

EDA Visualizations and Audio Exploration

Import Libraries

```
In [1]: import ast
import librosa
import numpy as np
import pandas as pd
import seaborn as sns
import plotly.express as px
import plotly.graph_objs as go
from collections import Counter
import matplotlib.pyplot as plt
from IPython.display import Audio
```

EDA

```
In [2]: trainmeta_df = pd.read_csv("/input/birdclef-2023/train_metadata.csv")
trainmeta_df.head()
```

	primary_label	secondary_labels	type	latitude	longitude	scientific_name	common_name	author	license	rating	link
0	abethr1	['song']	4.3906	38.2788	Turdus tephronotus	African Bare-eyed Thrush	Rolf A. de By	Creative Commons Attribution-NonCommercial-Sha...	4.0	https://www.canto.org/1	
1	abethr1	['call']	-2.9524	38.2921	Turdus tephronotus	African Bare-eyed Thrush	James Bradley	Creative Commons Attribution-NonCommercial-Sha...	3.5	https://www.canto.org/3	
2	abethr1	['song']	-2.9524	38.2921	Turdus tephronotus	African Bare-eyed Thrush	James Bradley	Creative Commons Attribution-NonCommercial-Sha...	3.5	https://www.canto.org/3	
3	abethr1	['song']	-2.9524	38.2921	Turdus tephronotus	African Bare-eyed Thrush	James Bradley	Creative Commons Attribution-NonCommercial-Sha...	5.0	https://www.canto.org/3	
4	abethr1	['call', 'song']	-2.9524	38.2921	Turdus tephronotus	African Bare-eyed Thrush	James Bradley	Creative Commons Attribution-NonCommercial-Sha...	4.5	https://www.canto.org/3	

```
In [3]: x = trainmeta_df[trainmeta_df["primary_label"] == "afgfly1"]
x
```

Out[3]:

	primary_label	secondary_labels	type	latitude	longitude	scientific_name	common_name	author	license	rating
356	afgfly1	[]	['song']	-7.5253	34.8521	Bradornis microrhynchus	African Gray Flycatcher	Martin St-Michel	Creative Commons Attribution-NonCommercial-Sha...	3.0
357	afgfly1	[]	['call']	8.9200	40.0430	Bradornis microrhynchus	African Gray Flycatcher	Andrew Spencer	Creative Commons Attribution-NonCommercial-Sha...	5.0
358	afgfly1	[]	['call']	8.9200	40.0430	Bradornis microrhynchus	African Gray Flycatcher	Andrew Spencer	Creative Commons Attribution-NonCommercial-Sha...	5.0
359	afgfly1	[]	['call']	-1.5765	36.6316	Bradornis microrhynchus	African Gray Flycatcher	James Bradley	Creative Commons Attribution-NonCommercial-Sha...	5.0
360	afgfly1	['combol2', 'kerspa2', 'ratcis1']	['call,begging call,juvenile']	-2.8145	37.4113	Bradornis microrhynchus	African Gray Flycatcher	Rory Nefdt	Creative Commons Attribution-NonCommercial-Sha...	3.5
361	afgfly1	[]	['adult', 'call', 'sex uncertain']	-3.1481	36.6951	Bradornis microrhynchus	African Gray Flycatcher	isaac kilusu	Creative Commons Attribution-NonCommercial-Sha...	4.0
362	afgfly1	[]	['call', 'juvenile', 'sex uncertain']	-3.1481	36.6951	Bradornis microrhynchus	African Gray Flycatcher	isaac kilusu	Creative Commons Attribution-NonCommercial-Sha...	5.0
363	afgfly1	[]	['adult', 'call', 'sex uncertain']	-3.1481	36.6951	Bradornis microrhynchus	African Gray Flycatcher	isaac kilusu	Creative Commons Attribution-NonCommercial-Sha...	5.0

