

□ HMS - Exploratory Data Analysis and Journey in Neurocritical Care

1. Understanding the EEG Domain

What is Electroencephalography (EEG)? Electroencephalography, commonly referred to as EEG, is a non-invasive method used to record electrical activity in the brain. This technique involves placing electrodes on the scalp, which detect tiny electrical charges that result from the activity of brain cells. The signals captured by these electrodes are amplified and recorded, typically resulting in a series of wavy lines that are analyzed by specialists.



How Electroencephalography Data Looks Like? Electroencephalography (EEG) data typically appears as a series of wavy lines, each representing the electrical activity recorded from different electrodes placed on the scalp. These lines, called traces, show the voltage changes over time. The patterns observed in EEG data are influenced by the brain activity of the individual, and they can vary significantly depending on the state of consciousness, activity, or any neurological conditions.

What Patterns May Be Observed in the EEG Data?

- **Wave Patterns:** EEG data is characterized by different types of wave patterns, such as alpha, beta, delta, and theta waves. Each type corresponds to different brain states. For example, alpha waves are often associated with a state of relaxation, while beta waves are linked with active thinking or concentration.
- **Amplitude and Frequency:** The waves have varying amplitudes (heights) and frequencies (speeds). The amplitude indicates the strength of the signal, and frequency shows how fast the brain waves are oscillating.
- **Artifacts:** These are non-brain waveforms that can appear in the data due to muscle movements, eye blinks, or electrical interference.
- **Abnormal Patterns:** In cases of neurological disorders like epilepsy, the EEG may show spikes, sharp waves, or other unusual patterns that indicate abnormal brain activity.

How Spectrograms are Related to EEG?

- **Frequency Analysis:** EEG signals consist of brain waves with different frequencies, like alpha, beta, theta, and delta waves. A spectrogram can visually display these frequencies, showing how they change over time during the EEG recording.
- **Identifying Patterns:** Spectrograms can help in identifying patterns that might not be easily discernible in the raw EEG waveforms. For example, they can be used to detect changes in brain activity during different sleep stages or to identify oscillatory activity associated with certain neurological disorders, like epilepsy.
- **Temporal and Frequency Resolution:** A key advantage of spectrograms in EEG analysis is their ability to provide information about both the timing (temporal resolution) and the frequency (frequency resolution) of brain waves. This is crucial for understanding dynamic changes in brain activity.
- **Data Visualization:** Spectrograms offer a more intuitive way to visualize and interpret complex EEG data. They can transform the EEG's time-domain data into a more accessible frequency-domain representation, which can be easier to analyze and understand, especially in research and clinical diagnostics.

What is LPD/GPD/LRDA/GRDA Conditions and How They May Be Seen in Data?

Lateralized Periodic Discharges (LPDs)

- **Lateralization:** LPDs are lateralized, meaning they predominantly occur on one side (hemisphere) of the brain.
- **Periodicity:** LPDs are periodic, meaning they display a repeating pattern. This periodicity is a key feature in their identification on an EEG.
- **Waveform Characteristics:** LPDs typically consist of sharp waveforms or complexes that are clearly distinguishable from the background EEG activity. These sharp waves are usually followed by a slow-wave component.
- **General Characteristics:** In EEG data, LPDs appear as regular, sharply contoured waveforms that stand out against the background brain activity and repeat at consistent intervals. They are usually unilateral, affecting either the left or right hemisphere, which is an important aspect in their interpretation.

Generalized Periodic Discharges (GPDs)

- **Generalized Distribution:** GPDs are characterized by their distribution across both hemispheres of the brain, rather than being localized to one side.
- **Periodicity:** Like LPDs, GPDs exhibit a repeating pattern. Their periodic nature is crucial for their identification on EEG and distinguishes them from other generalized EEG abnormalities.
- **Waveform Characteristics:** GPDs usually consist of repetitive, sharply contoured waveforms that can vary in shape and duration. They are often more synchronized compared to other EEG patterns.
- **Clinical Implications:** GPDs may be seen in various clinical scenarios, including severe diffuse brain injury, hypoxic-ischemic encephalopathy, and in association with certain drug toxicities or metabolic derangements. Their presence can indicate a severe underlying brain dysfunction.

Lateralized Rhythmic Delta Activity (LRDA)

- **Lateralization:** The key feature of LRDA is its lateralized nature, affecting predominantly one hemisphere of the brain, which can be crucial in localizing a neurological lesion or dysfunction.
- **Rhythmic Delta Waves:** Unlike the sharp waveforms of LPDs, LRDA is defined by smoother, more rhythmic waveforms, predominantly in the delta frequency range (1-4 Hz).
- **Clinical Context:** LRDA is often observed in patients with focal brain lesions, such as those caused by stroke, tumors, or inflammation. It may also be seen in the context of focal seizure activity.
- **Interpretation:** The presence of LRDA in an EEG reading can provide valuable information regarding the location and possibly the nature of brain pathology, aiding in diagnosis and treatment planning.

Generalized Rhythmic Delta Activity (GRDA)

- **Generalized Distribution:** GRDA differs from LRDA in that it is not lateralized but involves both hemispheres, often symmetrically.
- **Rhythmic Delta Waves:** This pattern is defined by continuous or quasi-continuous rhythmic delta waves. The activity is slower and more rhythmic compared to GPDs.
- **Associated Conditions:** GRDA can be seen in various clinical conditions, including encephalopathies of different etiologies (like toxic-metabolic disturbances), and in some cases, during certain sleep stages or in diffuse brain disorders.
- **Diagnostic Significance:** The presence of GRDA can be indicative of a global brain dysfunction and may warrant further investigation to identify its cause. It is particularly significant in assessing patients with altered levels of consciousness or diffuse neurological impairments.

