

LASER INTERFEROMETER GRAVITATIONAL WAVE OBSERVATORY
- LIGO -
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Technical Note	LIGO-T1700198-v1	2017/09/22
Online Detector Characterization using Neural Networks		
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1 Abstract

LIGO data has noise that comes from many sources. To be able to better distinguish gravitational wave signals from the noise, it is important to characterize the type of noise observed. Machine learning algorithms can be used to look for patterns within the data and to classify the noise into distinct categories. We apply clustering algorithms, such as kmeans, to identify earthquakes in seismic noise data. To test how well the clustering algorithms correctly pick out earthquake times, we compare the clusters to the times when earthquake waves reach a detector site. This comparison will be used to evaluate how well a neural network identifies earthquakes compared to the clustering algorithms.

2 Introduction

LIGO consists of two detectors, one in Hanford, Washington and one in Livingston, Louisiana that use laser interferometry to detect gravitational wave signals. LIGO data contains noise that comes from many sources. In order to be able to better distinguish signals from the noise, it is important to characterize the type of noise observed. Machine learning algorithms can be used to look for patterns within the data and to classify or cluster the data into different categories.

There are many sensors near the LIGO detectors that measure the behavior of the detector environment. For example, there are several seismometers around each LIGO detector that measure seismic noise in different frequency channels in each of the horizontal and vertical directions. Within the data, there are different types of seismic noise such as earthquakes and anthropogenic noise, which are found in different frequency bands and have different characteristic shapes.

In order to sort data, machine learning algorithms can use one of two approaches: classification or clustering. Classification algorithms search the data and sort the data into pre-defined categories. Clustering algorithms look for relationships within the data to create categories, called clusters, into which the data is then sorted. Classification algorithms are part of supervised learning since the computer determines the structure of the data from data that is already provided. Clustering algorithms are part of unsupervised learning since the computer determines the structure of the data without any previous information. Clustering algorithms can be used to characterize the noise by identifying common characteristics within the noise and therefore clustering algorithms can help improve categories of classification. [1]

Neural networks can be used to find relationships between the input data using layers of connections within the data. Neural networks consist of components called artificial neurons, which activate based on whether inputs into the unit meet a certain threshold. [1]

The aim of this project was to characterize different sources of noise in LIGO data using machine learning algorithms. First we tested clustering algorithms on seismic data, and then implemented a neural network to characterize the seismic noise data.

3 Clustering Algorithms Used

3.1 K-means Clustering

The k-means clustering algorithm creates clusters by separating data points into k number of groups. The value of k is input into the algorithm. The clusters are determined by minimizing the inertia, or the within-cluster sum-of-squares of the distances of each point from the mean. The inertia is a measure of how coherent the clusters are. By minimizing the inertia, the algorithm tries to minimize the difference between the mean value of a cluster and the values of points in the cluster. If a set of n samples x are inputted, the algorithm divides the samples into k clusters that are referred to as C. Each cluster is described by its

mean u_j , or centroid. The inertia of a cluster is calculated by the following expression:

$$\sum_{i=0}^n \min_{\mu_j \in C} (\|x_j - \mu_i\|^2)$$

The inertia is not normalized, but lower values are better and zero is the optimum value. The inertia assumes that the clusters are convex and isotropic, and would not work well to cluster irregular or elongated clusters. [2][3]

3.2 DBSCAN Clustering

The DBSCAN clustering algorithm creates clusters out of areas in the data of higher density. Unlike kmeans, it does not consider clusters to have any particular shapes, and the algorithm determines the number of clusters based on input parameters. Core samples are points that are in areas of high densities. The algorithm creates clusters around core samples so that the clusters consist of core samples, and non-core samples that are close to the core samples. The core samples are determined by two input parameters, the minimum samples and a specified distance, ε . A point is in the ε -neighborhood if the distance d from a point p to a point q is within a radius of ε . High density areas have the minimum sample of values within the ε -neighborhood. By increasing the number of minimum samples, and decreasing the distance, ε , a cluster's density is increased. [2][4]

3.3 Agglomerative Clustering

Agglomerative clustering is a type of hierarchical clustering algorithm. Hierarchical clustering builds clusters by merging and splitting clusters many times. Agglomerative clustering works by initially giving each data point its own cluster and then merging the clusters until the input number of clusters is reached. [2]

3.4 Birch Clustering

Birch clustering stands for balanced iterative reducing and clustering hierarchies. It is a hierarchical clustering algorithm which builds clusterings by merging and splitting clusters many times. [5]

3.5 Evaluating Clustering Algorithms

3.5.1 Calinsky Harabaz Index

The Calinsky-Harabaz index is a method used to evaluate clustering algorithm performance, that does not require input of external data. The Calinsky-Harabaz score is calculated by finding the ratio of the between-clusters dispersion mean to the within-cluster dispersion mean. This ratio is calculated as follows:

$$s(k) = \frac{Tr(B_k)}{Tr(W_k)} \times \frac{N - k}{k - 1}$$

Where k is the number of clusters, B_k is the between group dispersion matrix, W_k is the within group dispersion matrix and N is the number of data points. W_k and B_k are defined by:

$$W_k = \sum_{q=1}^k \sum_{x \in C_q} (x - c_q)(x - c_q)^T$$

$$B_k = \sum_q n_q (c_q - c)(c_q - c)^T$$

Where C_q is the number of set points in cluster q , c_q is the center of cluster q , c is the center of the clusters, and n_q is the number of points in cluster q . [2]

3.5.2 Earthquake Comparison Score

Another way to evaluate how well the clustering algorithms work is to compare the cluster labels to times of recorded earthquakes. This is done by adding up the cluster labels that occur five minutes before and after an earthquake Rayleigh wave arrives at the site and by adding the total amount of cluster labels and for each individual cluster. The number of cluster labels near an earthquake is divided by the total number of cluster labels. For each cluster k , the earthquake comparison score, $E(k)$ can be determined by:

$$E(k) = \frac{N_e}{N_t}$$

Where N_e is the number of cluster labels five minutes before and after an earthquake, and N_t is the total number of cluster labels. If a cluster corresponds to the presence of an earthquake then it will have a high percentage of its cluster labels present near an earthquake.

