## 2.1) Socket Programming with Data Buffering.

The code below gets the data from the sensor logger app using a socket server. I used a buffer for batch processing of the data which is to be parsed and exported. The data is parsed into a csy file.

```
import dash
In [ ]:
        from dash.dependencies import Output, Input
        from dash import dcc, html, dcc
        from datetime import datetime
        import json
        import plotly.graph_objs as go
        from collections import deque
        from flask import Flask, request
        import csv
        import numpy as np
        server = Flask(__name__)
        app = dash.Dash(__name__, server=server)
        MAX DATA POINTS = 1000
        UPDATE_FREQ_MS = 100
        buffer= []
        accMat= np.array([1,2,3])
        norm= np.array([1,2,3])
        totalaccMat= np.array([1,2,3,4,5,6,7,8,9,10,11,12,13,14,15])
        # Create a CSV file for storing sensor data
        csv_filename = "sensor_data.csv"
        with open(csv_filename, "w", newline="") as csv_file:
            fieldnames = ["Sensor", "Timestamp", "X", "Y", "Z"] # Define CSV column names
             csv_writer = csv.DictWriter(csv_file, fieldnames=fieldnames)
             csv_writer.writeheader() # Write the header row
        time = deque(maxlen=MAX_DATA_POINTS)
        sensor_name= deque(maxlen= MAX_DATA_POINTS)
        accel x = deque(maxlen=MAX DATA POINTS)
        accel y = deque(maxlen=MAX DATA POINTS)
        accel_z = deque(maxlen=MAX_DATA_POINTS)
        app.layout = html.Div(
                dcc.Markdown(
                                 children="""
                                 # Live Sensor Readings
                                 Streamed from Sensor Logger: tszheichoi.com/sensorlogger
                         ....
                         ),
                         dcc.Graph(id="live_graph"),
                         dcc.Interval(id="counter", interval=UPDATE_FREQ_MS),
                ]
```

```
@app.callback(Output("live graph", "figure"), Input("counter", "n intervals"))
def update_graph(_counter):
        data = [
                go.Scatter(x=list(time), y=list(d), name=name)
                for d, name in zip([accel x, accel y, accel z], ["X", "Y", "Z"])
        1
        graph = {
                "data": data,
                "layout": go.Layout(
                                "xaxis": {"type": "date"},
                                "yaxis": {"title": "Acceleration ms<sup>-2</sup>"},
                        }
                ),
        }
        if (
                len(time) > 0
        ): # Â cannot adjust plot ranges until there is at least one data point
                graph["layout"]["xaxis"]["range"] = [min(time), max(time)]
                graph["layout"]["yaxis"]["range"] = [
                        min(accel_x + accel_y + accel_z),
                        max(accel_x + accel_y + accel_z),
                1
        return graph
def csvMaker(filename, buf):
     with open(filename, mode = 'a+', newline='') as csv_file:
          csv writer = csv.writer(csv file)
          for row in buf:
               sensor= row[0]
               timestamp= row[1]
               x = row[2]
               y = row[3]
               z = row[4]
               csv writer.writerow([sensor,timestamp,x,y,z])
@server.route("/data", methods=["POST"])
def data():
    global accMat, totalaccMat, norm
    if str(request.method) == "POST":
        data = json.loads(request.data)
        for d in data['payload']:
            sensor= d.get("name", None)
            timestamp= datetime.fromtimestamp(d["time"] / 1000000000)
            xVal= (d["values"]["x"])
            yVal= (d["values"]["y"])
            zVal= (d["values"]["z"])
            row = [sensor, str(timestamp), str(xVal), str(yVal), str(zVal)]
            buffer.append(row)
            aRow= [xVal, yVal, zVal]
            accMat= np.vstack((accMat, aRow))
            norm= np.vstack((norm, aRow))
            if(len(buffer)==5):
                 csvMaker(csv filename, buffer)
                 buffer.clear()
```

```
aMat= accMat[1:]
    sampleARow= aMat.flatten()
    totalaccMat= np.vstack((totalaccMat,sampleARow))
    accMat= accMat[0]

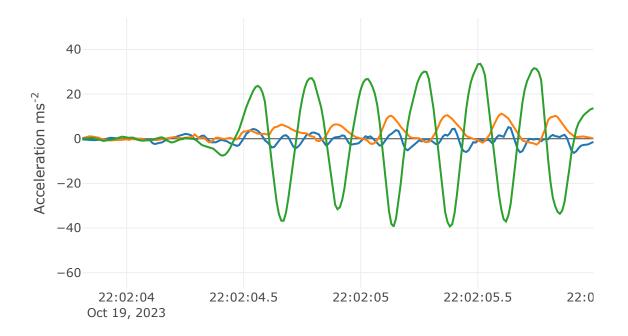
if d.get("name", None) == "accelerometer" or d.get("name", None) == "gravi
    ts = datetime.fromtimestamp(d["time"] / 10000000000)
    if len(time) == 0 or ts > time[-1]:
        time.append(ts)
        accel_x.append(d["values"]["x"])
        accel_y.append(d["values"]["y"])
        accel_y.append(d["values"]["z"])
        sensor_name.append(d.get("name", None)) # Proper indentation here

return "success"

if __name__ == "__main__":
    app.run_server(port=8000, host="172.31.193.134")
```