Disclaimer: I'm not an expert in this domain. For a more detailed explanation from the competition host, read this [paper](#). If you have questions or you think I'm wrong somewhere, feel free to leave a comment. And don't forget to upvote 😊.

```
In [ ]: # Required Libraries
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [ ]: # File Paths
train_path = '/kaggle/input/hms-harmful-brain-activity-classification/train.csv'
test_path = '/kaggle/input/hms-harmful-brain-activity-classification/test.csv'
sample_submission_path = '/kaggle/input/hms-harmful-brain-activity-classification/sample_submission.csv'
```

```
In [ ]: # Load Train, Test, and Sample Submission Data
train_data = pd.read_csv(train_path)
test_data = pd.read_csv(test_path)
sample_submission_data = pd.read_csv(sample_submission_path)
```

```
In [ ]: # Display General Information about Train Data
print("Train Data General Information:")
print(train_data.info())
```

```

Train Data General Information:
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 0 entries
Data columns (total 15 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   eeg_id                                0 non-null      object
1   eeg_sub_id                            0 non-null      object
2   eeg_label_offset_seconds              0 non-null      object
3   spectrogram_id                        0 non-null      object
4   spectrogram_sub_id                    0 non-null      object
5   spectrogram_label_offset_seconds      0 non-null      object
6   label_id                              0 non-null      object
7   patient_id                            0 non-null      object
8   expert_consensus                      0 non-null      object
9   seizure_vote                          0 non-null      object
10  lpd_vote                              0 non-null      object
11  gpd_vote                              0 non-null      object
12  lrda_vote                             0 non-null      object
13  grda_vote                             0 non-null      object
14  other_vote                            0 non-null      object
dtypes: object(15)
memory usage: 124.0+ bytes
None

```

```

In [ ]: # Display General Information about Test Data
print("\nTest Data General Information:")
print(test_data.info())

```

```

Test Data General Information:
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1 entries, 0 to 0
Data columns (total 3 columns):
#   Column          Non-Null Count  Dtype
---  -
0   spectrogram_id  1 non-null     int64
1   eeg_id          1 non-null     int64
2   patient_id      1 non-null     int64
dtypes: int64(3)
memory usage: 152.0 bytes
None

```

```

In [ ]: # Display General Information about Sample Submission Data
print("\nSample Submission Data General Information:")
print(sample_submission_data.info())

```

```

Sample Submission Data General Information:
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1 entries, 0 to 0
Data columns (total 7 columns):
#   Column          Non-Null Count  Dtype
---  -
0   eeg_id          1 non-null     int64
1   seizure_vote    1 non-null     float64
2   lpd_vote        1 non-null     float64
3   gpd_vote        1 non-null     float64
4   lrda_vote       1 non-null     float64
5   grda_vote       1 non-null     float64
6   other_vote      1 non-null     float64
dtypes: float64(6), int64(1)
memory usage: 184.0 bytes
None

```

```

In [ ]: # Display First Few Rows of Train Data
print("\nFirst Few Rows of Train Data:")
print(train_data.head())

```

```

First Few Rows of Train Data:
Empty DataFrame
Columns: [eeg_id, eeg_sub_id, eeg_label_offset_seconds, spectrogram_id, spectrogram_sub_id, spectrogram_label_offset_seconds, label_id, patient_id, expert_consensus, seizure_vote, lpd_vote, gpd_vote, lrda_vote, grda_vote, other_vote]
Index: []

```

```

In [ ]: # Display First Few Rows of Test Data
print("\nFirst Few Rows of Test Data:")
print(test_data.head())

```

```

First Few Rows of Test Data:
  spectrogram_id  eeg_id  patient_id
0          853520  3911565283        6885

```

```

In [ ]: # Display First Few Rows of Sample Submission Data
print("\nFirst Few Rows of Sample Submission Data:")
print(sample_submission_data.head())

```

First Few Rows of Sample Submission Data:

```
eeg_id seizure_vote lpd_vote gpd_vote lrda_vote grda_vote \
0 3911565283 0.166667 0.166667 0.166667 0.166667 0.166667

other_vote
0 0.166667
```

```
In [ ]: # Display General Statistics of Train Data
print("Train Data Statistics:")
print(train_data.describe())
```

Train Data Statistics:

```
eeg_id eeg_sub_id eeg_label_offset_seconds spectrogram_id \
count      0      0      0      0
unique      0      0      0      0
top        NaN    NaN    NaN    NaN
freq        NaN    NaN    NaN    NaN

spectrogram_sub_id spectrogram_label_offset_seconds label_id \
count      0      0      0
unique      0      0      0
top        NaN    NaN    NaN
freq        NaN    NaN    NaN

patient_id expert_consensus seizure_vote lpd_vote gpd_vote lrda_vote \
count      0      0      0      0      0      0
unique      0      0      0      0      0      0
top        NaN    NaN    NaN    NaN    NaN    NaN
freq        NaN    NaN    NaN    NaN    NaN    NaN

grda_vote other_vote
count      0      0
unique      0      0
top        NaN    NaN
freq        NaN    NaN
```

Explore train EEGs

```
In [ ]: import pyarrow.parquet as pq
```

```
In [ ]: # Path to the file you want to load
file_path = '/kaggle/input/hms-harmful-brain-activity-classification/train_eegs/1001369401.parquet'
```

```
In [ ]: # Load the Parquet file
eeg_data = pq.read_table(file_path).to_pandas()

# Explore the EEG data
print(eeg_data.head())
```

```
      Fp1      F3      C3      P3      F7      T3 \
0 -590.419983 -557.659973 -478.279999 -586.659973 -576.590027 -537.460022
1 -587.760010 -563.630005 -481.760010 -589.859985 -579.710022 -537.369995
2 -590.479980 -563.960022 -482.040009 -588.679993 -574.159973 -532.340027
3 -593.179993 -563.260010 -476.160004 -582.909973 -565.640015 -520.799988
4 -588.820007 -556.729980 -470.410004 -573.299988 -556.789978 -519.239990

      T5      O1      Fz      Cz      Pz      Fp2 \
0 -554.330017 -605.809998 -560.599976 -526.630005 -557.299988 -551.030029
1 -556.320007 -609.349976 -563.710022 -530.549988 -561.549988 -557.030029
2 -550.179993 -606.450012 -568.270020 -534.049988 -563.059998 -555.969971
3 -544.150024 -600.869995 -568.190002 -530.250000 -558.789978 -559.809998
4 -537.080017 -589.690002 -560.070007 -520.919983 -548.849976 -557.869995

      F4      C4      P4      F8      T4      T6 \
0 -498.549988 -460.649994 -548.750000 -546.299988 -525.330017 -593.190002
1 -507.019989 -466.730011 -555.469971 -548.190002 -534.900024 -604.090027
2 -511.809998 -473.109985 -560.419983 -559.169983 -538.059998 -606.919983
3 -516.119995 -468.730011 -554.599976 -563.510010 -520.880005 -599.190002
4 -520.229980 -455.420013 -540.690002 -557.200012 -514.859985 -586.950012

      O2      EKG
0 -491.980011 46.619999
1 -499.429993 15.770000
2 -501.010010 47.669998
3 -494.140015 105.309998
4 -482.350006 77.900002
```

```
In [ ]: # Plot the EEG data for the first 1000 samples
num_samples_to_plot = 1000

fig, ax = plt.subplots(figsize=(12, 8))
```