Train Meta Data

In [4]: `print(trainmeta_df.info())`

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 16941 entries, 0 to 16940
Data columns (total 12 columns):
 #   Column           Non-Null Count  Dtype  
--- 
 0   primary_label    16941 non-null   object  
 1   secondary_labels 16941 non-null   object  
 2   type              16941 non-null   object  
 3   latitude          16714 non-null   float64 
 4   longitude         16714 non-null   float64 
 5   scientific_name   16941 non-null   object  
 6   common_name       16941 non-null   object  
 7   author            16941 non-null   object  
 8   license           16941 non-null   object  
 9   rating            16941 non-null   float64 
 10  url               16941 non-null   object  
 11  filename          16941 non-null   object  
dtypes: float64(3), object(9)
memory usage: 1.6+ MB
None
```

Plots

In [5]: `fig = px.histogram(trainmeta_df, x="primary_label", nbins=len(trainmeta_df["primary_label"].unique()))
fig.update_layout(title_text="Distribution of Primary Labels")
fig.show()`

Distribution of Primary Labels



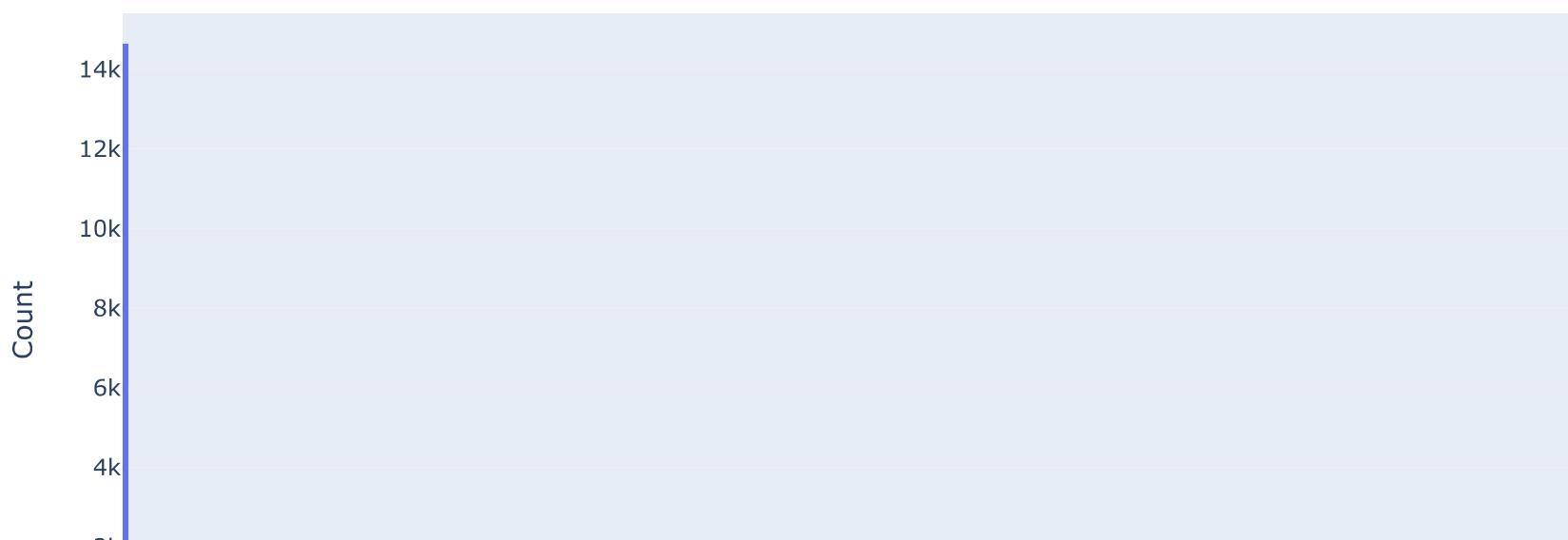
```
In [6]: secondary_df = trainmeta_df['secondary_labels'].str.replace('[', '').str.replace(']', '').str.replace('\\', '').str.split(',')
secondary_df = pd.merge(trainmeta_df.drop(columns=['secondary_labels']), secondary_df, left_index=True, right_index=True)

fig = px.histogram(secondary_df, x="secondary_label", title="Distribution of Secondary Labels")
fig.update_layout(xaxis_title="Secondary Label", yaxis_title="Count")
fig.show()
```

/opt/conda/lib/python3.7/site-packages/ipykernel_launcher.py:1: FutureWarning:

The default value of regex will change from True to False in a future version. In addition, single character regular expressions will *not* be treated as literal strings when regex=True.

Distribution of Secondary Labels



```
In [7]: # Flatten the list of labels in the "type" column
labels = [label.strip("[]") for sublist in trainmeta_df['type'].apply(ast.literal_eval) for label in sublist]

# Count the occurrence of each label
label_counts = Counter(labels)

# Create a bar plot of the label counts
fig = px.bar(x=list(label_counts.keys()), y=list(label_counts.values()))
fig.update_layout(title_text="Distribution of Types")
fig.show()
```

Distribution of Types



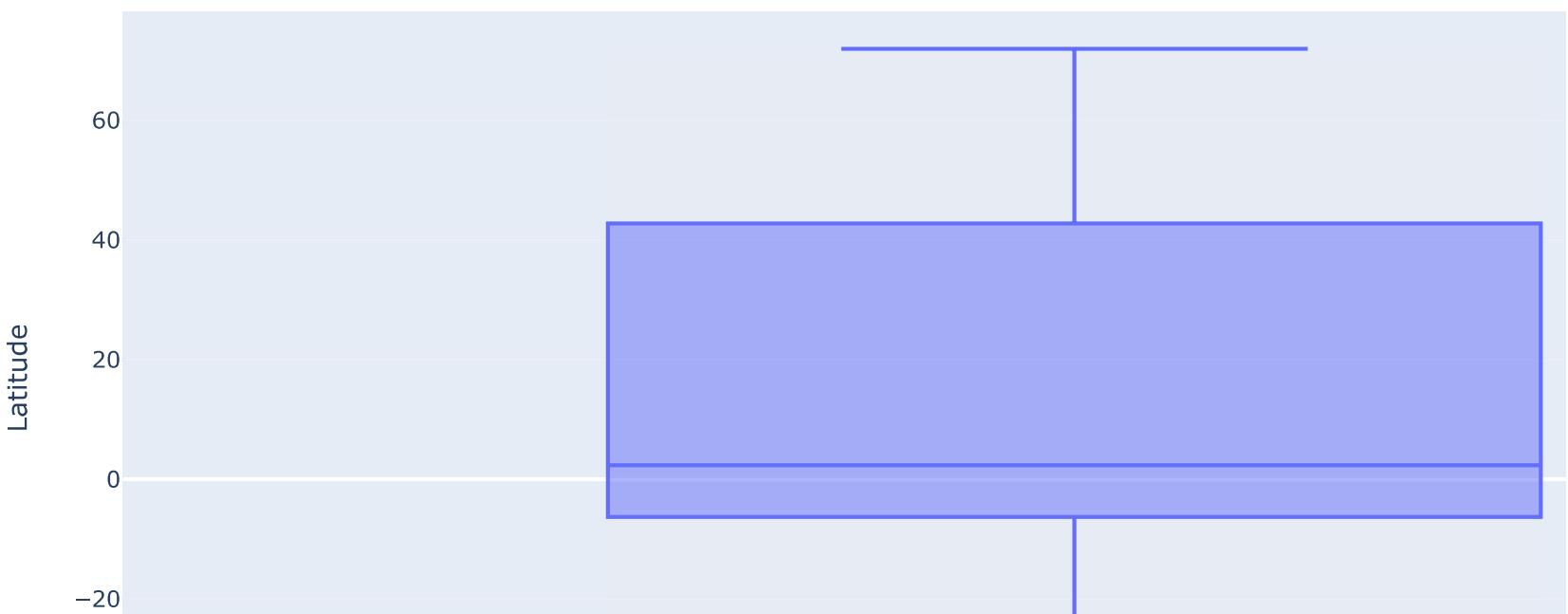
```
In [8]: fig1 = px.box(trainmeta_df, y="latitude")
fig1.update_layout(title_text="Distribution of Latitude",
                  yaxis=dict(title="Latitude"))