## **Live Sensor Readings**

Streamed from Sensor Logger: tszheichoi.com/sensorlogger



Part 2.2) PCA with covariance matrix.

1)The code below uses the numpy library to calculate the covariance data of the sensor data.

```
import numpy as np

def covarianceCalculate():
    totalAMat= totalaccMat[1:]
    totalAMat= totalAMat.T
    cov= np.cov(totalAMat)
    return cov, totalAMat, norm

cov= covarianceCalculate()
```

2)The code below uses sklearn to do Principle Component Analysis(PCA). First it standardizes the data and runs the fit\_transofrm method on the standardized data to return the pca\_features

matrix. It also returns the explained variance ratio for the pca features.

```
In []: from sklearn.preprocessing import StandardScaler
    from sklearn.decomposition import PCA

def PCAFunction():
        scaler= StandardScaler()
        std= scaler.fit_transform(totalaccMat)
        pca= PCA(n_components=2)
        pca_features= pca.fit_transform(std)
        explained_ratio= pca.explained_variance_ratio_

        return pca_features, explained_ratio

pca_feat, explained_rat= PCAFunction()
        pca_feat.shape
```

Out[ ]: (81, 2)

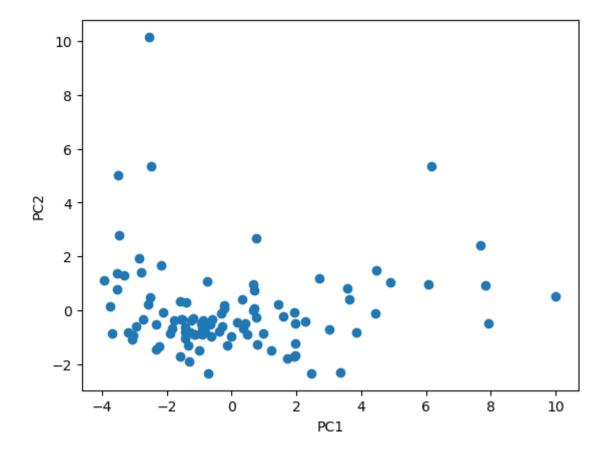
3)The code below gets the pca\_features of a sensor recording data and visualizes it onto a lower dimensional space.

```
import matplotlib.pyplot as plt
cov, accMat, normalArr= covarianceCalculate()
normalArr= normalArr[1:]
pca_features, explained_ratio= PCAFunction()

print(explained_ratio)

plt.scatter(pca_features[:,0], pca_features[:,1])
plt.xlabel('PC1')
plt.ylabel('PC2')
plt.show()
```

[0.53109284 0.19984808]