4 Clustering Results

I compared how well the clusters determined by clustering algorithms correspond to earthquakes that show up as peaks in the data. I read in seismic data taken from three seismometers. I read in the earthquake band channels (0.03 - 0.1 Hz) from the data and then clustered the channels using kmeans, and dbscan. The script counts the cluster labels five minutes before and after the time when earthquake Rayleigh waves arrive at the site as well as the total number of cluster labels. For each individual cluster, the earthquake comparison score is calculated. Only earthquakes with ground displacement greater than 65

percent of the recorded ground displacements are considered. This score is used to determine if cluster corresponds to an earthquake.

I have used seismic data from the Hanford observatory from March 2017. I have clustered the data from the earthquake channels using kmeans, and dbscan. I've also used the Calinsky-Harabaz index to evaluate how well the clustering works. Tables 1 and 2 show the clustering results for the kmeans and dbscan algorithms respectively.

Number of Clusters	Calinsky-Harabaz Score	Cluster of Earthquake Score	Earthquake Score
2	40192	1	0.5
3	37288	1	0.48
4	43960	2	0.31
5	44225	2	0.34
6	45618	2	0.33
7	46338	2	0.33
8	46349	1	0.44
9	46190	3	0.59
10	45323	6	0.75
Average	43943	N/A	0.45

Table 1: Results of kmeans clustering

Epsilon Value	Minimum Samples	Number of Clusters	Calinsky-Harabaz Score	Cluster of Earthquake Score	Earthquake Score
1	15	1	14	-1	0.01
2	10	15	5	-1	0.01
2	15	5	6	-1	0.01
2	20	1	14	-1	0.01
2	25	1	14	-1	0.01
2	30	1	14	-1	0.01
3	15	6	123	-1	0.01
4	15	8	194	-1	0.01

Table 2: Results of DBSCAN clustering

The dbscan clustering algorithm does not seem to work well because it places the majority of its points in the noise cluster (indicated as -1) which results in most of the points near earthquakes being classified as not-earthquakes.

A plot of the data clustered by kmeans is shown in Figure 2 and Figure 3. The lines that indicate the time that the Rayleigh waves from the earthquake reach the sight do not align with the peaks of the earthquakes on the plot. As a result, a new criteria was defined to evaluate how well the clusters align with earthquake predictions. The peaks in the data are taken to be earthquakes and the peak earthquake score counts the cluster labels five minutes before and after the center of the ppeaks. This score is used to determine how well

a cluster corresponds to an earthquake as determined from peaks in the data. Tables 3, 4, 5, and 6 show the clustering results for the kmeans, dbscan, agglomerative clustering, and birch algorithms, respectively. A plot of the data clustered by kmeans, and compared with peaks in the data, is shown in Figure 4 and Figure 5.

Number of Clusters	Calinsky-Harabaz Score	Cluster of Earthquake Score	Earthquake Score
2	40172	1	0.03
3	37282	1	0.04
4	43960	1	0.07
5	44225	4	0.08
6	45616	3	0.08
7	46338	3	0.08
8	46349	7	0.11
9	46095	1	0.11
10	46747	6	0.13
Average	46747	N/A	0.08

Table 3: Results of kmeans clustering (Earthquake score determined by peaks)

Epsilon Value	Minimum Samples	Number of Clusters	Calinsky-Harabaz Score	Cluster of Earthquake Score	Earthquake Score
1	15	1	14	-1	0.0125
2	10	15	5	-1	0.0126
2	15	5	6	-1	0.0125
2	20	1	14	-1	0.0125
2	25	1	14	-1	0.0125
2	30	1	14	-1	0.0125
3	15	6	123	-1	0.0141
4	15	6	194	-1	0.0159
5	15	8	373	-1	0.0176

Table 4: Results of DBSCAN clustering (Earthquake score determined by peaks)

Based on the earthquake score based on peaks, the kmeans clustering algorithm does not appear to agree very well with results. The agglomerative clustering and birch clustering algorithms give the same results, which are not very different from the kmeans clustering results. None of the clustering algorithms have clusters that clearly identify earthquakes.

In order to obtain clusters that better correspond to earthquakes, I added rows to the data that are shifted by time and inputted this timeshifted data into the clustering algorithms. Table 7 shows the average comparison of different timeshifts for the kmeans clustering algorithm.

Shifting the data and applying the kmeans clustering algorithm does not improve the performance of the kmeans clustering algorithm.

Number of Clusters	Calinsky-Harabaz Score	Cluster of Earthquake Score	Earthquake Score
2	37801	1	0.04
3	33282	0	0.04
4	40010	1	0.08
5	40007	1	0.08
6	40339	1	0.08
7	43396	0	0.08
8	42621	3	0.13
9	42045	8	0.13
10	41586	8	0.13
Average	46747	N/A	0.08

Table 5: Results of agglomerative clustering (Earthquake score determined by peaks)

Number of Clusters	Calinsky-Harabaz Score	Cluster of Earthquake Score	Earthquake Score
2	37801	1	0.04
3	33282	0	0.04
4	40010	1	0.08
5	40007	1	0.08
6	40339	1	0.08
7	43396	0	0.08
8	42621	3	0.13
9	42045	8	0.13
10	41586	8	0.13
Average	46747	N/A	0.08

Table 6: Results of birch clustering (Earthquake score determined by peaks)

Timeshift (minutes)	Calinsky-Harabaz Average	Maximum Earthquake Score Average
0	44087	0.08
10	49251	0.08
30	44081	0.09
60	44066	0.08

Table 7: Results of Timeshifted Kmeans Clustering (Earthquake score determined by peaks)

5 Neural Network Results

The neural network was implemented using the keras package with tensorflow as a backend. Earthquake channel seismic BLRMS data from Hanford during March 2017 was timeshifted by 30 minutes and then read into the neural network. For each point in the data it was indicated whether or not it corresponds to an earthquake. The peaks in the

data were used to determine which points correspond to earthquakes. Points that occur 5 minutes before and after a peak in the data were determined to be earthquake points.

Neural networks can be used to find relationships between the input data using layers of connections within the data. Neural networks consist of components called artificial neurons, which activate based on whether inputs into the unit meet a certain threshold. [1] A diagram of a neural network is shown in 1.