# create a box plot for longitude
fig2 = px.box(trainmeta_df, y="longitude")
fig2.update_layout(title_text="Distribution of Longitude",
                  yaxis=dict(title="Longitude"))

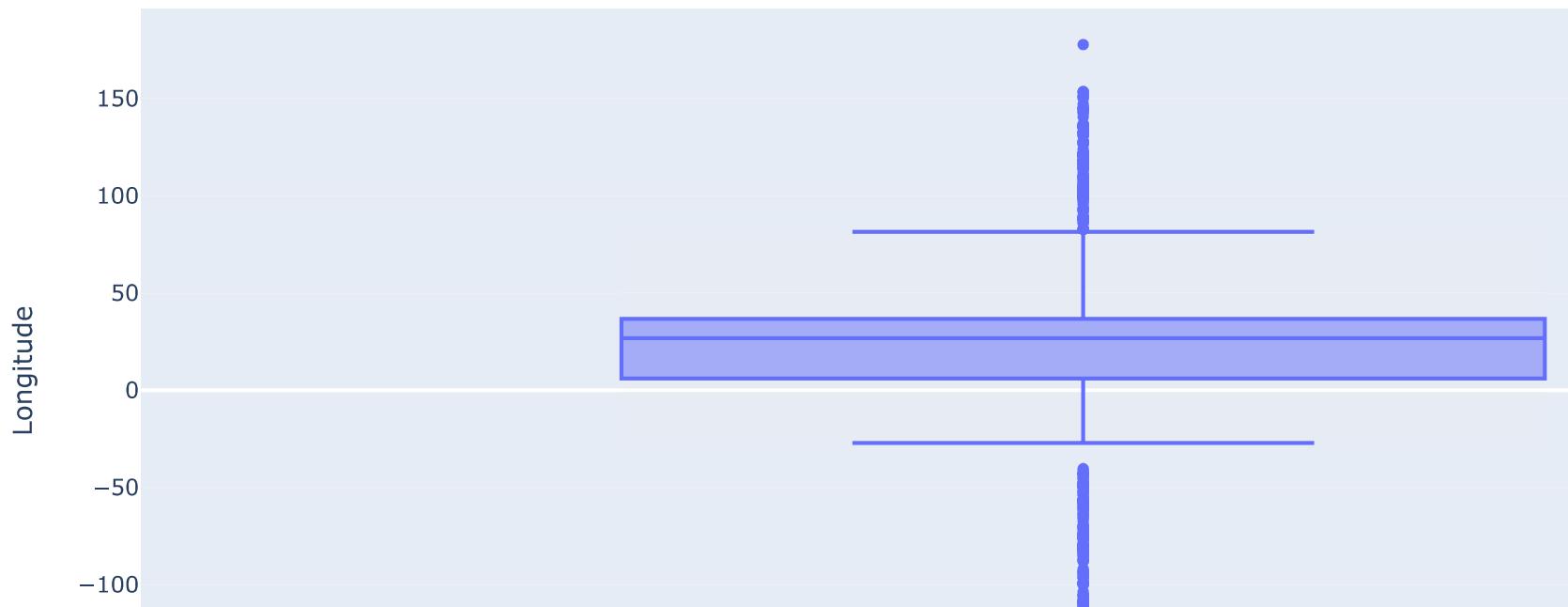
# create a box plot for rating
fig3 = px.box(trainmeta_df, y="rating")
fig3.update_layout(title_text="Distribution of Ratings",
                  yaxis=dict(title="Rating"))