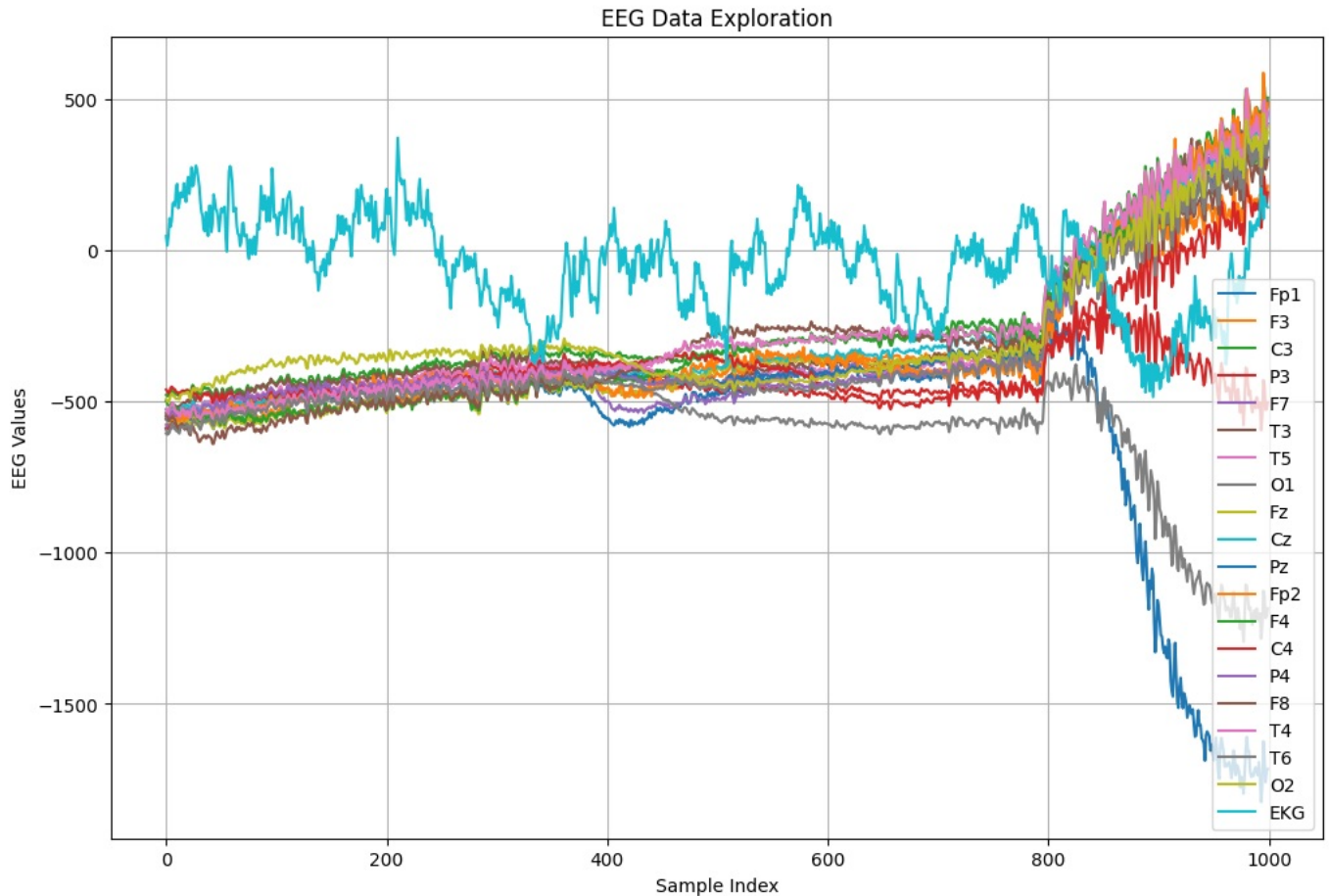
```

for column in eeg_data.columns:
    ax.plot(eeg_data.index[:num_samples_to_plot], eeg_data[column][:num_samples_to_plot], label=column)

ax.set_title('EEG Data Exploration')
ax.set_xlabel('Sample Index')
ax.set_ylabel('EEG Values')
ax.legend()
ax.grid(True) # Add grid lines for better readability

plt.show()

```



```

In [ ]: # Frequency Analysis
def plot_frequency_spectrum(eeg_data, sampling_rate=1000):
    n = len(eeg_data)
    k = np.arange(n)
    T = n / sampling_rate
    frq = k / T
    frq = frq[range(n//2)]

    fft_values = np.fft.fft(eeg_data.values)
    fft_values = fft_values[range(n//2)]

    plt.figure(figsize=(12, 6))
    plt.plot(frq, abs(fft_values))
    plt.title('Frequency Spectrum of EEG Data')
    plt.xlabel('Frequency (Hz)')
    plt.ylabel('Amplitude')
    plt.show()

```

```

In [ ]: # Temporal Analysis
def plot_temporal_analysis(eeg_data):
    plt.figure(figsize=(12, 6))
    for column in eeg_data.columns:
        plt.plot(eeg_data.index, eeg_data[column], label=column)

    plt.title('Temporal Analysis of EEG Data')
    plt.xlabel('Sample Index')
    plt.ylabel('EEG Values')
    plt.legend()
    plt.show()