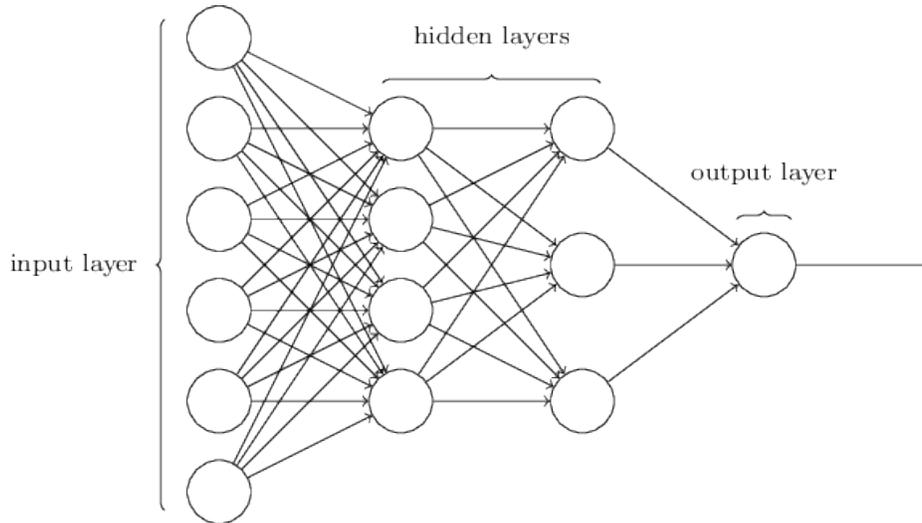


Figure 1: Diagram of Neural Network (from <http://neuralnetworksanddeeplearning.com/chap1.html>)

The neural network consists of a sequential model with three dense layers. The sequential model consists of a linear stack of layers. The first layer has an output equal to the number of channels in the input data. The second layer has an output of 9 and the last layer has an output of one. The last output indicates whether each point is an earthquake. The first two layers use elu activation and the last layer uses softmax activation. Each layer is followed by a dropout statement which helps prevent overfitting. The network was compiled using binary crossentropy as the loss.

An accuracy of 0.998 is obtained when using part of the data set to validate the trained neural network. This value indicates 99 percent of the validating data was identified correctly. A plot of the neural network is shown in figures 6 and 7. The vertical lines correspond to peaks in data that are caused by earthquakes. The dotted lines indicate where the neural network detects an earthquake. The neural network does a better job of detecting earthquakes than the clustering algorithms based on comparing how well the neural network detects earthquakes to how the clustering algorithm detects earthquakes. However, more data is needed in order to test the neural network, and to improve the accuracy of the neural network.

6 Acknowledgments

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References

- [1] Aurelien Geron, *Hands-On Machine Learning with Scikit-Learn and TensorFlow*. O'Reilly Media Inc., (2017).
- [2] <http://scikit-learn.org/stable/modules/clustering.html>
- [3] David Arthur, Sergei Vassilvitskii, *k-means++: The Advantages of Careful Seeding*. Proceedings of the Eighteenth Annual ACM-SIAM Symposium on Discrete Algorithms, Society of Industrial and Applied Mathematics, (2007).
- [4] Martin Ester, Hans-Peter Kriegel, Jorg Sander, Xiaowei Xu, *A Density-Based Algorithm for Discovering Clusters in Large Spatial Databases with Noise*. Proceedings of the 2nd International Conference on Knowledge Discovery and Data Mining, (1996).
- [5] Tian Zhang, Raghu Ramakrishnan, Miron Liviny *BIRCH: An Efficient Data Clustering Method for Very Large Databases* Proceedings of the 1996 ACM SIGMOD international conference on Management of data, (1996).

A Script for Evaluating Clustering Algorithms

```
'''
```

```
This script reads in seismic noise data from March 2017 and earthquake data.
```

```
It shifts the data by time for clustering
```

```
It determined earthquake times by looking at peaks in data
```

```
It clusters earthquake channels using kmeans and dbscan.
```

```
It compares the clusters around the earthquake times to deterime effectiveness of clus
```

```
It plots the data as clustered by kmeans and dbscan
```

```
'''
```

```
from __future__ import division
from sklearn.cluster import KMeans
from sklearn.cluster import DBSCAN
from sklearn.cluster import AffinityPropagation
from sklearn.cluster import MeanShift, estimate_bandwidth
from sklearn.cluster import spectral_clustering
from sklearn.cluster import AgglomerativeClustering
from sklearn.cluster import Birch
from sklearn import metrics
from sklearn.preprocessing import StandardScaler
```

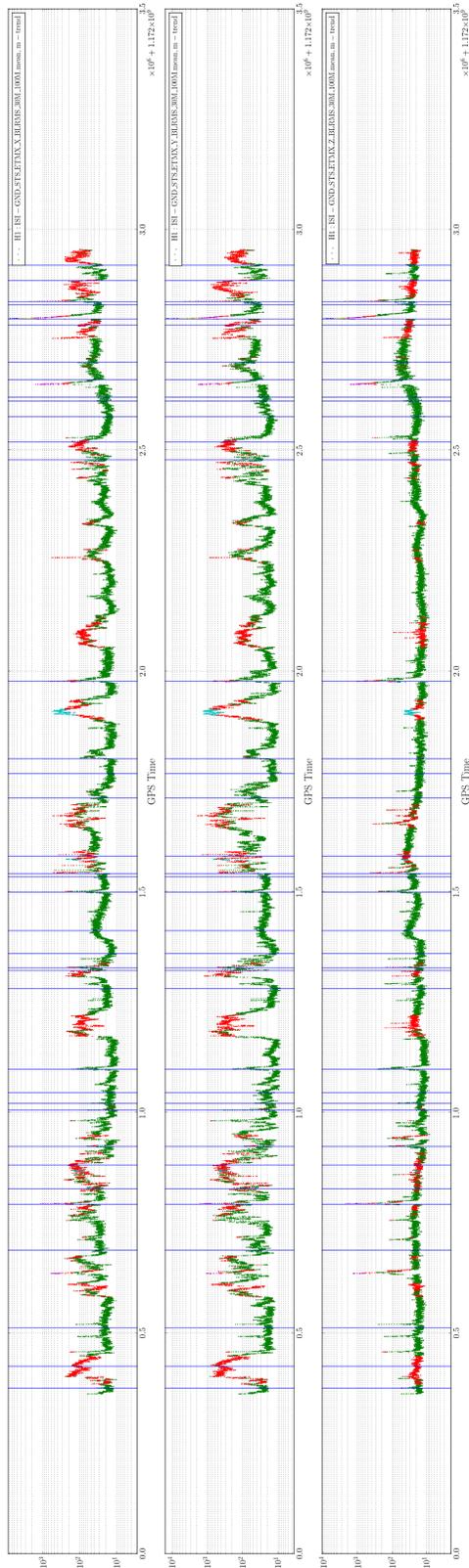


Figure 2: Plot of data from earthquake channels clustered using kmeans with $k=6$ (earthquakes indicated)