# show the figures
fig1.show()
fig2.show()
fig3.show()
```

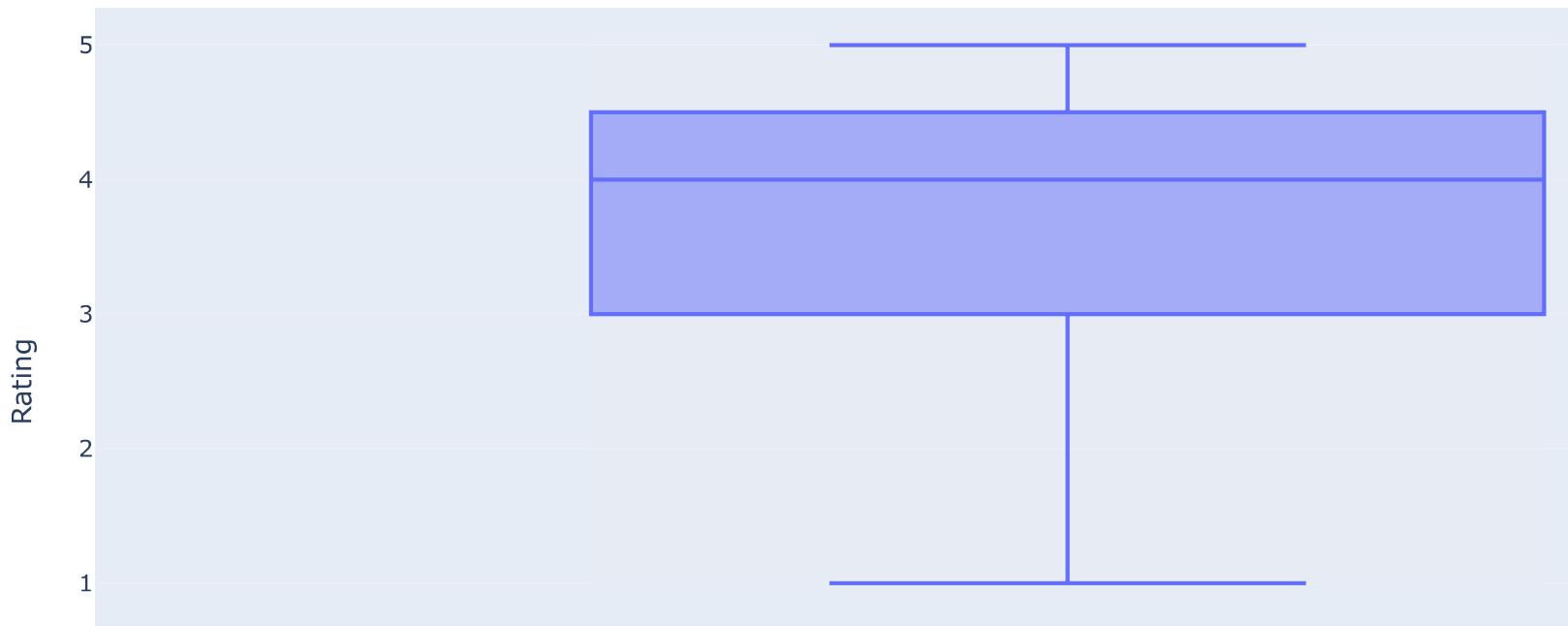
Distribution of Latitude



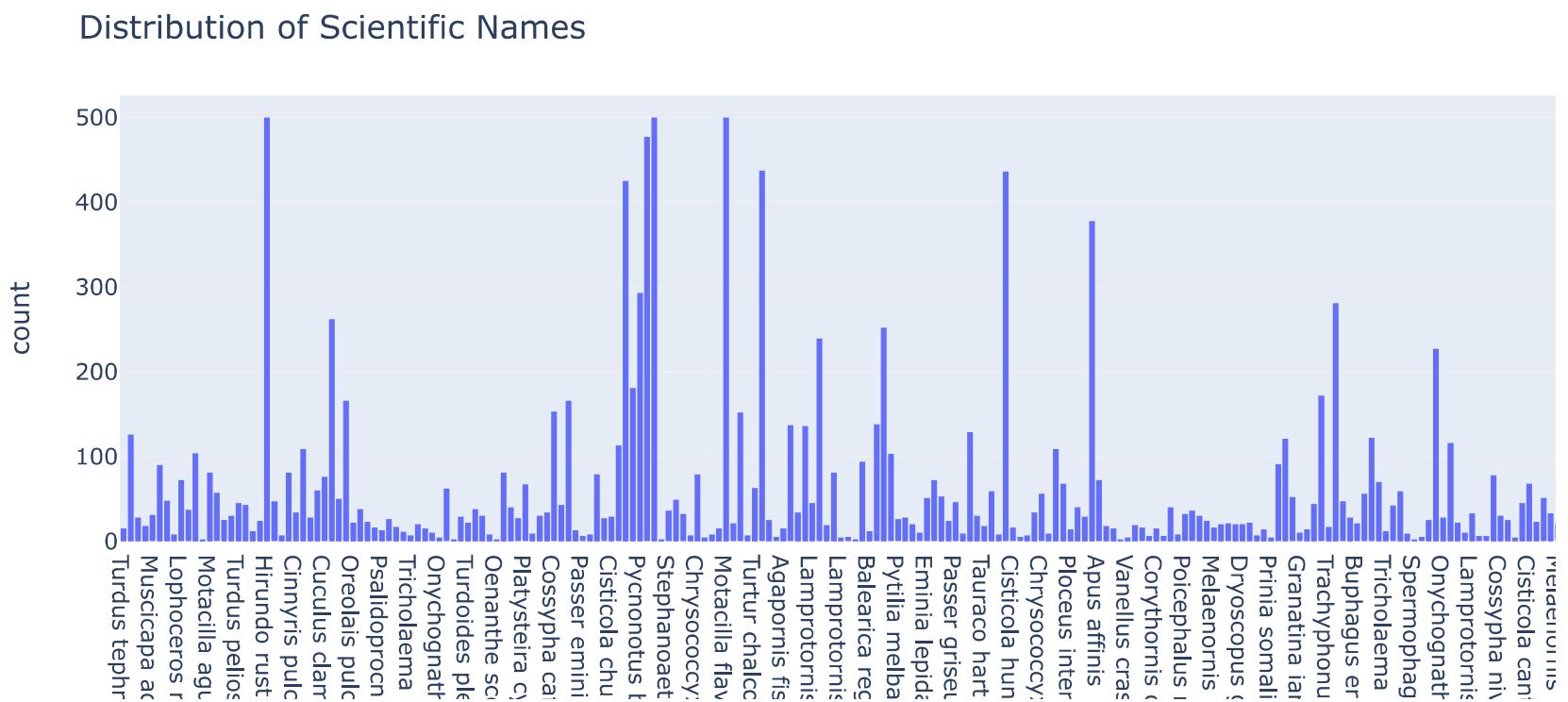
Distribution of Longitude



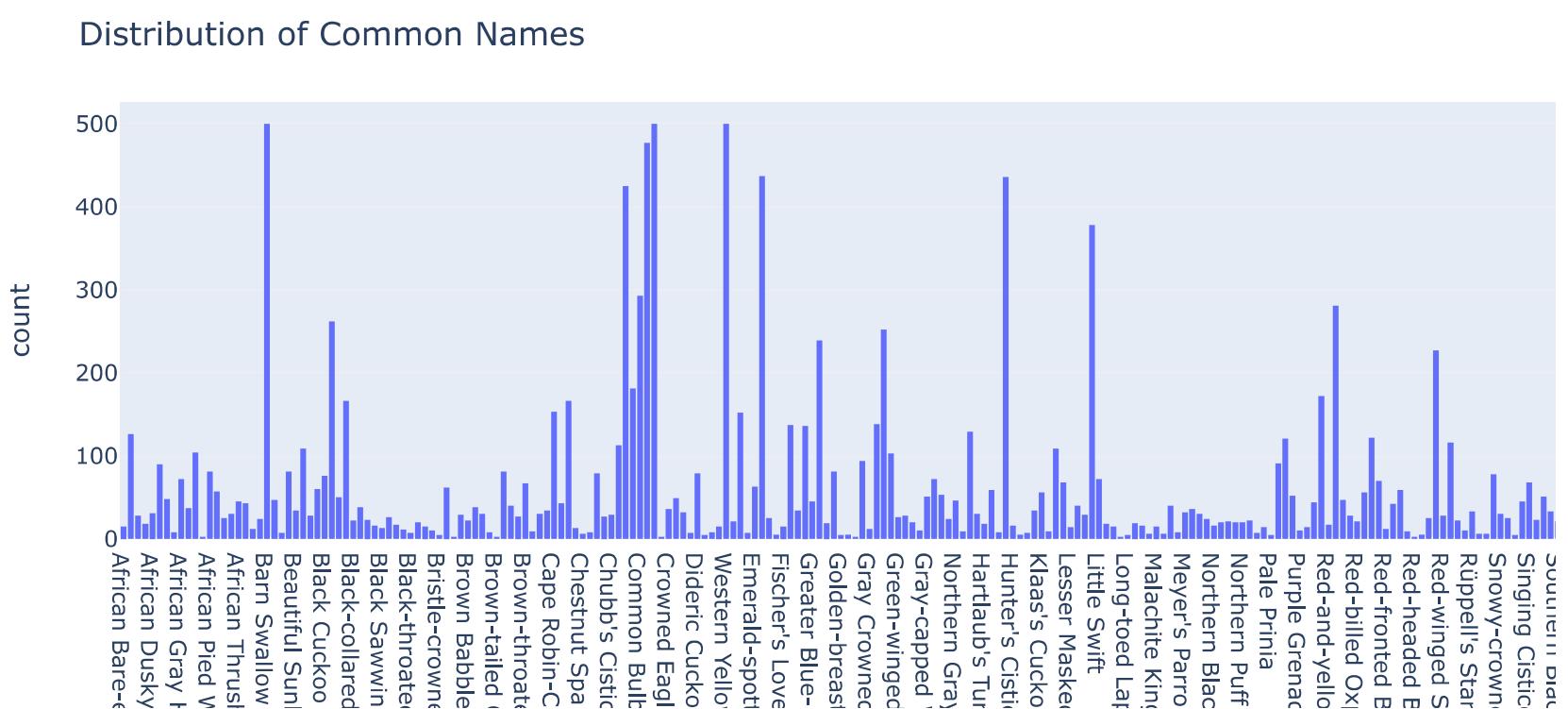
Distribution of Ratings



