

WH002-read-and-separate

November 4, 2019

```
[210]: from os import listdir
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import scipy.cluster.hierarchy as sch
import pylab
```

```
[3]: files = listdir('data')
print(files)
```

```
['017.TXT', '074.TXT', '041.TXT', '009.TXT', '014.TXT', '004.TXT', '046.TXT',
'099.TXT', '081.TXT', '065.TXT', '025.TXT', '072.TXT', '042.TXT', '066.TXT',
'024.TXT', '015 coughing after approx 90s.TXT', '028.TXT', '069.TXT', '090.TXT',
'036.TXT', '084.TXT', '022.TXT', '052.TXT', '047.TXT']
```

```
[4]: with open('data/'+files[0]) as f:
    first_line = f.readline()
    print(first_line)
```

```
-0.210  -0.111  7.010   21.260  10.071  23.000  0.000   0.992
```

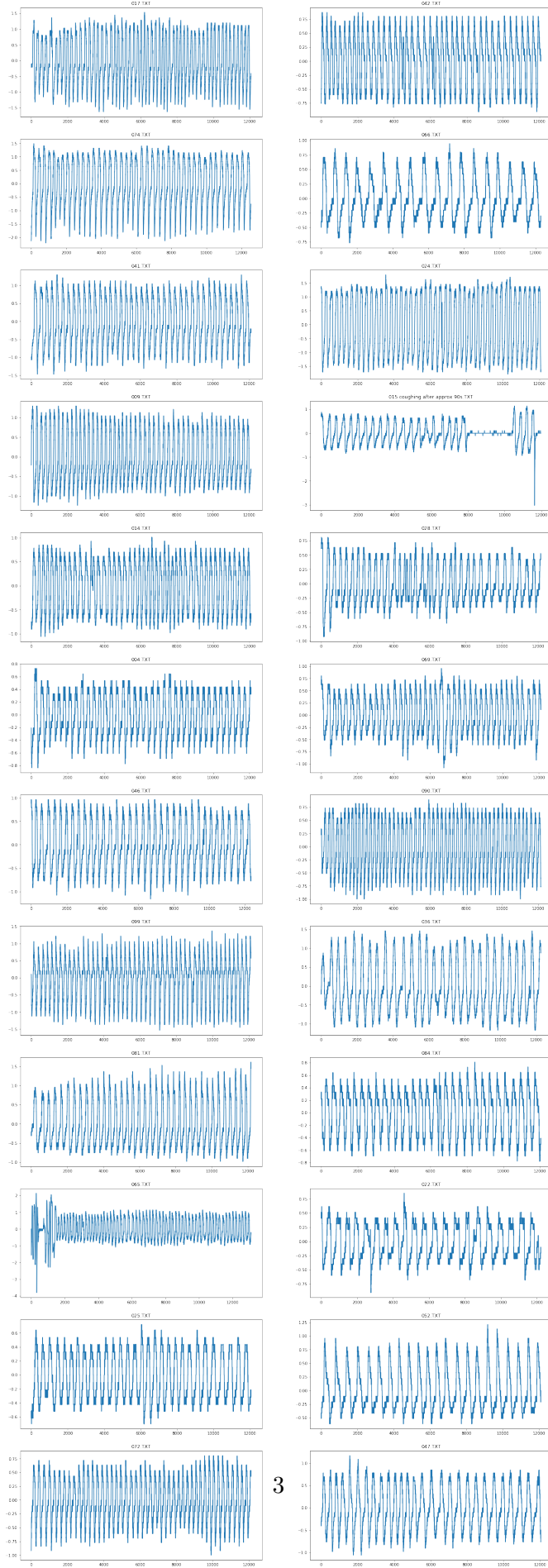
```
[5]: data = np.genfromtxt('data/'+files[0], delimiter='\t')
```

```
[6]: data[0]
```

```
[6]: array([-0.21 , -0.111,  7.01 , 21.26 , 10.071, 23.    ,  0.    ,  0.992])
```

```
[7]: fig, axs = plt.subplots(len(files)//2,2,figsize=(24, len(files)//2 * 6))
p=0
for i in range (2):
    for j in range(len(files)//2):
        if p < len(files):
            data = np.genfromtxt('data/'+files[p], delimiter='\t')
            print(files[p],data.shape)
            axs[j,i].plot(data[:,0])
            axs[j,i].set_title(files[p])
            p+=1
```