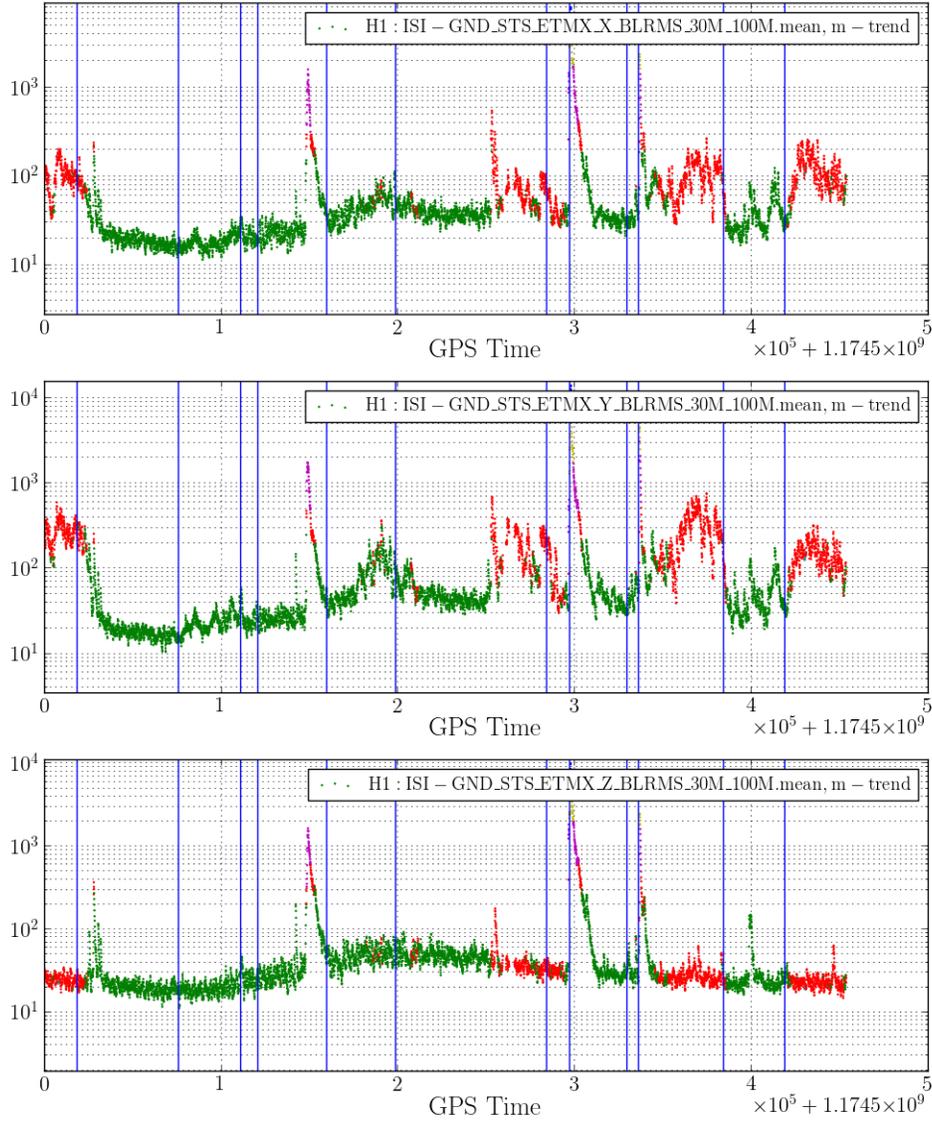


Figure 3: Figure 1 zoomed in for detail

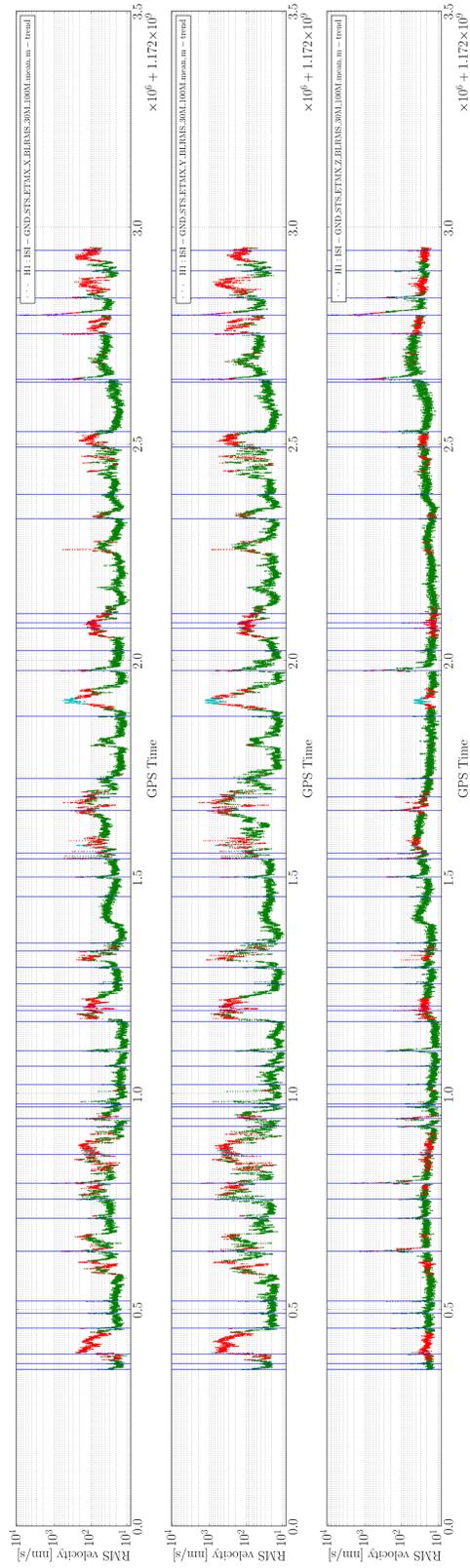


Figure 4: Plot of data from earthquake channels clustered using kmeans with $k=6$ (peaks indicated)