```
In [9]: fig = px.histogram(trainmeta_df, x="scientific_name", nbins=len(trainmeta_df["scientific_name"].unique()))
fig.update_layout(title_text="Distribution of Scientific Names")
fig.show()
```

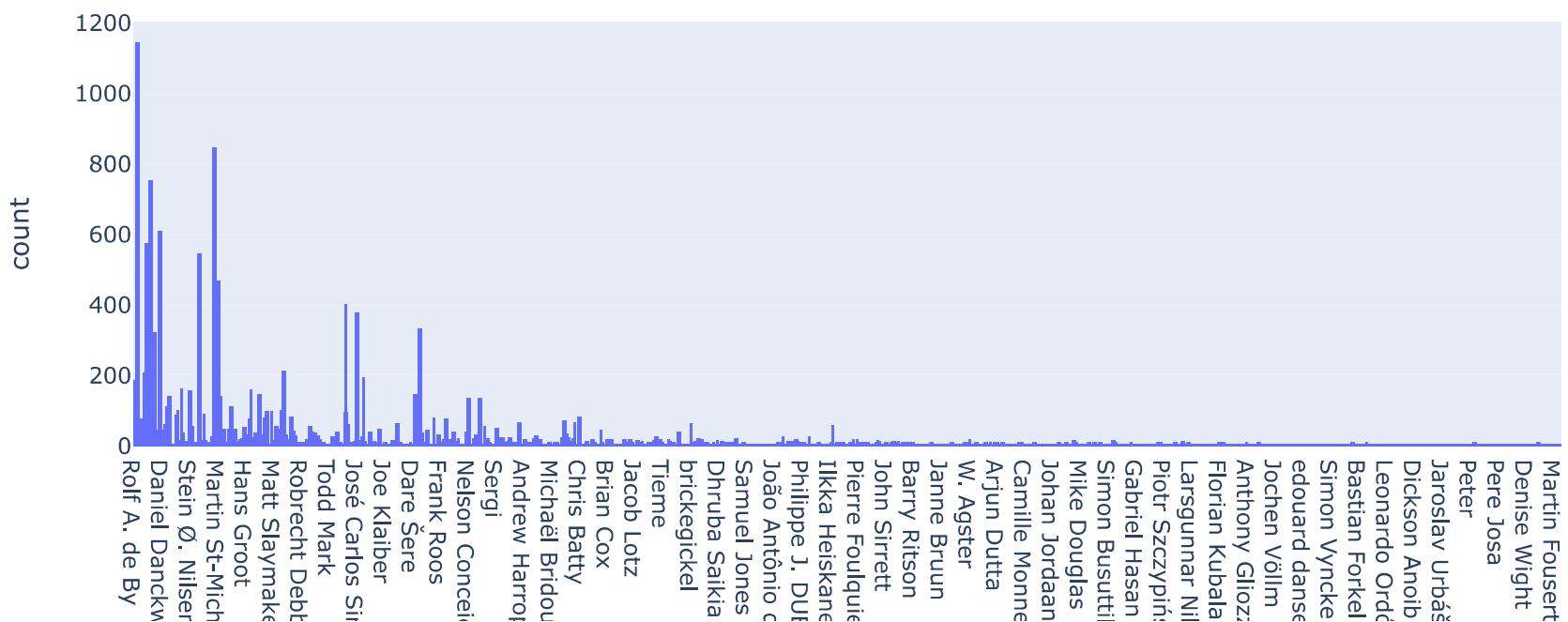