017.TXT (12069, 8)
074.TXT (12549, 8)
041.TXT (12090, 8)
009.TXT (12092, 8)
014.TXT (12079, 8)
004.TXT (12076, 8)
046.TXT (12336, 8)
099.TXT (12114, 8)
081.TXT (12168, 8)
065.TXT (13092, 8)
025.TXT (12049, 8)
072.TXT (12107, 8)
042.TXT (12075, 8)
066.TXT (12301, 8)
024.TXT (12082, 8)
015 coughing after approx 90s.TXT (11971, 8)
028.TXT (12155, 8)
069.TXT (12111, 8)
090.TXT (12143, 8)
036.TXT (12261, 8)
084.TXT (12060, 8)
022.TXT (12219, 8)
052.TXT (12099, 8)
047.TXT (12042, 8)



```
[104]: def find_start(signal_data,pos):
        flag = True
        for i in range(1, 4):
            if signal_data[pos - i] > 0:
                flag = False
        if flag:
            for i in range(1, 4):
                if signal_data[pos + i] < 0:
                    flag = False
        return (flag,pos)
```

```
[105]: def separate_cycles(signal_data):
        data_min = 0
        min_list=[]
        flag = False
        for i in range(signal_data.shape[0]-5):
            if signal_data[i]<0:
                flag,pos = find_start(signal_data,i)
            if flag:
                min_list.append(i)
                flag = False
        return min_list
```

```
[153]: def split_cycles(min_list,raw_data):
        cycles=[]
        for i in range(len(min_list)-1):
            d = np.zeros(50)
            c = np.zeros(50)
            j = 0
            l = min_list[i+1]-min_list[i]
            for k in range(min_list[i],min_list[i+1]-1):
                m = int(((k-min_list[i])*50.0)/l)
                d[m]+=raw_data[k]
                c[m]+=1
            for k in range(50):
                d[k] = d[k]/c[k]
            if True in np.isnan(np.array(d)):
                d = []
            else:
                cycles.append(d)
        return cycles
```

```
[154]: def get_file(file_number):
        data = np.genfromtxt('data/'+files[file_number], delimiter='\t')[:,0]
        #data_average = data.mean()
```

```

#data = data - data_average
mins = separate_cycles(data)
cycles = split_cycles(mins,data)
return cycles

```

```
[155]: cycles = get_file(6)
```

```

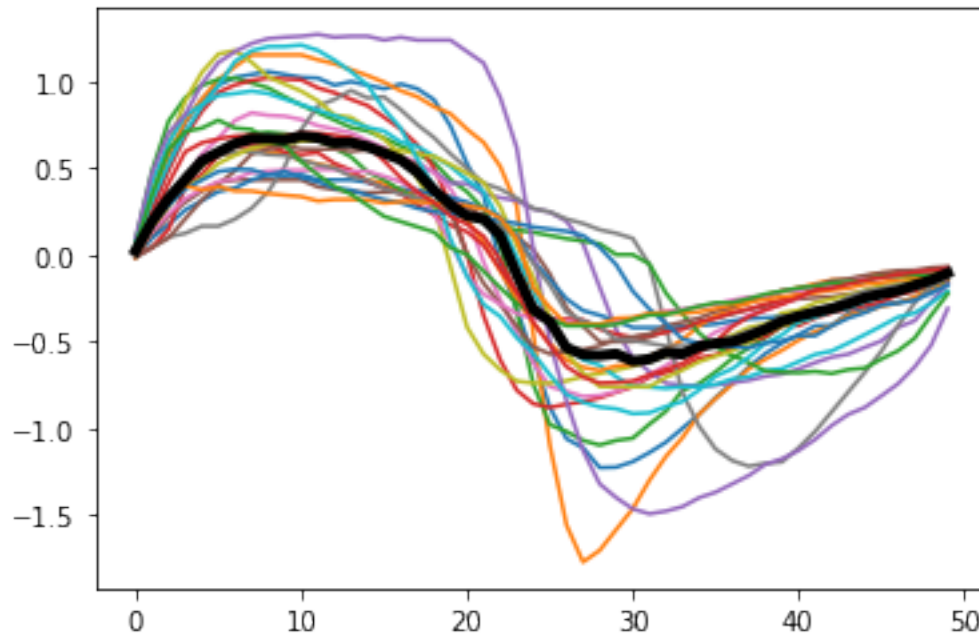
/home/ric/Projects/Python/env/lib/python3.7/site-
packages/ipykernel_launcher.py:13: RuntimeWarning: invalid value encountered in
double_scalars
del sys.path[0]