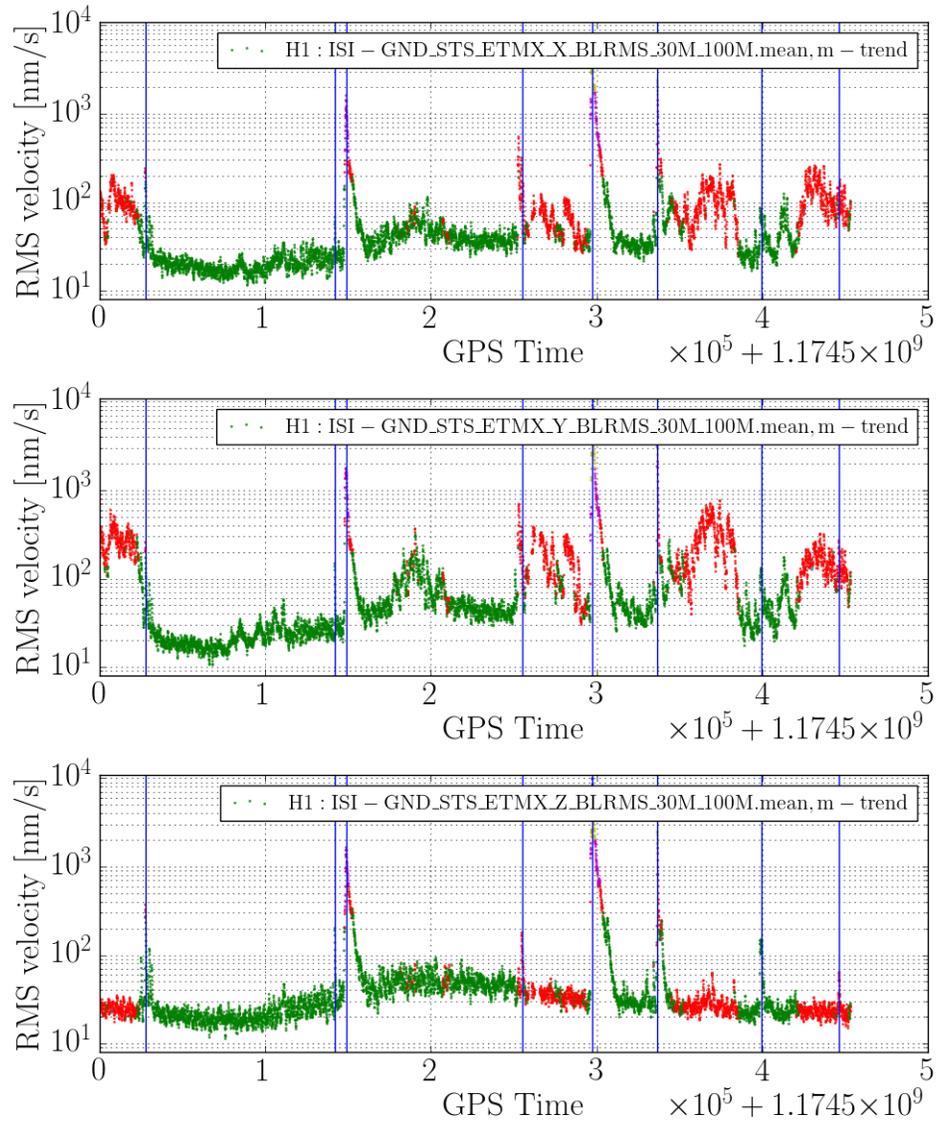


Figure 5: Figure 3 zoomed in for detail

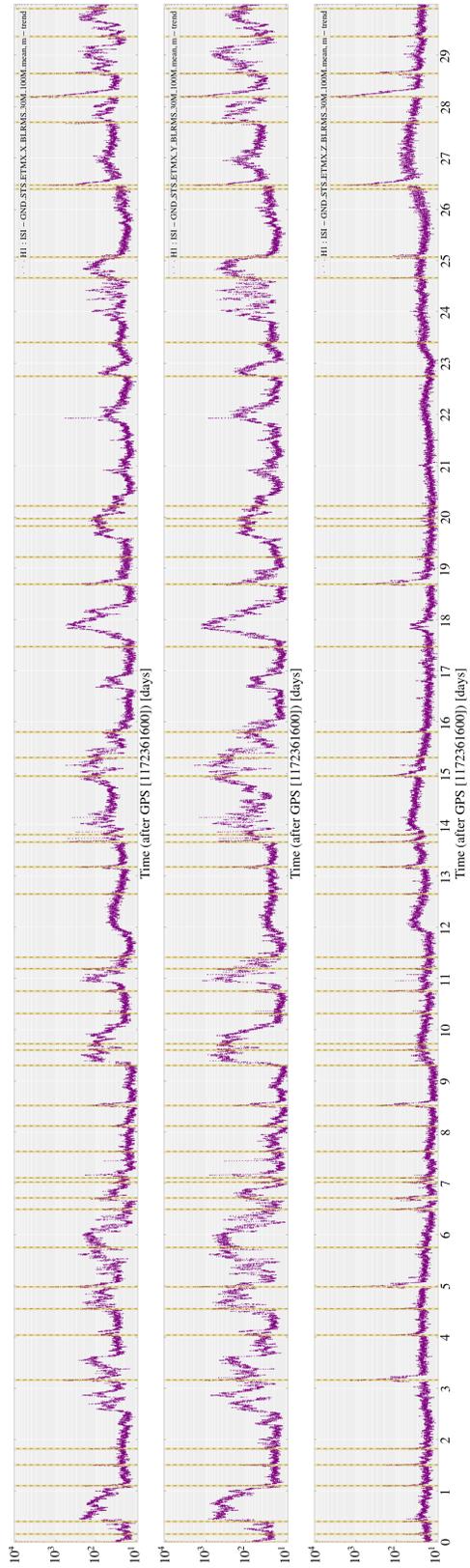


Figure 6: Plot of data from earthquake channels classified by neural net (peaks indicated)