```
In [10]: fig = px.histogram(trainmeta_df, x="common_name", nbins=len(trainmeta_df["common_name"].unique()))
fig.update_layout(title_text="Distribution of Common Names")
fig.show()
```



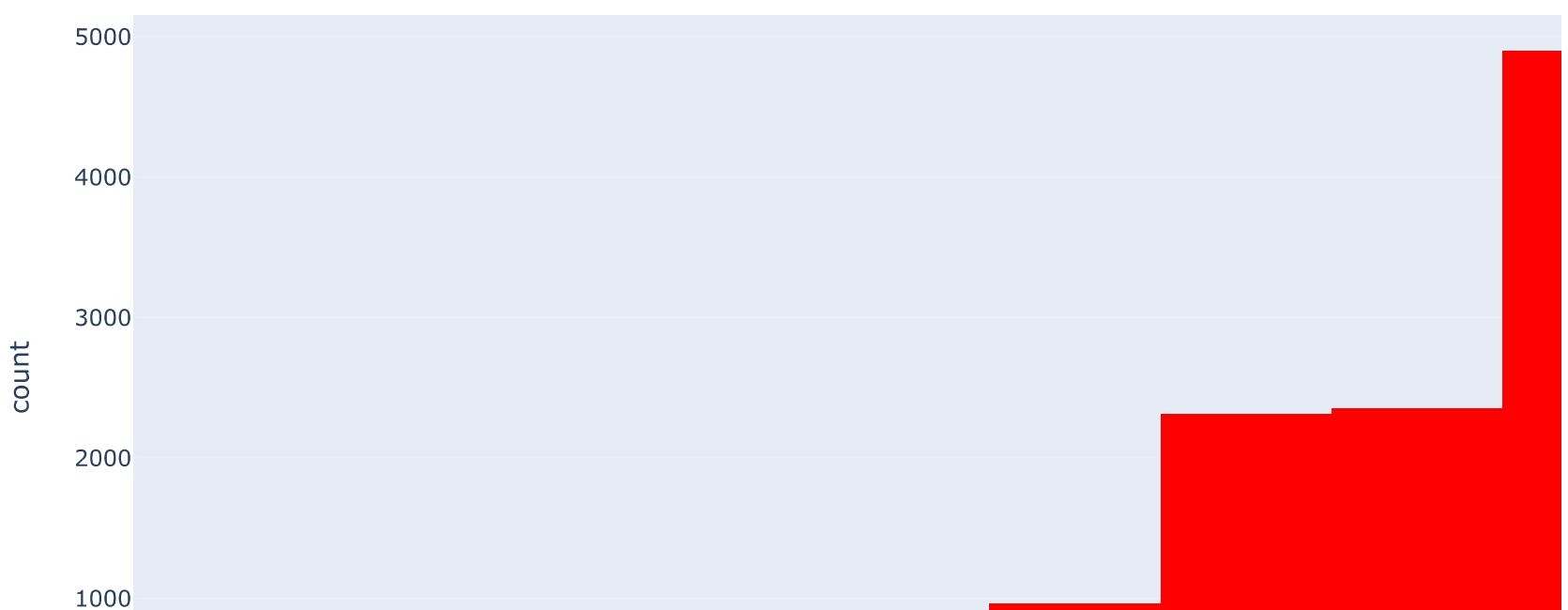
```
In [11]: fig = px.histogram(trainmeta_df, x="author", nbins=len(trainmeta_df["author"].unique()))
fig.update_layout(title_text="Distribution of Authors")
fig.show()
```

Distribution of Authors



```
In [12]: fig = px.histogram(trainmeta_df, x="rating", nbins=len(trainmeta_df["rating"].unique()), color_discrete_sequence=['red'])
fig.update_layout(title_text="Distribution of Ratings")
fig.show()
```

Distribution of Ratings



Correlation Plots

```
In [13]: # drop columns from correlation matrix
corr = trainmeta_df.corr()