```

```

[224]: for c in cycles:
        plt.plot(c)
plt.plot(np.median(np.array(cycles),axis=0),linewidth=4, color='black')
plt.show()

```



```

[202]: def sig_plot():
        median_plots = []
        fig, axs = plt.subplots(len(files)//2,2,figsize=(24, len(files)//2 * 6))
        p=0
        for i in range (2):
            for j in range(len(files)//2):
                if p < len(files):
                    cycles = get_file(p)
                    for c in cycles:

```

```

        axs[j,i].plot(c)
        m = np.median(np.array(cycles),axis=0)
        axs[j,i].plot(m,linewidth=4, color='black')
        axs[j,i].set_title(files[p])
        median_plots.append(m)
        p+=1
    return median_plots

```

```

[203]: def difference(a,b):
        count = 0
        for i in range(a.shape[0]-1):
            #print(a[i],b[i])
            count += np.absolute(a[i]-b[i])
        return count/a.shape[0]

```

```

[214]: def dendrogram_heat_map(distance_data,map_data,tags):
        """ Create a heat map with associated dendrogram from a distance N*N numpy_
        ↪matrix and a scipy
        scipy.cluster.hierarchy linkage 2d array"""
        # Adapted from http://stackoverflow.com/users/208339/steve-tjoa
        fig = pylab.figure(figsize=(12,12))

        # Plot first dendrogram.
        ax1 = fig.add_axes([0.09,0.1,0.2,0.6])#set position
        den1 = sch.dendrogram(map_data, orientation='right',labels=tags)
        # No axis labels
        ax1.set_xticks([])
        #ax1.set_yticks([])

        # Plot second dendrogram.
        ax2 = fig.add_axes([0.31,0.74,0.61,0.2])#set position
        den2 = sch.dendrogram(map_data,labels=tags)
        # No axis labels
        #ax2.set_xticks([])
        ax2.set_yticks([])

        # Plot distance matrix as heat map.
        heat_map = fig.add_axes([0.32,0.1,0.6,0.6])
        idx1 = den1['leaves']
        idx2 = den2['leaves']
        # Reorder the distance data so that it matches with the dendrogram order
        distance_data = distance_data[idx1,:]
        distance_data = distance_data[:,idx2]
        hm = heat_map.matshow(distance_data, aspect='auto', origin='lower',
        ↪cmap=pylab.cm.YlGnBu)
        # No Y axis
        heat_map.set_yticks([])