```

```

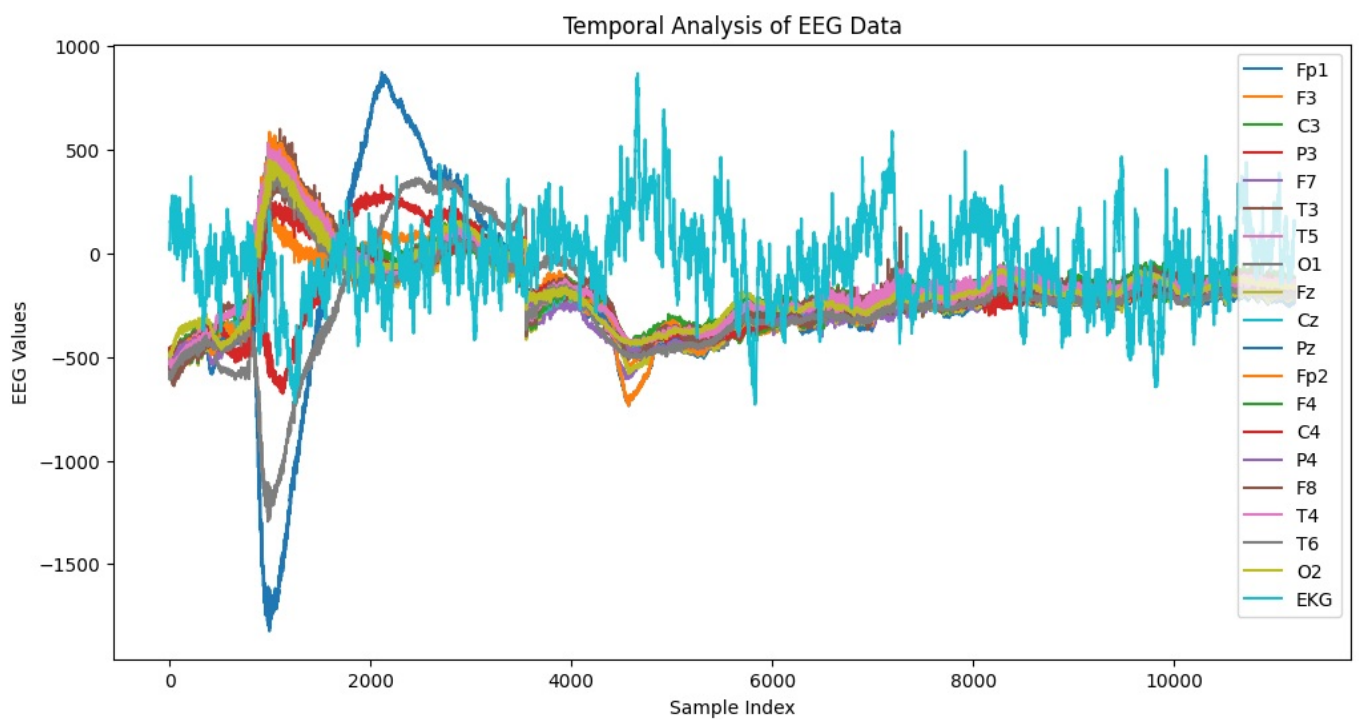
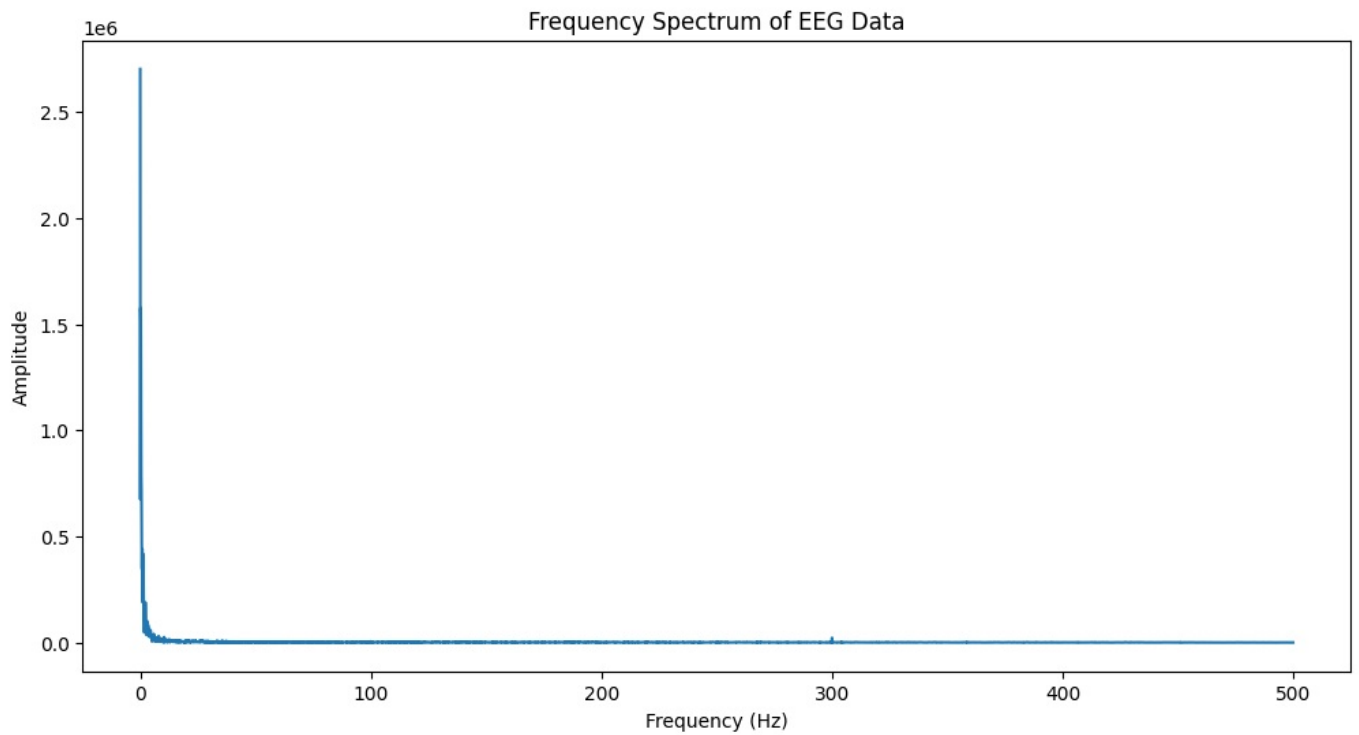
In [ ]: # Load EEG data
file_path = "/kaggle/input/hms-harmful-brain-activity-classification/train_eegs/1001369401.parquet"
eeg_data = pq.read_table(file_path).to_pandas()

# Visualize Frequency Analysis

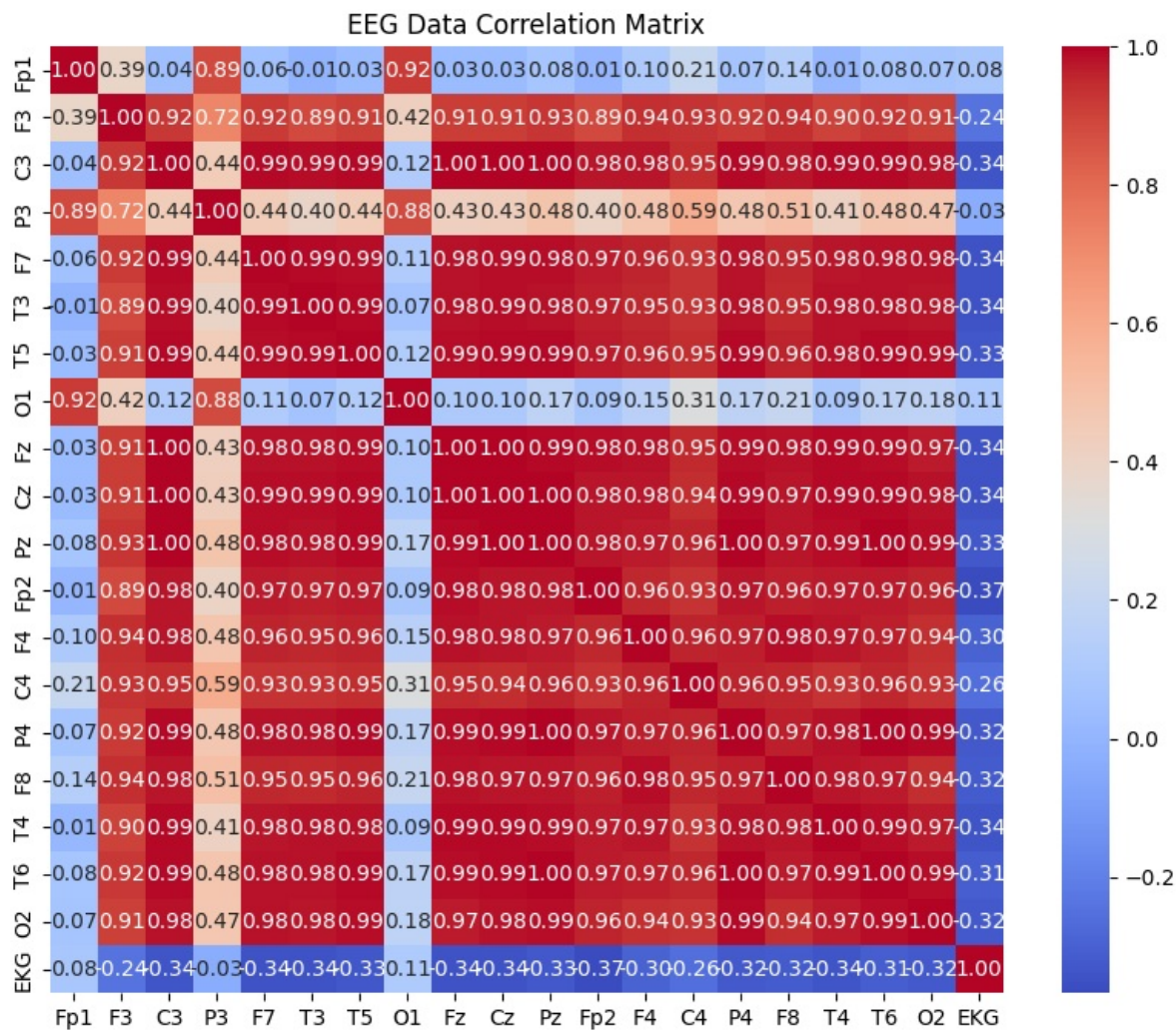
```

```
plot_frequency_spectrum(eeg_data.iloc[:, 0])
```

```
# Visualize Temporal Analysis
plot_temporal_analysis(eeg_data)
```



```
In [ ]: # EEG Data Correlation Matrix
correlation_matrix = eeg_data.corr()
plt.figure(figsize=(10, 8))
sns.heatmap(correlation_matrix, annot=True, cmap='coolwarm', fmt=".2f")
plt.title('EEG Data Correlation Matrix')
plt.show()
```