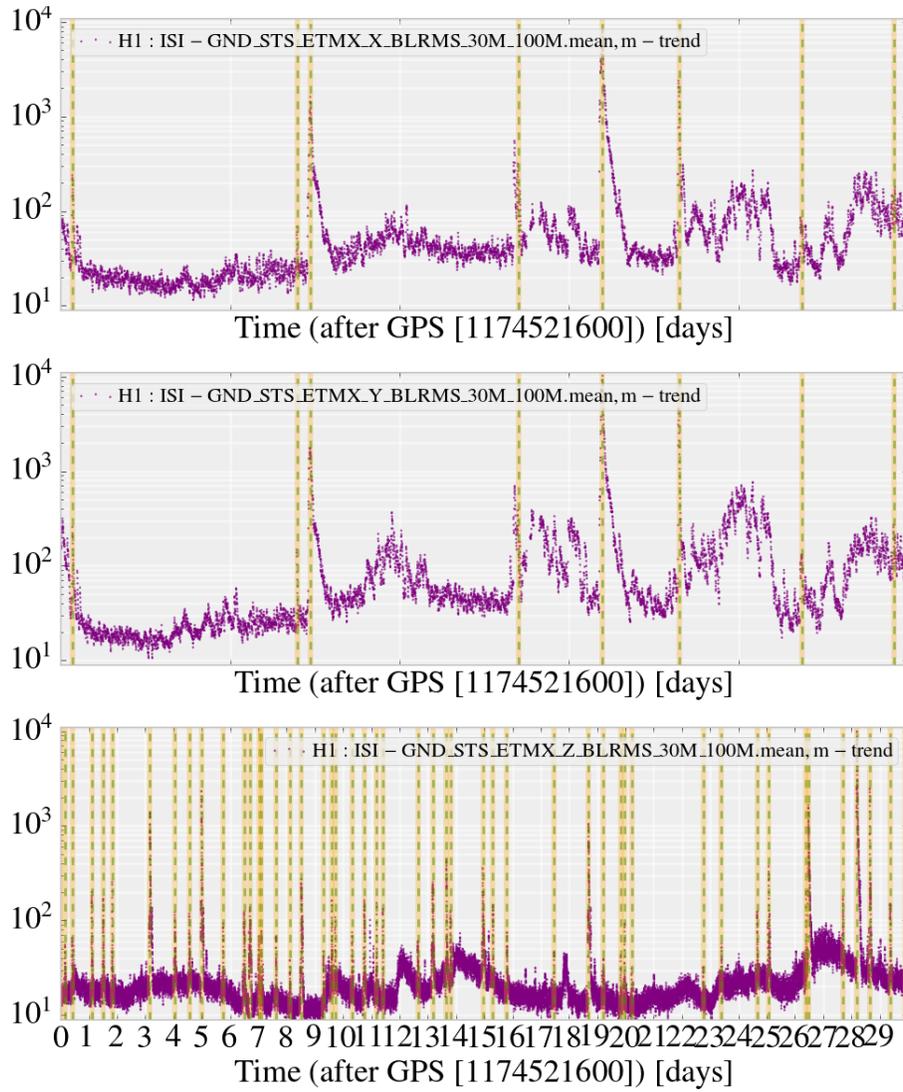


Figure 7: Figure 6 zoomed in for detail

```

import numpy as np
from scipy.io import loadmat
import matplotlib
matplotlib.use('Agg')
import matplotlib.pyplot as plt
from matplotlib.pyplot import cm
import scipy.signal as sig
from astropy.time import Time
import collections

plt.rc('text', usetex = True)
plt.rc('font', **{'family': 'serif', 'serif': ['Computer Modern']})
plt.rc('axes', labelsz = 20.0)
plt.rc('axes', axisbelow = True)
plt.rc('axes.formatter', limits=[-3,4])
plt.rc('legend', fontsize = 14.0)
plt.rc('xtick', labelsz = 16.0)
plt.rc('ytick', labelsz = 16.0)
plt.rc('figure', dpi = 100)

# colors for clusters
colors = np.array(['r', 'g', 'b', 'y', 'c', 'm', 'darkgreen', 'plum',
                  'darkblue', 'pink', 'orangered', 'indigo'])

cl          = 6    # number of clusters for kmeans
eps         = 2    # min distance for density for DBscan
min_samples = 15  # min samples for DBscan

#read in data
H1dat = loadmat('Data/' + 'H1_SeismicBLRMS.mat')
#edat  = np.loadtxt('Data/H1_earthquakes.txt')

# read in earthquake channels
cols   = [6,12,18,24,30,36,42,48]      # NEED comment here
vdat   = np.array(H1dat['data'][0])
vchans = np.array(H1dat['chans'][0])
for i in cols:
    add = np.array(H1dat['data'][i])
    vdat = np.vstack((vdat, add))
for i in cols:
    vchans = np.append(vchans, H1dat['chans'][i])
timetuples = vdat.T

# shift the data
vdat2   = vdat
vchans2 = vchans
num     = 10

```

```

t_shift = 10 # how many minutes to shift the data by
for i in cols:
    add = np.array(H1dat['data'][i])
    for j in range(1, t_shift+1):
        add_shift = add[j:]
        add_values = np.zeros((j,1))
        add_shift = np.append(add_shift, add_values)
        vdat2 = np.vstack((vdat2, add_shift))
        chan = 'Time_Shift_' + str(j) + '_Min_EQ_Band_' + str(i)
        vchans2 = np.append(vchans2, chan)
print(np.shape(vdat2))
vdat2 = vdat[:, :43200-t_shift]
print(np.shape(vdat2))
timetuples2 = vdat.T
timetuples3 = vdat[0:num].T

#convert time to gps time
times      = '2017-03-01 00:00:00'
ti         = Time(times,format='iso',scale='utc')
t_start    = int(np.floor(ti.gps/60)*60)
dur_in_days = 30
dur_in_minutes = dur_in_days*24*60
dur         = dur_in_minutes*60
t_end      = t_start + dur
t          = np.arange(t_start, t_end, 60)

# create list of earthquake times from peaks
# find peaks in all three z channel directions
widths     = np.arange(5, 140) # range of widths in minutes
min_snr    = 5
noise_perc = 15
peaks1     = sig.find_peaks_cwt(vdat[2], widths,
                               min_snr = min_snr, noise_perc=noise_perc)
peaks2     = sig.find_peaks_cwt(vdat[5], widths,
                               min_snr = min_snr, noise_perc=noise_perc)
peaks3     = sig.find_peaks_cwt(vdat[8], widths,
                               min_snr = min_snr, noise_perc=noise_perc)

# takes average time for earthquake times from three channels
# that are within dtau minutes of each other
dtau = 3
peak_list = np.array([])
for i in peaks1:
    for j in peaks2:
        for k in peaks3:
            if (abs(i-j) <= dtau and abs(i-k) <= dtau):