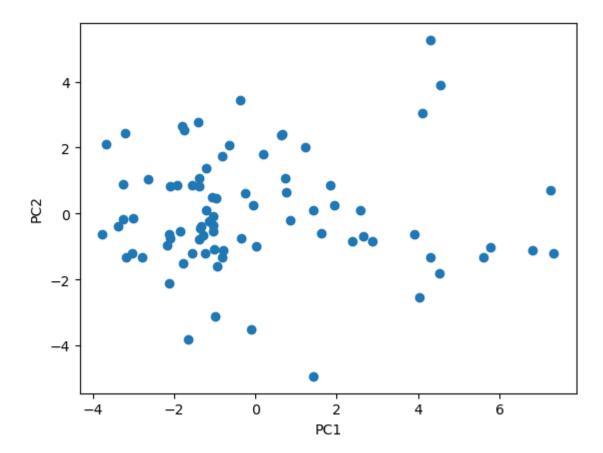
The code below is the same as the code above but uses new variables to store the data of a different recording.

```
import matplotlib.pyplot as plt
cov2, accMat2, normalArr2= covarianceCalculate()
normalArr2= normalArr2[1:]
pca_features2, explained_ratio2= PCAFunction()

print(explained_ratio2)

plt.scatter(pca_features2[:,0], pca_features2[:,1])
plt.xlabel('PC1')
plt.ylabel('PC2')
plt.show()

[0.47775693 0.19863499]
```



Principal Component Analysis was performed on 2 separate recordings above and the explained variance ratios were printed out. As you can see in the output for the first recording above, the variance explained by the first principal component is about 55% and the variance explained by the second principal component is about 23%. For the second recording, the variance explained by the first principal component is about 62% and the variance explained by the second principal component is about 22%.

Part2.3) Dynamic Time Warping (with attempted early abandoning).

1)The code below defines a dtw\_function which does dynamic time warping on specific dimensions. It compares the x-values, y-values and z-values from 2 separate recordings. The function also finds the dtw distances for each dimension. The DTW is used for the purpose of finding the similarity between two time sequences in the same dimension which may vary.

```
In []: from dtw import *
    import numpy as np

def dtw_function(x_arr1, x_arr2, y_arr1, y_arr2, z_arr1, z_arr2):
        x_dtw= dtw(x_arr1, x_arr2, keep_internals= True)
        x_dtw.plot(type="twoway")
        y_dtw= dtw(y_arr1, y_arr2, keep_internals= True)
        y_dtw.plot(type="twoway")
        z_dtw= dtw(z_arr1, z_arr2, keep_internals= True)
        z_dtw.plot(type="twoway")

# Get the alignment path as a list of pairs of indices
        x_alignment_path = list(zip(x_dtw.index1, x_dtw.index2))
        y_alignment_path = list(zip(y_dtw.index1, y_dtw.index2))
```

```
z_alignment_path = list(zip(z_dtw.index1, z_dtw.index2))

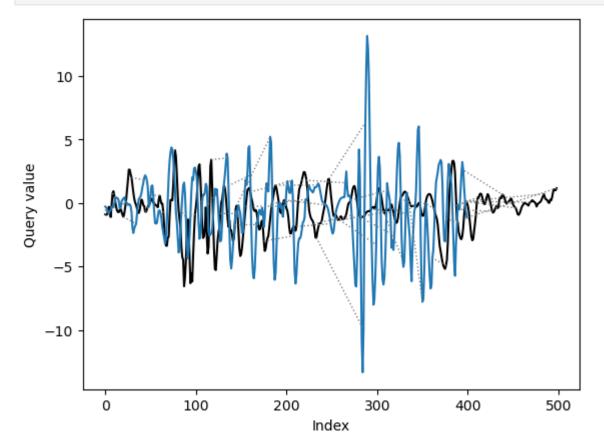
# Calculate distances between corresponding points
x_distances = [abs(x_arr1[i] - x_arr2[j]) for i, j in x_alignment_path]
y_distances = [abs(y_arr1[i] - y_arr2[j]) for i, j in y_alignment_path]
z_distances = [abs(z_arr1[i] - z_arr2[j]) for i, j in z_alignment_path]
return x_distances, y_distances, z_distances
```