Explore train spectrograms

```
In [ ]: # File path for the spectrogram in Parquet format
spectrogram_file_path = pd.read_parquet("/kaggle/input/hms-harmful-brain-activity-classification/train_spectrog
spectrogram_file_path
```

```
Out[ ]:
```

	time	LL_0.59	LL_0.78	LL_0.98	LL_1.17	LL_1.37	LL_1.56	LL_1.76	LL_1.95	LL_2.15	...	RP_18.16	RP_1
0	1	28.680000	53.990002	67.629997	59.880001	50.880001	74.309998	78.480003	63.080002	59.869999	...	0.13	
1	3	29.639999	38.959999	44.009998	66.800003	48.509998	42.180000	47.340000	48.599998	40.880001	...	0.15	
2	5	8.890000	9.020000	16.360001	23.559999	27.340000	30.040001	27.559999	23.290001	15.120000	...	0.12	
3	7	1.770000	1.930000	1.810000	1.600000	1.430000	1.280000	1.190000	1.110000	1.010000	...	0.03	
4	9	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	...	NaN	
...	
295	591	2.210000	2.280000	2.200000	1.280000	1.350000	1.930000	2.300000	2.440000	2.310000	...	0.03	
296	593	2.490000	2.540000	2.150000	1.490000	1.360000	1.570000	1.970000	2.050000	1.890000	...	0.02	
297	595	0.240000	0.190000	0.210000	0.120000	0.110000	0.080000	0.060000	0.040000	0.040000	...	0.00	
298	597	0.990000	1.230000	1.370000	1.620000	1.940000	2.190000	2.270000	2.310000	2.300000	...	0.02	
299	599	1.810000	2.110000	1.950000	1.800000	1.950000	2.120000	2.330000	2.450000	2.430000	...	0.02	