```

```

        avg = (i+j+k)/3
        peak_list = np.append(peak_list, avg)
EQ_times = np.array([])
for i in peak_list:
    EQ_times = np.append(EQ_times, t[int(i)])

# kmeans clustering loop

Nmin = 2
num = 9
Nmax = Nmin + num
for cl in range(Nmin, Nmax):
    kmeans = KMeans(n_clusters=cl, random_state=13).fit(timetuples2)
    kpoints = np.array([])
    xvals = np.arange(t_start, t_end, 60)
    for t in EQ_times: #for each EQ: collect indices within 30 min of EQ
        tmin = int(t - 10)
        tmax = int(t + 10)
        for j in range(tmin, tmax):
            val = abs(xvals - j)
            aval = np.argmin(val)
            kpoints = np.append(kpoints, aval)
    kpoints = np.unique(kpoints) # make sure there are no repeating indices
    kclusters = np.array([])
    for i in kpoints:
        #for each index find the corresponding cluster and store them in array
        kclusters = np.append(kclusters, kmeans.labels_[int(i)])
        # kmeans score determined by ratio of points in
        # cluster/points near EQ to points in cluster/all points
    k_count = collections.Counter(kclusters).most_common()
    ktot_count = collections.Counter(kmeans.labels_).most_common()
    k_list_cl = [x[0] for x in k_count] #cluster number
    k_list = [x[1] for x in k_count] #occurences of cluster
    ktot_list_cl = [x[0] for x in ktot_count]
    ktot_list = [x[1] for x in ktot_count]
    k_clusters = np.array([])
    k_compare = np.array([])
    k_list2 = np.array([])
    ktot_list2 = np.array([])
    # arrange so that k_clusters k_list2 and k_compare are in the same order
    for i in range(len(k_list_cl)):
        for j in range(len(ktot_list_cl)):
            if k_list_cl[i] == ktot_list_cl[j]:
                k_clusters = np.append(k_clusters, k_list_cl[i])
                compare = k_list[i]/ktot_list[j]
                k_compare = np.append(k_compare, compare)

```

```

        k_list2    = np.append(k_list2, k_list[i])
        ktot_list2 = np.append(ktot_list2, k_list[i])
np.set_printoptions(precision=3)
#print(k_clusters)
#print(k_compare)
max_val = max(k_compare)
max_index = np.argmax(k_compare)
max_cluster = int(k_clusters[max_index])
k_cal_score = metrics.calinski_harabaz_score(timetuples, kmeans.labels_)
#print('K-means ' + str(cl) + ': C-H score = {:.6g}'.format(k_cal_score))
print(str(cl) + ' & {:.6g}'.format(k_cal_score) + ' & ' + str(max_cluster) + ' & {:.6g}'.format(k_cal_score))
print('\n\hline')

# dbscan clustering loop

min_samples_list = [10,20,25,30]
eps_list = [1,2,3,4,5]
#for min_samples in min_samples_list:
for eps in eps_list:
    db = DBSCAN(eps=eps,min_samples=min_samples).fit(timetuples)
    #number of clusters
    n_clusters = len(set(db.labels_)) - (1 if -1 in db.labels_ else 0)
    #add up number of clusters that appear next to each earthquake
    xvals = np.arange(t_start,t_end,60)
    dbpoints = np.array([])
    for t in EQ_times: #for each EQ: collect indices within 5 min of EQ
        tmin = int(t-5*60)
        tmax = int(t+5*60)
        for j in range(tmin,tmax):
            val = abs(xvals-j)
            aval = np.argmin(val)
            dbpoints = np.append(dbpoints, aval)
    dbpoints = np.unique(dbpoints)
    dbclusters = np.array([])
    for i in dbpoints: dbclusters = np.append(dbclusters,db.labels_[int(i)]) #for each EQ
    #dbscan score determined by percent of points sorted into one cluster near EQ
    db_count = collections.Counter(dbclusters).most_common()
    dbtot_count = collections.Counter(db.labels_).most_common()
    db_list_cl = [x[0] for x in db_count]
    db_list = [x[1] for x in db_count]
    dbtot_list_cl = [x[0] for x in dbtot_count]
    dbtot_list = [x[1] for x in dbtot_count]
    db_clusters = np.array([])
    db_compare = np.array([])
    db_list2 = np.array([])

```

```

dbtot_list2 = np.array([])
for i in range(len(db_list_cl)):
    for j in range(len(dbtot_list_cl)):
        if db_list_cl[i] == dbtot_list_cl[j]:
            db_clusters = np.append(db_clusters, db_list_cl[i])
            compare = db_list[i]/dbtot_list[j]
            db_compare = np.append(db_compare, compare)
            db_list2 = np.append(db_list2, db_list[i])
            dbtot_list2 = np.append(dbtot_list2, db_list[i])
    #print(db_clusters)
    #print(db_compare)
max_val = max(db_compare)
max_index = np.argmax(db_compare)
max_cluster = int(db_clusters[max_index])
db_cal_score = metrics.calinski_harabaz_score(timetuples, db.labels_)
print(str(eps) + ' & ' + str(min_samples) + ' & ' + str(n_clusters) + ' & {:.6g}',
print('\hline')

```

#ag clustering loop

```

Nmin = 2
num = 9
Nmax = Nmin + num
for cl in range(Nmin, Nmax):
    ag = AgglomerativeClustering(n_clusters=cl).fit(timetuples)
    agpoints = np.array([])
    xvals = np.arange(t_start, t_end, 60)
    for t in EQ_times: #for each EQ: collect indices within 30 min of EQ
        tmin = int(t - 10)
        tmax = int(t + 10)
        for j in range(tmin, tmax):
            val = abs(xvals - j)
            aval = np.argmin(val)
            agpoints = np.append(agpoints, aval)
    agpoints = np.unique(agpoints) # make sure there are no repeating indices
    agclusters = np.array([])
    for i in agpoints:
        #for each index find the corresponding cluster and store them in array
        agclusters = np.append(agclusters, ag.labels_[int(i)])
    ag_count = collections.Counter(agclusters).most_common()
    agtot_count = collections.Counter(ag.labels_).most_common()
    ag_list_cl = [x[0] for x in ag_count] #cluster number
    ag_list = [x[1] for x in ag_count] #occurences of cluster
    agtot_list_cl = [x[0] for x in agtot_count]
    agtot_list = [x[1] for x in agtot_count]