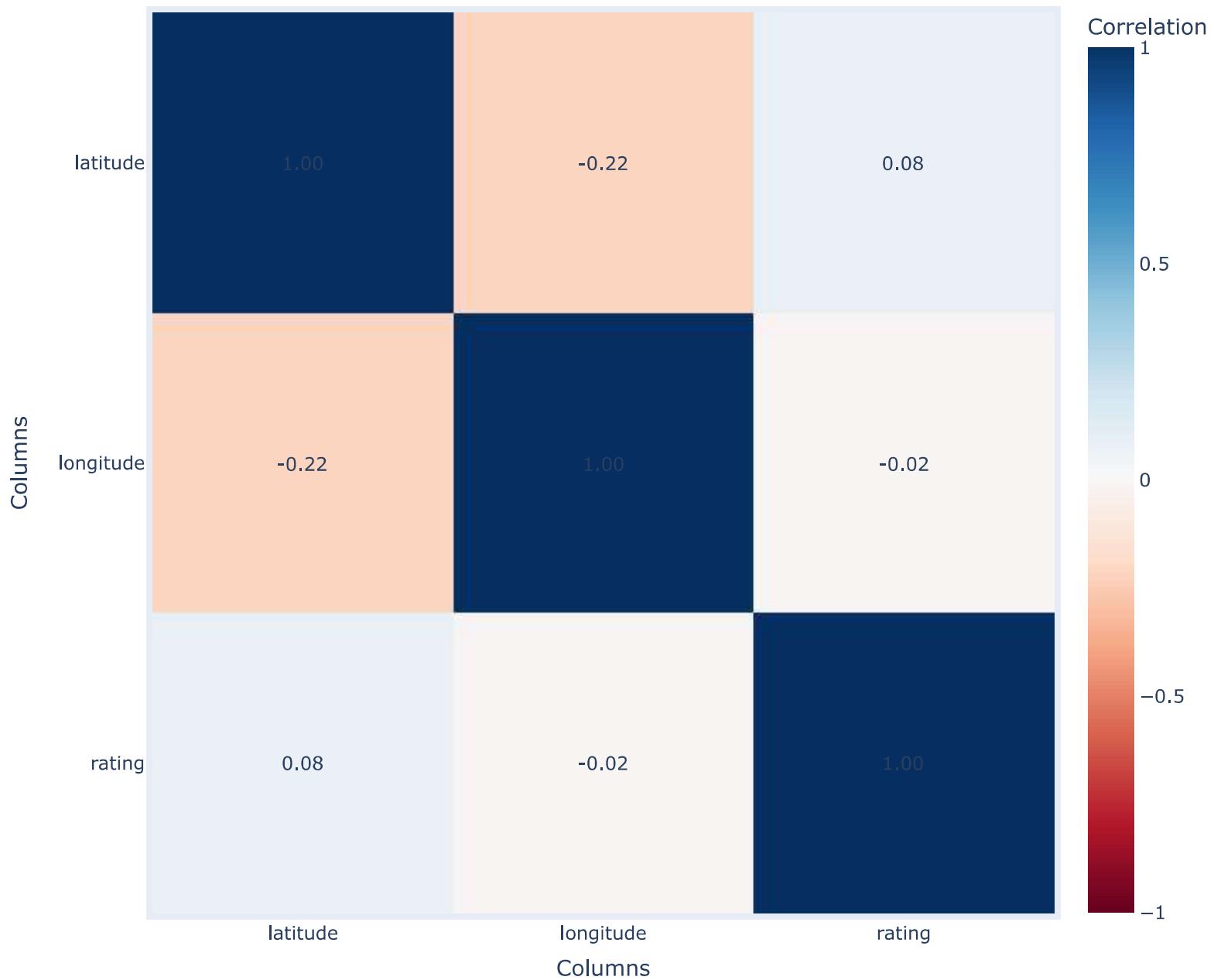
# create correlation heatmap
fig = px.imshow(corr,
                 labels=dict(x="Columns", y="Columns", color="Correlation"),
                 x=corr.columns,
                 y=corr.columns,
                 color_continuous_scale='RdBu',
                 zmin=-1,
                 zmax=1,
                 title="Correlation Heatmap")

# add text annotations
annotations = []
for i, row in enumerate(corr.values):
    for j, value in enumerate(row):
        text = '{:.2f}'.format(value)
        annotations.append(dict(x=corr.columns[j], y=corr.columns[i], text=text, showarrow=False))

fig.update_layout(width=800, height=800)
```

```
fig.update_traces(showscale=True, colorbar_thickness=25, colorbar_len=0.75)
fig.update_layout(margin=dict(l=50, r=50, b=100, t=100, pad=4))
fig.update_layout(annotations=annotations)
fig.show()
```

Correlation Heatmap

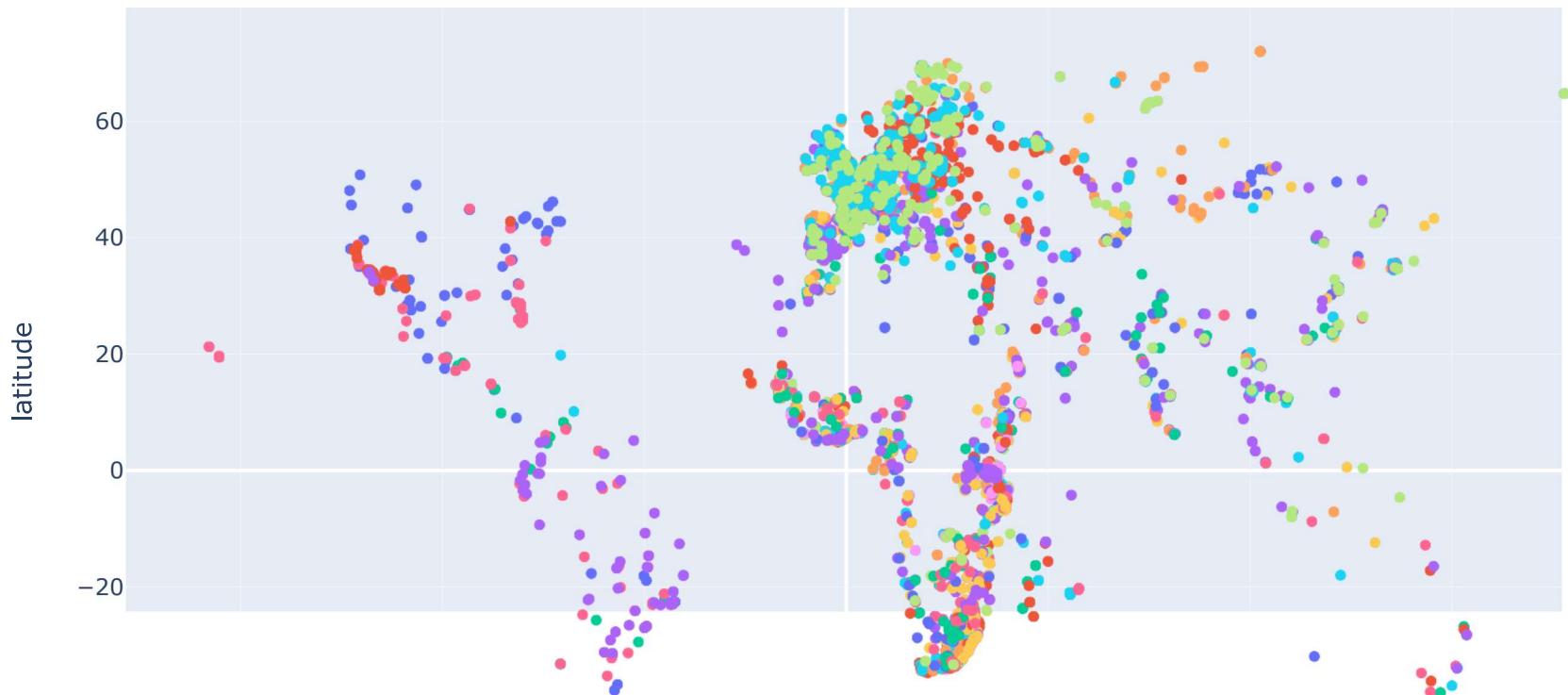


Interactive Map Plots

Scatter Plot