300 rows × 401 columns

```
In [ ]: # Basic Information
print(f"Spectrogram Shape: {spectrogram_file_path.shape}")
print(f"Minimum Intensity: {spectrogram_file_path.min().min()} dB")
print(f"Maximum Intensity: {spectrogram_file_path.max().max()} dB")
print(f"Average Intensity: {spectrogram_file_path.mean().mean()} dB")
```

```
Spectrogram Shape: (300, 401)
Minimum Intensity: 0.0 dB
Maximum Intensity: 206415.046875 dB
Average Intensity: 55.734953598339956 dB
```

```
In [ ]: def plot_spectrogram(spectrogram_path):
    sample_spect = pd.read_parquet(spectrogram_path)

    split_spect = {
        "LL": sample_spect.filter(regex='^LL', axis=1),
        "RL": sample_spect.filter(regex='^RL', axis=1),
        "RP": sample_spect.filter(regex='^RP', axis=1),
        "LP": sample_spect.filter(regex='^LP', axis=1),
    }

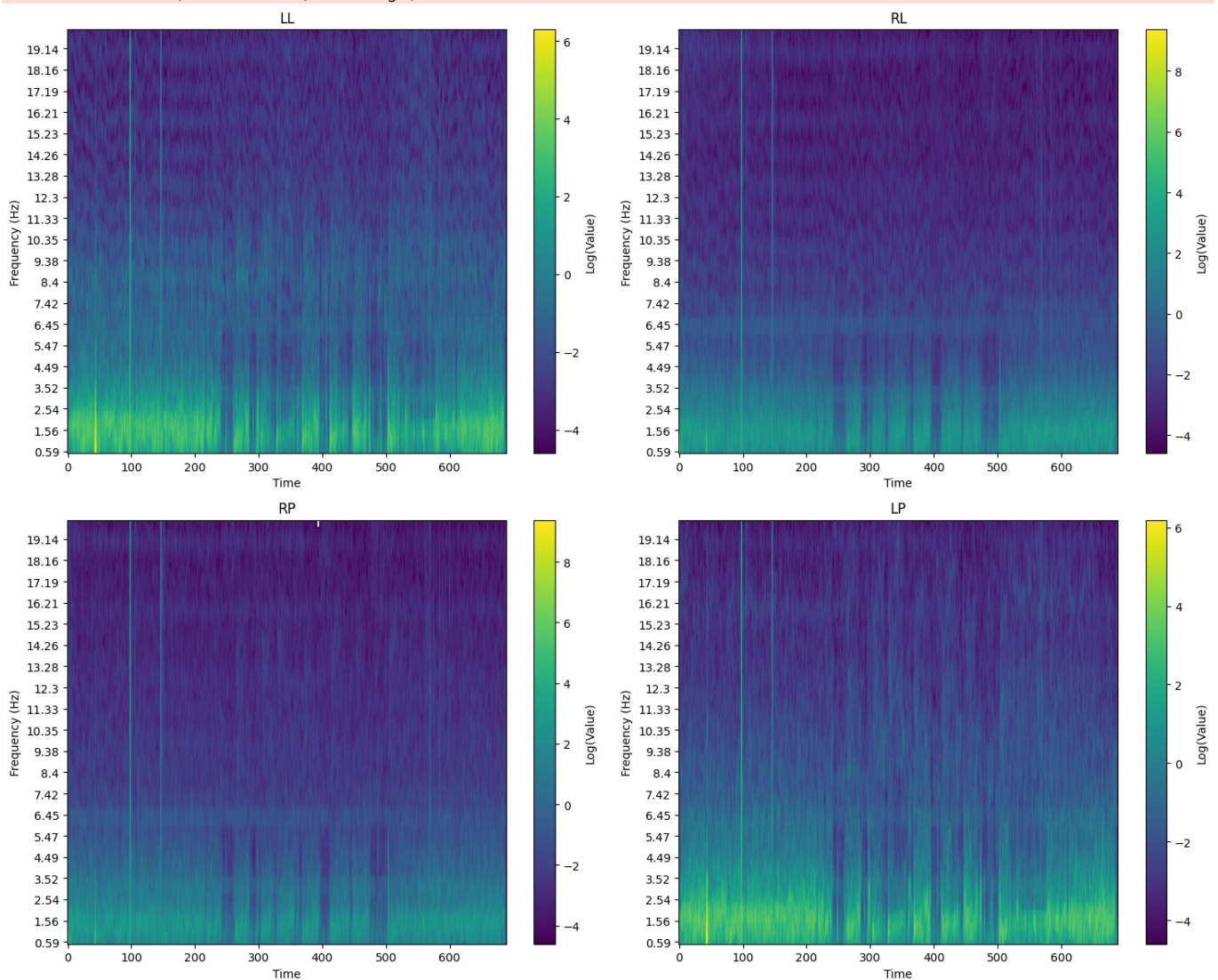
    fig, axes = plt.subplots(nrows=2, ncols=2, figsize=(15, 12))
    axes = axes.flatten()
    label_interval = 5
    for i, split_name in enumerate(split_spect.keys()):
        ax = axes[i]
        img = ax.imshow(np.log(split_spect[split_name]).T, cmap='viridis', aspect='auto', origin='lower')
        cbar = fig.colorbar(img, ax=ax)
        cbar.set_label('Log(Value)')
        ax.set_title(split_name)
        ax.set_ylabel("Frequency (Hz)")
        ax.set_xlabel("Time")

        ax.set_yticks(np.arange(len(split_spect[split_name].columns)))
        ax.set_yticklabels([column_name[3:] for column_name in split_spect[split_name].columns])
        frequencies = [column_name[3:] for column_name in split_spect[split_name].columns]
        ax.set_xticks(np.arange(0, len(split_spect[split_name].columns), label_interval))
        ax.set_xticklabels(frequencies[::label_interval])

    plt.tight_layout()
    plt.show()
```

```
In [ ]: # Corrected function call
plot_spectrogram("/kaggle/input/hms-harmful-brain-activity-classification/train_spectrograms/1000189855.parquet")
```

/opt/conda/lib/python3.10/site-packages/pandas/core/internals/blocks.py:329: RuntimeWarning: divide by zero encountered in log
result = func(self.values, **kwargs)