```

```

ag_clusters = np.array([])
ag_compare  = np.array([])
ag_list2    = np.array([])
agtot_list2 = np.array([])
# arrange so that k_clusters k_list2 and k_compare are in the same order
for i in range(len(ag_list_cl)):
    for j in range(len(agtot_list_cl)):
        if ag_list_cl[i] == agtot_list_cl[j]:
            ag_clusters = np.append(ag_clusters, ag_list_cl[i])
            compare     = ag_list[i]/agtot_list[j]
            ag_compare  = np.append(ag_compare, compare)
            ag_list2    = np.append(ag_list2, ag_list[i])
            agtot_list2 = np.append(agtot_list2, ag_list[i])
np.set_printoptions(precision=3)
max_val = max(ag_compare)
max_index = np.argmax(ag_compare)
max_cluster = int(ag_clusters[max_index])
ag_cal_score = metrics.calinski_harabaz_score(timetuples, ag_labels_)
print(str(cl) + ' & {:.6g}'.format(ag_cal_score) + ' & ' + str(max_cluster) + ' & {')
print('\nhline')

```

#birch clustering loop

```

Nmin = 2
num = 9
Nmax = Nmin + num
for cl in range(Nmin, Nmax):
    birch = Birch(n_clusters=cl).fit(timetuples)
    bpoints = np.array([])
    xvals = np.arange(t_start, t_end, 60)
    for t in EQ_times: #for each EQ: collect indices within 30 min of EQ
        tmin = int(t - 10)
        tmax = int(t + 10)
        for j in range(tmin, tmax):
            val = abs(xvals - j)
            aval = np.argmin(val)
            bpoints = np.append(bpoints, aval)
    bpoints = np.unique(bpoints) # make sure there are no repeating indices
    bclusters = np.array([])
    for i in bpoints:
        #for each index find the corresponding cluster and store them in array
        bclusters = np.append(bclusters, birch.labels_[int(i)])
    b_count = collections.Counter(bclusters).most_common()
    btot_count = collections.Counter(birch.labels_).most_common()
    b_list_cl = [x[0] for x in b_count] #cluster number

```

```

b_list      = [x[1] for x in b_count] #occurrences of cluster
btot_list_cl = [x[0] for x in btot_count]
btot_list   = [x[1] for x in btot_count]
b_clusters  = np.array([])
b_compare   = np.array([])
b_list2     = np.array([])
btot_list2  = np.array([])
# arrange so that b_clusters b_list2 and k_compare are in the same order
for i in range(len(b_list_cl)):
    for j in range(len(btot_list_cl)):
        if b_list_cl[i] == btot_list_cl[j]:
            b_clusters = np.append(b_clusters, b_list_cl[i])
            compare     = b_list[i]/btot_list[j]
            b_compare   = np.append(b_compare, compare)
            b_list2     = np.append(b_list2, b_list[i])
            btot_list2  = np.append(btot_list2, btot_list[j])
np.set_printoptions(precision=3)
max_val = max(b_compare)
max_index = np.argmax(b_compare)
max_cluster = int(b_clusters[max_index])
b_cal_score = metrics.calinski_harabaz_score(timetuples, birch.labels_)
print(str(c1) + ' & {:.6g}'.format(b_cal_score) + ' & ' + str(max_cluster) + ' & {:.6g}'.format(max_index))
print('\n\\hline')

```

B Script for Neural Network

```

'''
Reads in data from mat file
Creates a neural network using keras
Plots data with prediction labels in a graph
'''

from keras.models import Sequential
from keras.layers import Dense, Dropout
from keras import optimizers
import numpy as np
import scipy.io as sio
from astropy.time import Time
import collections
import matplotlib
matplotlib.use('Agg')
import matplotlib.pyplot as plt
from matplotlib.pyplot import cm

np.random.seed(7)

```

```

# read in data
EQ_data = sio.loadmat('Data/EQ_info.mat')
vdat    = EQ_data['vdat']
vchans  = EQ_data['vchans']
EQ_times = EQ_data['EQ_times']
EQ_times = EQ_times.reshape(49,1)
X       = EQ_data['X']
points, size = np.shape(X)
Y       = EQ_data['EQ_labels']
Y       = Y.reshape(43200,)
t       = EQ_data['t']

# print info about data
print('size is ' + str(size))
print('Shape of X is ' + str(np.shape(X)))
print('Shape of Y is ' + str(np.shape(Y)))
print(np.shape(EQ_times))

# neural network
optimizer = optimizers.Adam(lr = 1e-5)
model = Sequential()
model.add(Dense(size, input_shape = (size,), activation = 'elu'))
model.add(Dropout(.1))
model.add(Dense(9, activation = 'elu'))
model.add(Dropout(.1))
model.add(Dense(1, activation = 'softmax'))

# model.output_shape
model.compile(loss = 'binary_crossentropy',
              optimizer = optimizer,
              metrics = ['accuracy'])
model.fit(X, Y,
          epochs = 10,
          batch_size = 256,
          validation_split=0.1,
          verbose = 1)

#score = model.evaluate(X,Y)
#print(score)

model.summary()

# prediction values
Y_pred = model.predict(X)
Y_pred2 = Y_pred.T

```

```

Y_pred3 = np.array([])
for i in Y_pred2:
    Y_pred3 = np.append(Y_pred3, i)
Y_pred3 = Y_pred3.astype(int)

# Plot of data points
colors = np.array(['r', 'b', 'm', 'g'])
labels = Y.T
labels = labels.astype(int)
num = size

fig, axes = plt.subplots(len(vdat[0:num]), figsize=(40, 4*len(vdat[0:num])))
for ax, data, chan in zip(axes, vdat[0:num], vchans):
    ax.scatter(t, data, c=colors[Y_pred3], edgecolor='',
              s=3, label=r'$\mathrm{\%s}$' % chan.replace('_', '\_'))
    ax.set_yscale('log')
    ax.set_ylim(np.median(data)*0.1, max(data)*1.1)
    ax.set_xlabel('GPS Time')
    ax.grid(True, which='both')
    ax.legend()
    for e in range(len(EQ_times)):
        ax.axvline(x = EQ_times[e], color = 'r')

fig.tight_layout()

print("Saving plot...")
fig.savefig('Figures/NeuralNetworkComparison3.pdf')

```