```

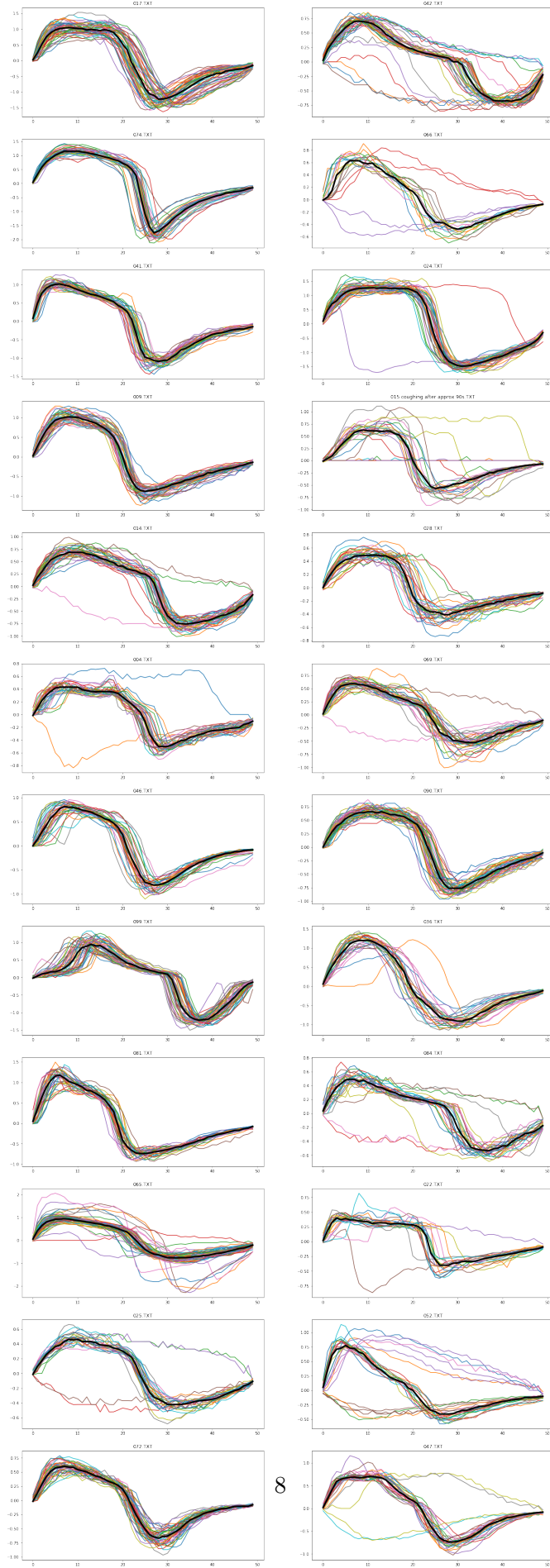
```
heat_map.set_xticks([])

# Plot colorbar.
axcolor = fig.add_axes([0.93,0.1,0.02,0.6])
pylab.colorbar(hm, cax=axcolor)
```

```
[215]: def difference(a,b):
        count = 0
        for i in range(a.shape[0]-1):
            #print(a[i],b[i])
            count += np.absolute(a[i]-b[i])
        return count/a.shape[0]
```

```
[216]: cycles = sig_plot()
```

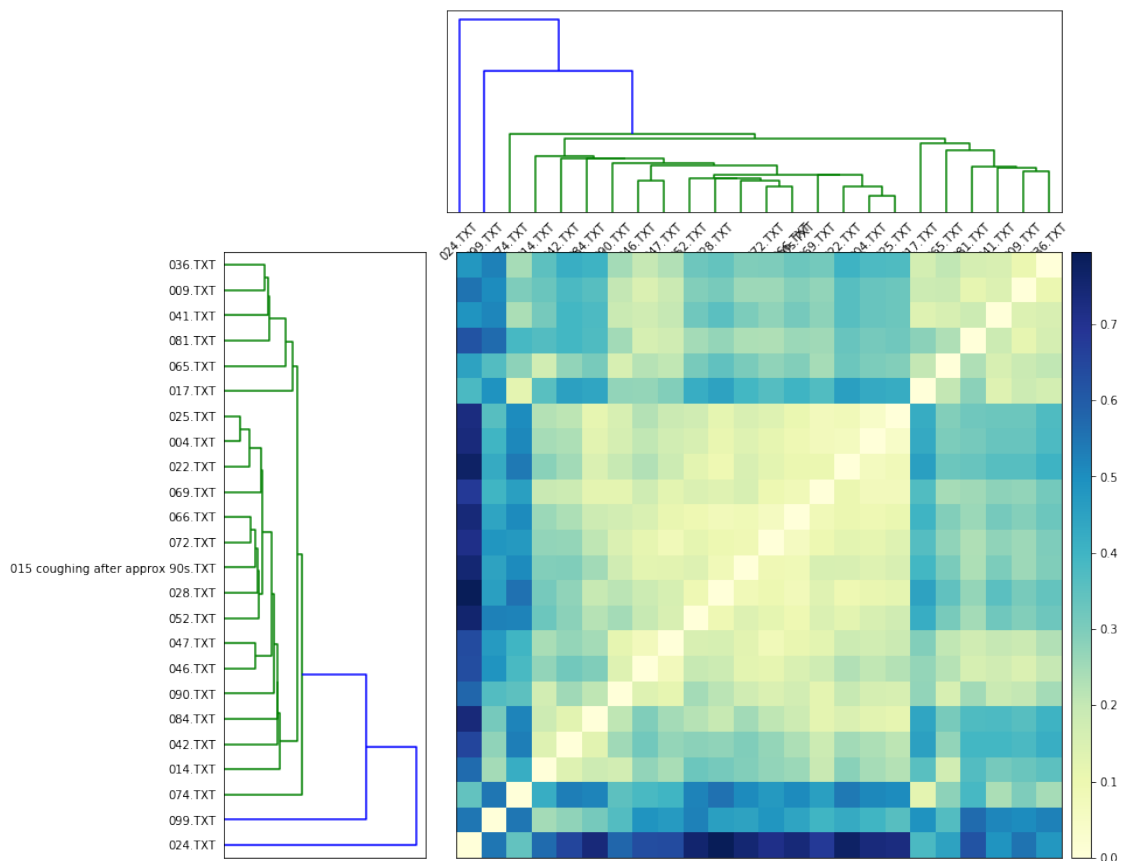
```
/home/ric/Projects/Python/env/lib/python3.7/site-
packages/ipykernel_launcher.py:13: RuntimeWarning: invalid value encountered in
double_scalars
  del sys.path[0]
```




```
[217]: distance = np.zeros((len(cycles),len(cycles)))
for i in range(len(cycles)):
    for j in range(len(cycles)):
        distance[i,j]=difference(cycles[i],cycles[j])
kmap = sch.linkage(distance, method='single')
```