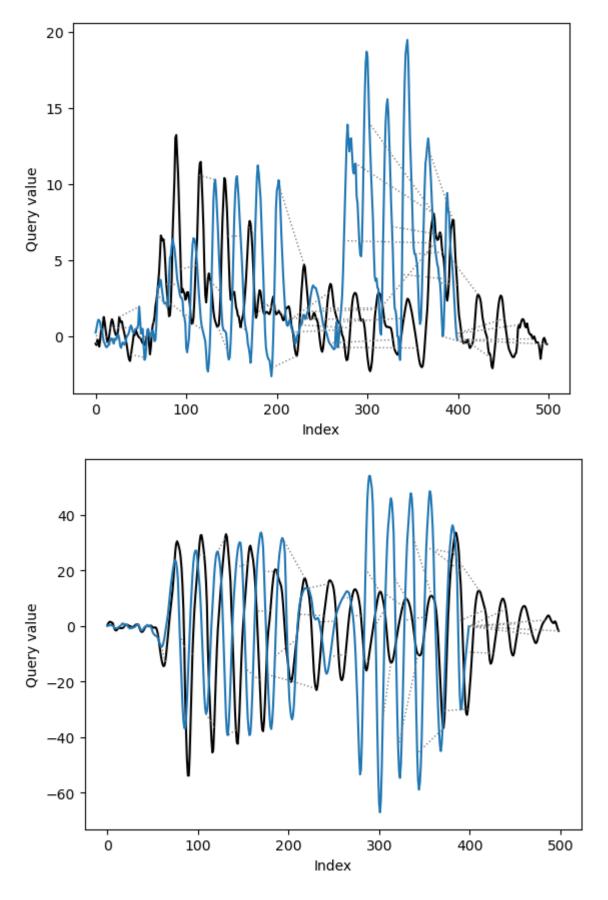
2)The below code block runs the dtw\_function on all dimensions from 2 different recordings.

```
In []: x_arr1= normalArr[:,0]
x_arr2= normalArr2[:,0]

y_arr1= normalArr[:,1]
y_arr2= normalArr2[:,1]

z_arr1= normalArr[:,2]
z_arr2= normalArr2[:,2]
x_dtw_distance, y_dtw_distance, z_dtw_distance= dtw_function(x_arr1, x_arr2, y_arr1, y)
```





Part 2.4) Anomaly Detection. In this part, I used sklearn and OneClassSVM to do anomoly detection using a one class svm model.

The libraries used are imported below.

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt

from sklearn.model_selection import train_test_split
from sklearn.svm import OneClassSVM
```

1)The code below splits the data into training data and testing data from the pca features matrix which was found in part 2.2.

```
In [ ]: acc_train, acc_test= train_test_split(pca_features2, test_size=0.2, random_state=42)
    print(acc_train.shape[0])
    print(acc_test.shape[0])

    one_class_svm= OneClassSVM(nu=0.05, kernel = 'rbf', gamma= 'auto').fit(acc_train)
    score= one_class_svm.score_samples(acc_test)
    score_threshold= np.percentile(score,2)

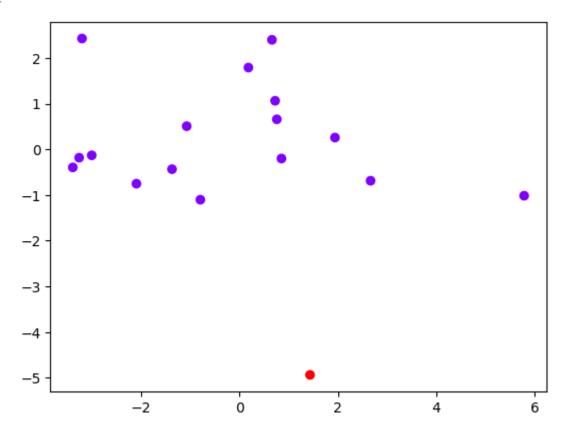
    prediction= [1 if i<score_threshold else 0 for i in score]</pre>
```

The code below visualizes the anomaly detection model.

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```
In [ ]: df_test= pd.DataFrame(acc_test, columns=['feat1', 'feat2'])
    df_test['anomoly detection']= prediction
    plt.scatter(df_test['feat1'], df_test['feat2'], c=df_test['anomoly detection'], cmap='
```