Listen to Spectrogram


```
In [ ]: !pip install librosa
```

```
Requirement already satisfied: librosa in /opt/conda/lib/python3.10/site-packages (0.10.1)
Requirement already satisfied: audioread>=2.1.9 in /opt/conda/lib/python3.10/site-packages (from librosa) (3.0.1)
Requirement already satisfied: numpy!=1.22.0,!1.22.1,!1.22.2,>=1.20.3 in /opt/conda/lib/python3.10/site-packages (from librosa) (1.24.3)
Requirement already satisfied: scipy>=1.2.0 in /opt/conda/lib/python3.10/site-packages (from librosa) (1.11.4)
Requirement already satisfied: scikit-learn>=0.20.0 in /opt/conda/lib/python3.10/site-packages (from librosa) (1.2.2)
Requirement already satisfied: joblib>=0.14 in /opt/conda/lib/python3.10/site-packages (from librosa) (1.3.2)
Requirement already satisfied: decorator>=4.3.0 in /opt/conda/lib/python3.10/site-packages (from librosa) (5.1.1)
Requirement already satisfied: numba>=0.51.0 in /opt/conda/lib/python3.10/site-packages (from librosa) (0.57.1)
Requirement already satisfied: soundfile>=0.12.1 in /opt/conda/lib/python3.10/site-packages (from librosa) (0.12.1)
Requirement already satisfied: pooch>=1.0 in /opt/conda/lib/python3.10/site-packages (from librosa) (1.8.0)
Requirement already satisfied: soxr>=0.3.2 in /opt/conda/lib/python3.10/site-packages (from librosa) (0.3.7)
Requirement already satisfied: typing-extensions>=4.1.1 in /opt/conda/lib/python3.10/site-packages (from librosa) (4.5.0)
Requirement already satisfied: lazy-loader>=0.1 in /opt/conda/lib/python3.10/site-packages (from librosa) (0.3)
Requirement already satisfied: msgpack>=1.0 in /opt/conda/lib/python3.10/site-packages (from librosa) (1.0.5)
Requirement already satisfied: llvmlite<0.41,>=0.40.0dev0 in /opt/conda/lib/python3.10/site-packages (from numba>=0.51.0->librosa) (0.40.1)
Requirement already satisfied: platformdirs>=2.5.0 in /opt/conda/lib/python3.10/site-packages (from pooch>=1.0->librosa) (4.1.0)
Requirement already satisfied: packaging>=20.0 in /opt/conda/lib/python3.10/site-packages (from pooch>=1.0->librosa) (21.3)
Requirement already satisfied: requests>=2.19.0 in /opt/conda/lib/python3.10/site-packages (from pooch>=1.0->librosa) (2.31.0)
Requirement already satisfied: threadpoolctl>=2.0.0 in /opt/conda/lib/python3.10/site-packages (from scikit-learn>=0.20.0->librosa) (3.2.0)
Requirement already satisfied: cffi>=1.0 in /opt/conda/lib/python3.10/site-packages (from soundfile>=0.12.1->librosa) (1.15.1)
Requirement already satisfied: pycparser in /opt/conda/lib/python3.10/site-packages (from cffi>=1.0->soundfile>=0.12.1->librosa) (2.21)
Requirement already satisfied: pyparsing!=3.0.5,>=2.0.2 in /opt/conda/lib/python3.10/site-packages (from packaging>=20.0->pooch>=1.0->librosa) (3.0.9)
Requirement already satisfied: charset-normalizer<4,>=2 in /opt/conda/lib/python3.10/site-packages (from requests>=2.19.0->pooch>=1.0->librosa) (3.2.0)
Requirement already satisfied: idna<4,>=2.5 in /opt/conda/lib/python3.10/site-packages (from requests>=2.19.0->pooch>=1.0->librosa) (3.4)
Requirement already satisfied: urllib3<3,>=1.21.1 in /opt/conda/lib/python3.10/site-packages (from requests>=2.19.0->pooch>=1.0->librosa) (1.26.15)
Requirement already satisfied: certifi>=2017.4.17 in /opt/conda/lib/python3.10/site-packages (from requests>=2.19.0->pooch>=1.0->librosa) (2023.11.17)
```

```
In [ ]: import librosa.display
import librosa
import soundfile as sf # Import soundfile module for writing WAV files
```

```
In [ ]: # Replace this with your actual Parquet file path
parquet_file_path = '/kaggle/input/hms-harmful-brain-activity-classification/train_spectrograms/1001616430.parq

# Read the Parquet file into a DataFrame
df = pd.read_parquet(parquet_file_path)

# Extract the spectrogram data from the DataFrame
spectrogram = df.to_numpy()

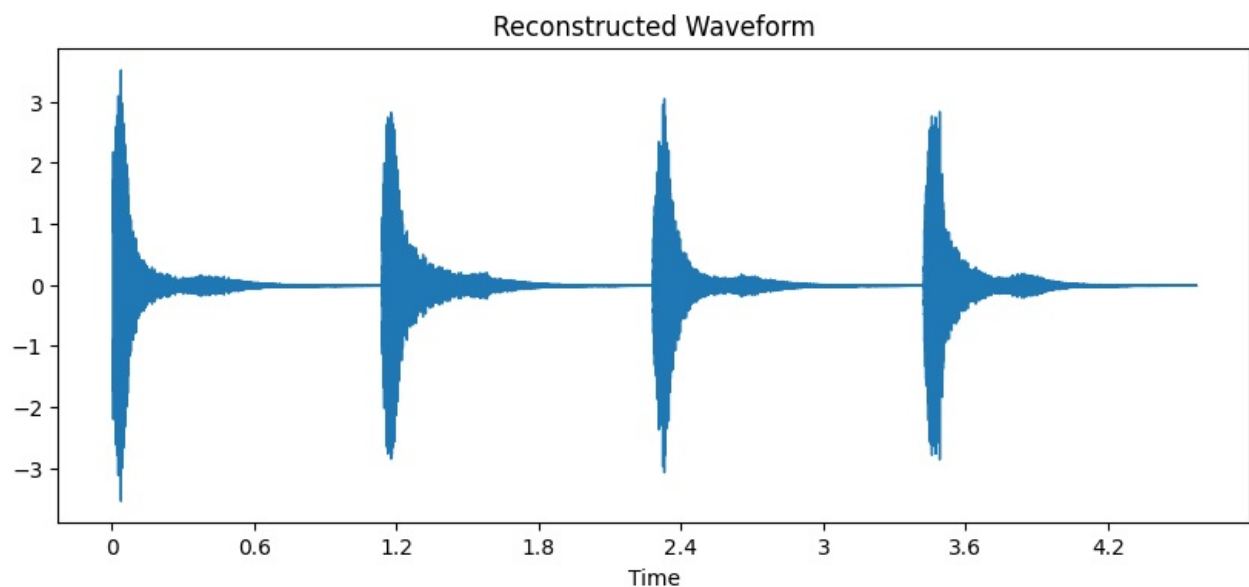
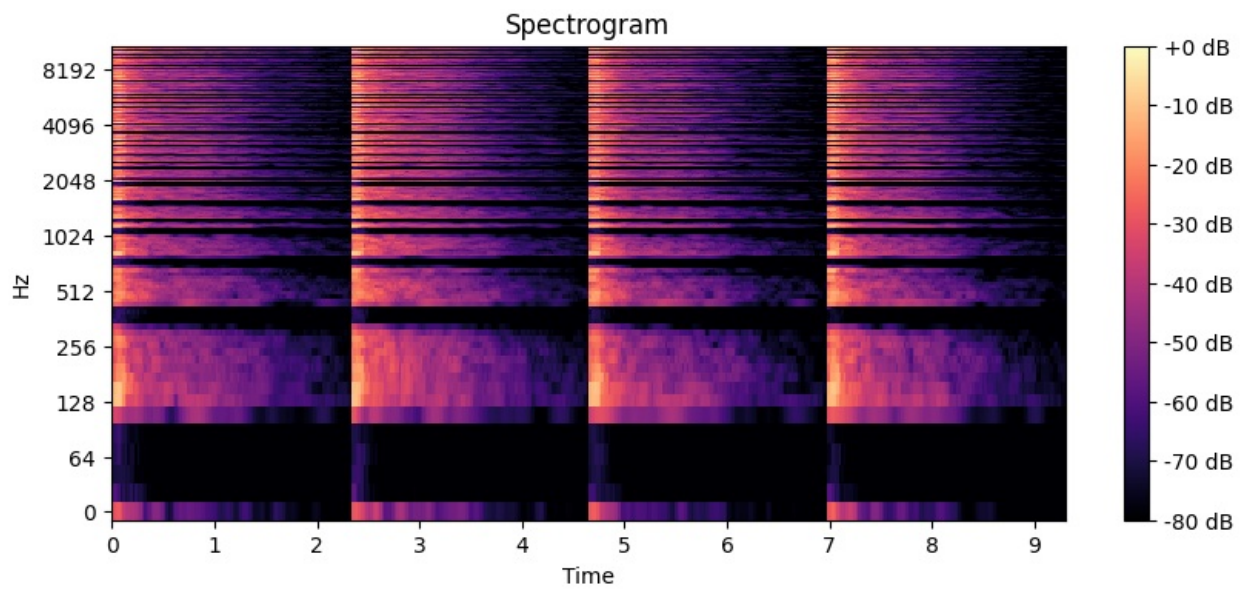
# Assuming 'sr' is the sampling rate of your original audio
# Make sure to replace this with your actual sampling rate
sr = 22050

# Use librosa's iSTFT to convert the spectrogram back to a time-domain signal
audio_signal = librosa.istft(spectrogram)

# Display the original spectrogram
plt.figure(figsize=(10, 4))
librosa.display.specshow(librosa.amplitude_to_db(np.abs(spectrogram), ref=np.max), y_axis='log', x_axis='time')
plt.colorbar(format='%+2.0f dB')
plt.title('Spectrogram')
plt.show()

# Display the reconstructed waveform
plt.figure(figsize=(10, 4))
librosa.display.waveshow(audio_signal, sr=sr)
plt.title('Reconstructed Waveform')
plt.show()

# Save the reconstructed waveform as an audio file
sf.write('reconstructed_audio.wav', audio_signal, sr)
```



```
In [ ]: from IPython.display import Audio

# Specify the path to the reconstructed audio file
reconstructed_audio_path = 'reconstructed_audio.wav'

# Play the reconstructed audio
Audio(reconstructed_audio_path)
```