/home/ric/Projects/Python/env/lib/python3.7/site-packages/ipykernel_launcher.py:5: ClusterWarning: scipy.cluster: The symmetric non-negative hollow observation matrix looks suspiciously like an uncondensed distance matrix
 """

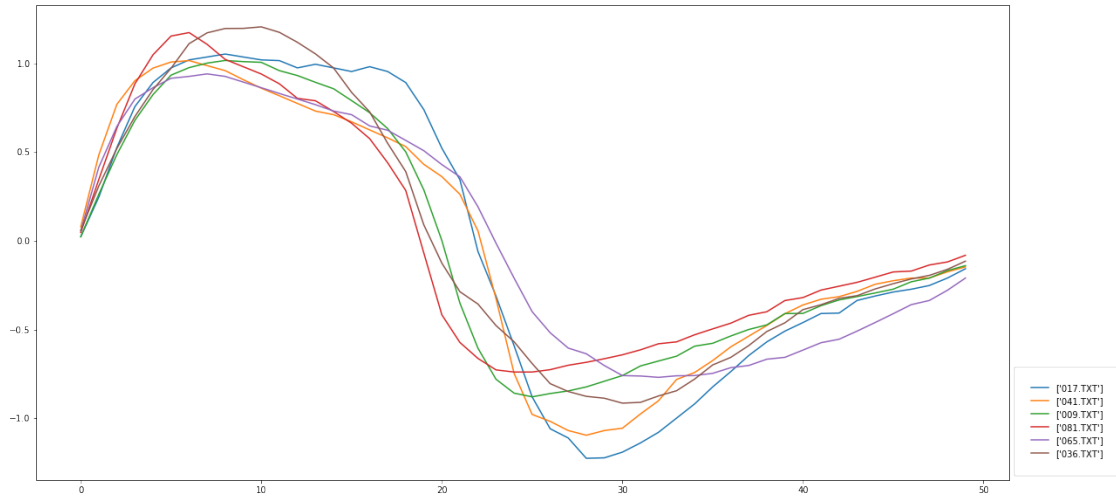
```
[218]: dendrogram_heat_map(distance,kmap,files)
```