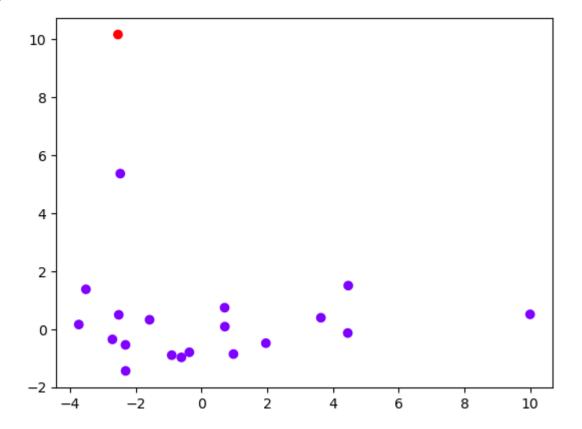
Out[ ]: <matplotlib.collections.PathCollection at 0x1faff8c3820>



The code below runs anomaly detection on the second recording.

```
In [ ]:
        acc_train, acc_test= train_test_split(pca_features, test_size=0.2, random_state=42)
        print(acc train.shape[0])
        print(acc_test.shape[0])
        one_class_svm= OneClassSVM(nu=0.05, kernel = 'rbf', gamma= 'auto').fit(acc_train)
         score= one_class_svm.score_samples(acc_test)
         score threshold= np.percentile(score,2)
        prediction= [1 if i<score threshold else 0 for i in score]</pre>
        df_test= pd.DataFrame(acc_test, columns=['feat1', 'feat2'])
        df_test['anomoly detection'] = prediction
        plt.scatter(df_test['feat1'], df_test['feat2'], c=df_test['anomoly detection'], cmap='
        80
        20
```

<matplotlib.collections.PathCollection at 0x1fa8a48e5b0> Out[ ]:



2) An attempt on using DTW distances between timeseries as features and improving the anomaly detection model. As mentioned above, the dtw\_function which was implemented returns the lists of distances between each pair of corresponding points in the time series data for the X,Y and Z dimensions. The 3 variables x\_dtw\_distance, y-dtw\_distance and z\_dtw\_distance, hold the dtw distances. In this code below, dtw\_distances are included as features in the anomaly detection model. The DTW distances are combined into one matrix, and the One-Class SVM model is trained on this combined dataset. This approach allows the model to leverage both the PCA features and the DTW distances for improved anomaly detection.

```
In []: min_len= min(len(x_dtw_distance), len(y_dtw_distance), len(z_dtw_distance))
    x_dtw_distance= x_dtw_distance[0:min_len]
    y_dtw_distance= y_dtw_distance[0:min_len]
    z_dtw_distance= z_dtw_distance[0:min_len]

    dist_mat= np.array((x_dtw_distance, y_dtw_distance, z_dtw_distance), dtype=float)
    dist_mat= dist_mat.T
    acc_train, acc_test = train_test_split(dist_mat, test_size=0.2, random_state=42)

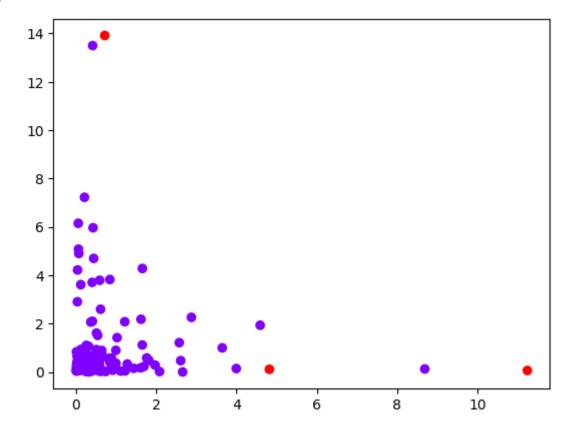
one_class_svm = OneClassSVM(nu=0.05, kernel='rbf', gamma='auto').fit(acc_train)

score = one_class_svm.score_samples(acc_test)

score_threshold = np.percentile(score, 2)
    prediction= [1 if i<score_threshold else 0 for i in score]
    df_test= pd.DataFrame(acc_test, columns=['feat1', 'feat2', 'feat3'])
    df_test['anomoly detection']= prediction

plt.scatter(df_test['feat1'], df_test['feat2'], c=df_test['anomoly detection'], cmap='</pre>
```

Out[ ]: <matplotlib.collections.PathCollection at 0x1fa88e9e940>



The plot above shows multiple anomalies with this improved anomaly detection model. Since this is only a 2D plot, the anomalies shown within the cluster may have obscure values in the Z dimension.