Out[]: Your browser does not support the audio element.

Final Submission

```
In [ ]: # Assuming you have predictions for each class from your model
# Replace these lines with your actual predictions
seizure_predictions = [0.2, 0.1, 0.3, 0.4, 0.5] # Replace with your seizure predictions
lpd_predictions = [0.1, 0.2, 0.4, 0.3, 0.5] # Replace with your LPD predictions
gpd_predictions = [0.3, 0.1, 0.2, 0.5, 0.4] # Replace with your GPD predictions
lrda_predictions = [0.4, 0.5, 0.1, 0.2, 0.3] # Replace with your LRDA predictions
grda_predictions = [0.5, 0.4, 0.3, 0.1, 0.2] # Replace with your GRDA predictions

# Combine predictions into a DataFrame
predictions_df = pd.DataFrame({
    'seizure_vote': seizure_predictions,
    'lpd_vote': lpd_predictions,
    'gpd_vote': gpd_predictions,
    'lrda_vote': lrda_predictions,
    'grda_vote': grda_predictions,
    'other_vote': [0.0] * len(seizure_predictions) # Add placeholder for 'other_vote'
})

# Add 'eeg_id' column from the test_df
predictions_df['eeg_id'] = test_data['eeg_id']
```

```

# Define the columns order
columns_order = ['eeg_id', 'seizure_vote', 'lpd_vote', 'gpd_vote', 'lrda_vote', 'grda_vote', 'other_vote']

# Reorder columns
predictions_df = predictions_df[columns_order]

# Normalize the predictions to sum up to 1 for each row
normalized_predictions = predictions_df.iloc[:, 1:].div(predictions_df.iloc[:, 1:].sum(axis=1), axis=0)
predictions_df.iloc[:, 1:] = normalized_predictions

# Save the predictions to a CSV file
predictions_df.to_csv("submission.csv", index=False)

# Display the final DataFrame
print(predictions_df)

```

	eeg_id	seizure_vote	lpd_vote	gpd_vote	lrda_vote	grda_vote	\
0	3.911565e+09	0.133333	0.066667	0.200000	0.266667	0.333333	
1	NaN	0.076923	0.153846	0.076923	0.384615	0.307692	
2	NaN	0.230769	0.307692	0.153846	0.076923	0.230769	
3	NaN	0.266667	0.200000	0.333333	0.133333	0.066667	
4	NaN	0.263158	0.263158	0.210526	0.157895	0.105263	

	other_vote
0	0.0
1	0.0
2	0.0
3	0.0
4	0.0

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