name']
          top10_soundex = babynames.sort_values('soundex_distance').head(10)['Name']
         # Printing the top ten names for each distance metric
          # print('Top 10 names based on Hamming distance:')
          # print(top10_hamming)
          # print('')
          print('Top 10 names based on longest common substring distance:')
          print(top10 lcs)
         print('')
          print('Top 10 names based on Levenshtein distance:')
          print(top10_levenshtein)
          print('')
          print('Top 10 names based on full Damerau-Levenshtein distance:')
          print(top10 dl)
          print('')
          print('Top 10 names based on restricted Damerau-Levenshtein distance:')
          print(top10 restricted dl)
```

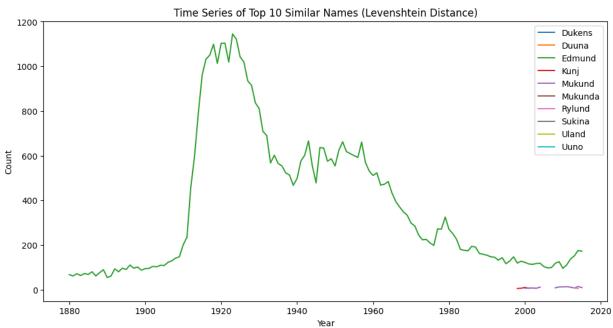
```
print('')
print('Top 10 names based on q-gram distance:')
print(top10_qgram)
print('')
print('Top 10 names based on Jaro distance:')
print(top10_jaro)
print('')
print('Top 10 names based on Jaro-Winkler distance:')
print(top10_jaro_winkler)
print('')
print('Top 10 names based on Jaccard distance:')
print(top10_jaccard)
print('')
print('Top 10 names based on cosine distance:')
print(top10_cosine)
print('')
print('Top 10 names based on Soundex distance:')
print(top10_soundex)
```

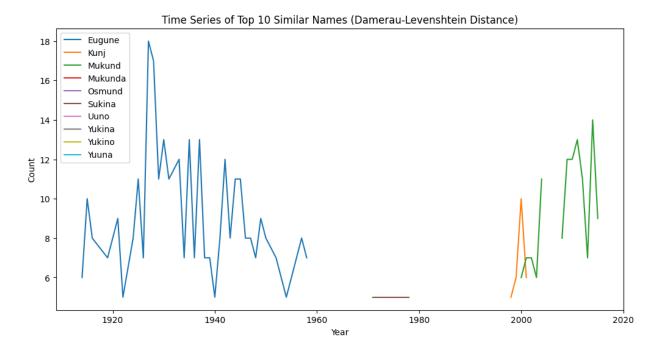
```
Top 10 names based on longest common substring distance:
            Mukund
1272422
1858055
           Mukunda
1587092
              Kuno
              Kunj
1303322
1001470
                Ku
835026
                Un
1405960
              Rund
805735
             Lukus
62489
               Sun
734248
             Funda
Name: Name, dtype: object
Top 10 names based on Levenshtein distance:
1272422
            Mukund
1858055
           Mukunda
725904
            Uland
1303322
              Kunj
1689414
            Rylund
625751
             Duuna
1128
            Edmund
1452989
            Dukens
720816
            Sukina
179470
              Uuno
Name: Name, dtype: object
Top 10 names based on full Damerau-Levenshtein distance:
1272422
           Mukund
           Mukunda
1858055
179470
              Uuno
1318435
            Yukino
720816
            Sukina
1303322
              Kunj
547798
            Osmund
1505530
            Yukina
118981
            Eugune
1711226
             Yuuna
Name: Name, dtype: object
Top 10 names based on restricted Damerau-Levenshtein distance:
1272422
            Mukund
185676
             Ikuko
1858055
           Mukunda
1380291
             Yuuka
922049
            Nikunj
1484079
             Mukul
625751
             Duuna
1331942
             Luken
1587092
              Kuno
720816
            Sukina
Name: Name, dtype: object
Top 10 names based on q-gram distance:
1272422
            Mukund
1858055
           Mukunda
1777325
           Kundana
1303322
              Kunj
1587092
              Kuno
1405960
              Rund
101680
             Lunda
```

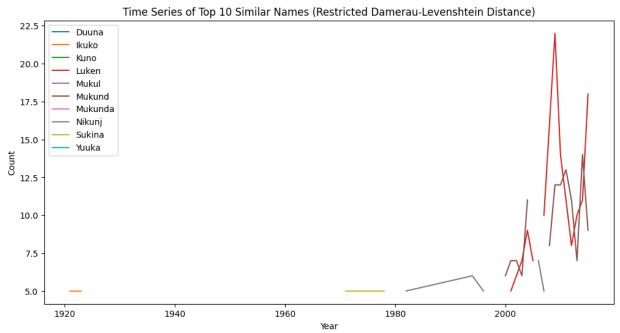
```
704285
             Mundy
1044456
             Tunde
15887
             Gunda
Name: Name, dtype: object
Top 10 names based on Jaro distance:
1272422
             Mukund
1858055
            Mukunda
1405960
               Rund
179470
               Uuno
721013
            Yulunda
757396
              Kindu
              Kendu
941209
919282
           Chukwudi
718862
           Burgundy
           Katelund
1036098
Name: Name, dtype: object
Top 10 names based on Jaro-Winkler distance:
1272422
             Mukund
            Mukunda
1858055
1405960
               Rund
179470
               Uuno
            Yulunda
721013
757396
              Kindu
941209
              Kendu
919282
           Chukwudi
           Burgundy
718862
           Katelund
1036098
Name: Name, dtype: object
Top 10 names based on Jaccard distance:
1272422
            Mukund
1858055
           Mukunda
1777325
           Kundana
1303322
              Kunj
1587092
              Kuno
1405960
              Rund
101680
             Lunda
704285
             Mundy
             Tunde
1044456
15887
             Gunda
Name: Name, dtype: object
Top 10 names based on cosine distance:
            Mukund
1272422
1858055
           Mukunda
           Kundana
1777325
1405960
              Rund
1303322
              Kunj
1587092
              Kuno
657408
             Rundy
1758444
             Kunsh
1448640
             Kunga
822179
             Kunte
Name: Name, dtype: object
Top 10 names based on Soundex distance:
1272422
              Mukund
1207957
             Majenta
```

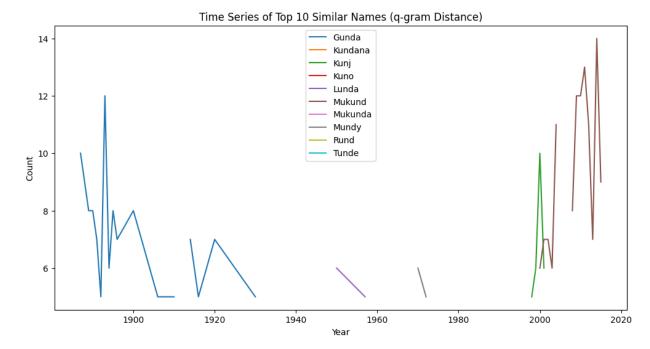
```
719346
                     Mashonda
         875175
                      Magenta
         835816
                     Mishonda
         1285959
                     Makendra
         933673
                    Mashaunda
         677682
                     Mashanda
         837281
                    Mashawnda
                     Mashanti
         1405750
         Name: Name, dtype: object
In [15]: # Plotting the time series
         def plot time series(a, distance metric):
             # Subset the data for the top 10 names
             subset = babynames[babynames['Name'].isin(a)]
             # subset = subset[subset['Name'] != input name] # excluding the input name
             # Pivot the data to create a time series
             pivoted = subset.groupby(['YearOfBirth', 'Name'])['Number'].sum().unstack()
             # Plot
             pivoted.plot(figsize=(12,6))
             plt.title('Time Series of Top 10 Similar Names (' + distance_metric + ')')
              plt.xlabel('Year')
             plt.ylabel('Count')
              plt.legend()
              plt.show()
         # Plotting the time series for each distance metric
          # plot_time_series(top10_hamming, 'Hamming Distance')
          plot_time_series(top10_lcs, 'Longest Common Substring Distance')
         plot_time_series(top10_levenshtein, 'Levenshtein Distance')
          plot_time_series(top10_dl, 'Damerau-Levenshtein Distance')
          plot time series(top10 restricted dl, 'Restricted Damerau-Levenshtein Distance')
         plot_time_series(top10_qgram, 'q-gram Distance')
          plot_time_series(top10_jaro, 'Jaro Distance')