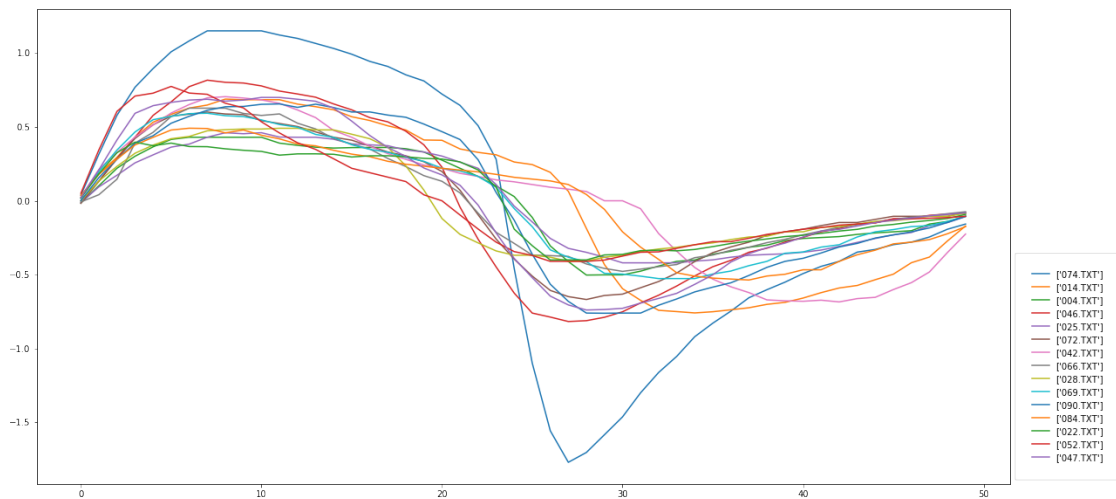
```
[ ]: ['069.TXT', '066.TXT', '072.TXT', '028.TXT', '052.TXT', '047.TXT', '046.TXT', '090.
      ↪TXT', '084.TXT', '042.TXT', '014.TXT', '074.TXT']
```

```
[244]: def plt_group(in_list):
plt.figure(figsize=[20,10])
for i in range(len(cycles)):
    if files[i] in in_list:
        plt.plot(cycles[i],label=[files[i]])
plt.legend(loc=3,borderpad=2,bbox_to_anchor=(1, 0))
plt.show()
```

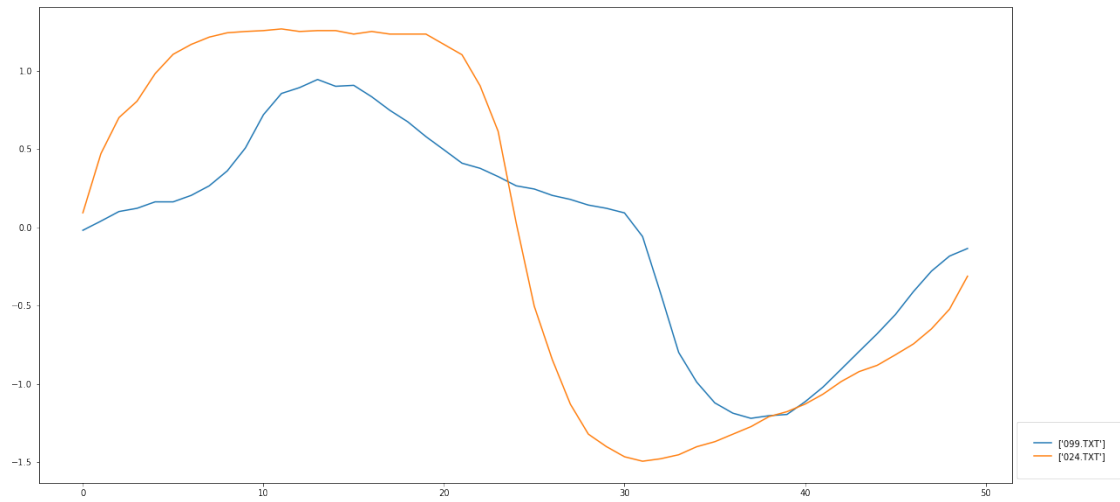
```
[247]: plt_group(['036.TXT','009.TXT','041.TXT','081.TXT','065.TXT','017.TXT'])
```



```
[251]: plt_group(['025.TXT','004.TXT','022.TXT','069.TXT','066.TXT','072.TXT','028.
↪TXT','052.TXT','047.TXT','046.TXT','090.TXT','084.TXT','042.TXT','014.
↪TXT','074.TXT'])
```



```
[253]: plt_group(['099.TXT', '024.TXT'])
```



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[ ]:
```