          plot time series(top10 jaro winkler, 'Jaro-Winkler Distance')
          plot time series(top10 jaccard, 'Jaccard Distance')
         plot_time_series(top10_cosine, 'Cosine Distance')
          plot_time_series(top10_soundex, 'Soundex Distance')
```

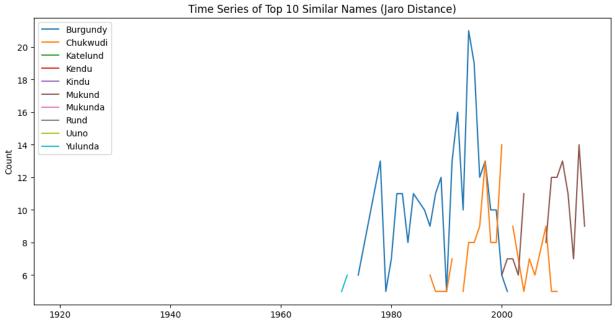




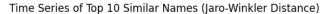


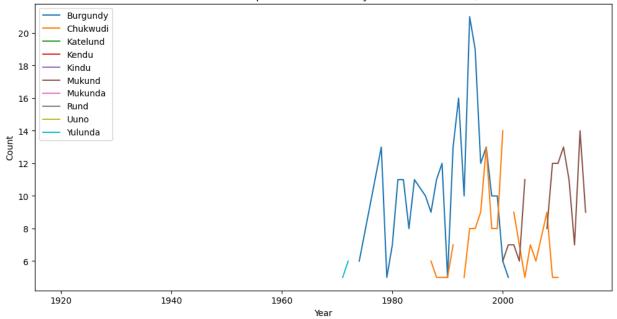




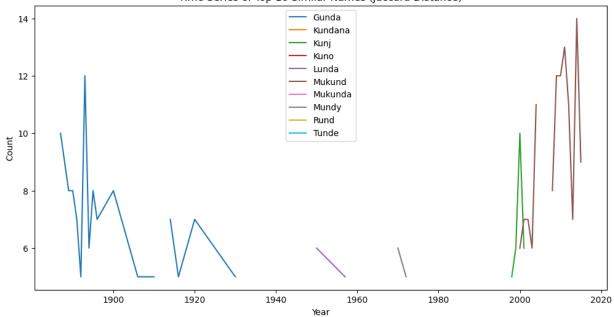


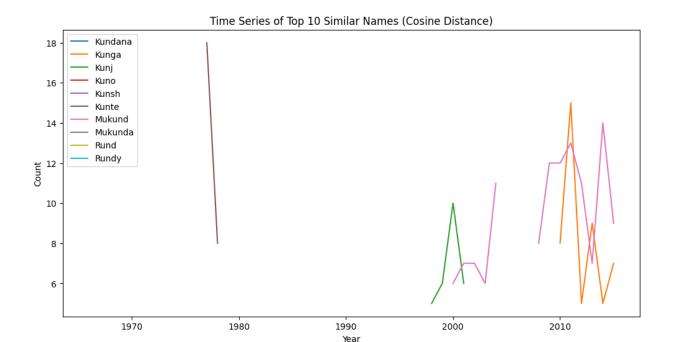
Year

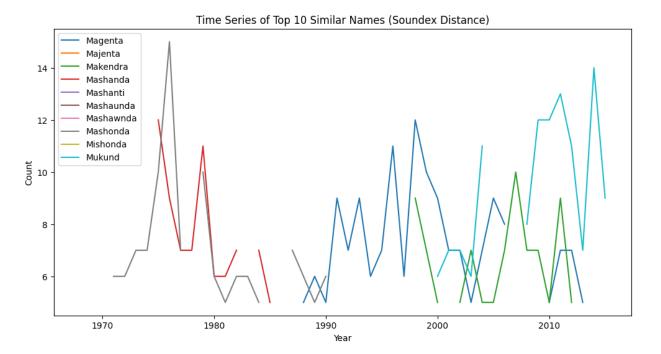




Time Series of Top 10 Similar Names (Jaccard Distance)







It is easy to note that the first name "Mukund" is not so common in the US and it has only started to appear after the 2000s. The "babynames" dataset consists of a large collection of names and their frequency of occurrence over the years. Therefore, the distance metrics that work well with longer strings and are capable of taking into account the semantic meaning of the symbols are likely to perform better on this dataset. Regarding which distance metrics to use for finding the related baby names and their similarities and differences, following are my observations:

The Levenshtein distance, Full Damerau-Levenshtein distance, and q-gram distance are examples of metrics that work well with longer strings and are capable of taking into account the semantic meaning of the symbols. The Levenshtein distance and Full Damerau-Levenshtein distance are capable of handling the insertion, deletion, and substitution of symbols in a string, which is especially useful for measuring the distance between names with different lengths or

with common phonemes but different spellings. The q-gram distance, on the other hand, breaks the names into substrings of a fixed length, which helps to capture the semantic meaning of the symbols and is less sensitive to the order of the symbols.

The Longest Common Substring distance and Jaro distance, while still useful, may not be as effective on this dataset due to the large variation in the length of names and the high degree of variation in the order of the symbols. However, the Jaro-Winkler distance, which gives more weight to the initial characters, may be more effective as it can capture the similarities between names that share a common prefix.

The Hamming distance, while useful in certain contexts, is not as applicable to this dataset as it only works on strings of the same length, and doesn't take into account the semantic meaning of the symbols.

The Jaccard distance and cosine distance between q-gram profiles, while useful for comparing the similarity of short strings, may not be the best choice for the babynames dataset, as names can vary significantly in length, and the size of the q-grams may need to be adjusted to ensure accurate results.

Finally, the distance based on Soundex encoding is useful for comparing the similarity of names that sound alike but are spelled differently. It can help to identify common phonemes and group similar sounding names together, but it may not be as effective in capturing the semantic meaning of the symbols as some of the other distance metrics.