```
In [14]: fig = px.scatter(trainmeta_df, x="longitude", y="latitude", color="common_name")
fig.update_layout(title="Distribution of Recordings by Location")
fig.show()
```

Distribution of Recordings by Location



Map Box (Open Street Map)

```
In [15]: fig = px.scatter_mapbox(trainmeta_df, lat="latitude", lon="longitude", color="common_name",
                           hover_name="filename", hover_data=["common_name", "author", "rating"],
                           zoom=3, height=500)
fig.update_layout(mapbox_style="open-street-map")
fig.update_layout(margin={"r":0, "t":0, "l":0, "b":0})
fig.show()
```

EBird Taxonomy

```
In [17]: train_species_df = pd.read_csv("/input/birdclef-2023/eBird_Taxonomy_v2021.csv")
train_species_df.head()
```

	TAXON_ORDER	CATEGORY	SPECIES_CODE	PRIMARY_COM_NAME	SCI_NAME	ORDER1	FAMILY	SPECIES_GROUP
0	1	species	ostric2	Common Ostrich	Struthio camelus	Struthioniformes	Struthionidae (Ostriches)	Ostriches
1	6	species	ostric3	Somali Ostrich	Struthio molybdophanes	Struthioniformes	Struthionidae (Ostriches)	NaN
2	7	slash	y00934	Common/Somali Ostrich	Struthio camelus/molybdophanes	Struthioniformes	Struthionidae (Ostriches)	NaN
3	8	species	grerhe1	Greater Rhea	Rhea americana	Rheiformes	Rheidae (Rheas)	Rheas
4	14	species	lesrhe2	Lesser Rhea	Rhea pennata	Rheiformes	Rheidae (Rheas)	NaN

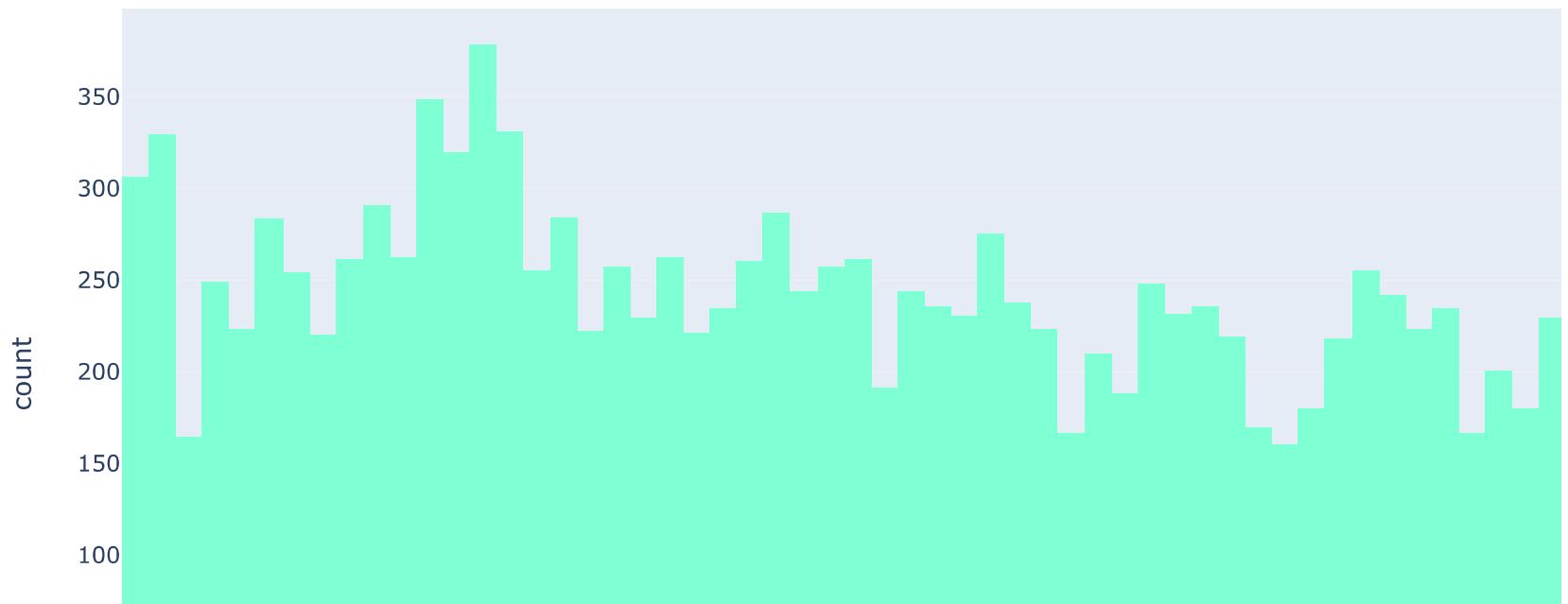
```
In [18]: # Histogram of the taxonomic order counts
fig1 = px.histogram(train_species_df, x="TAXON_ORDER", color_discrete_sequence=['aquamarine'])
fig1.update_layout(title_text="Distribution of Taxonomic Orders")

# Bar plot of the species group counts
# Box plot of the taxonomic order counts by category
fig3 = px.box(train_species_df, x="CATEGORY", y="TAXON_ORDER", color_discrete_sequence=['red'])
fig3.update_layout(title_text="Taxonomic Order Distribution by Category")

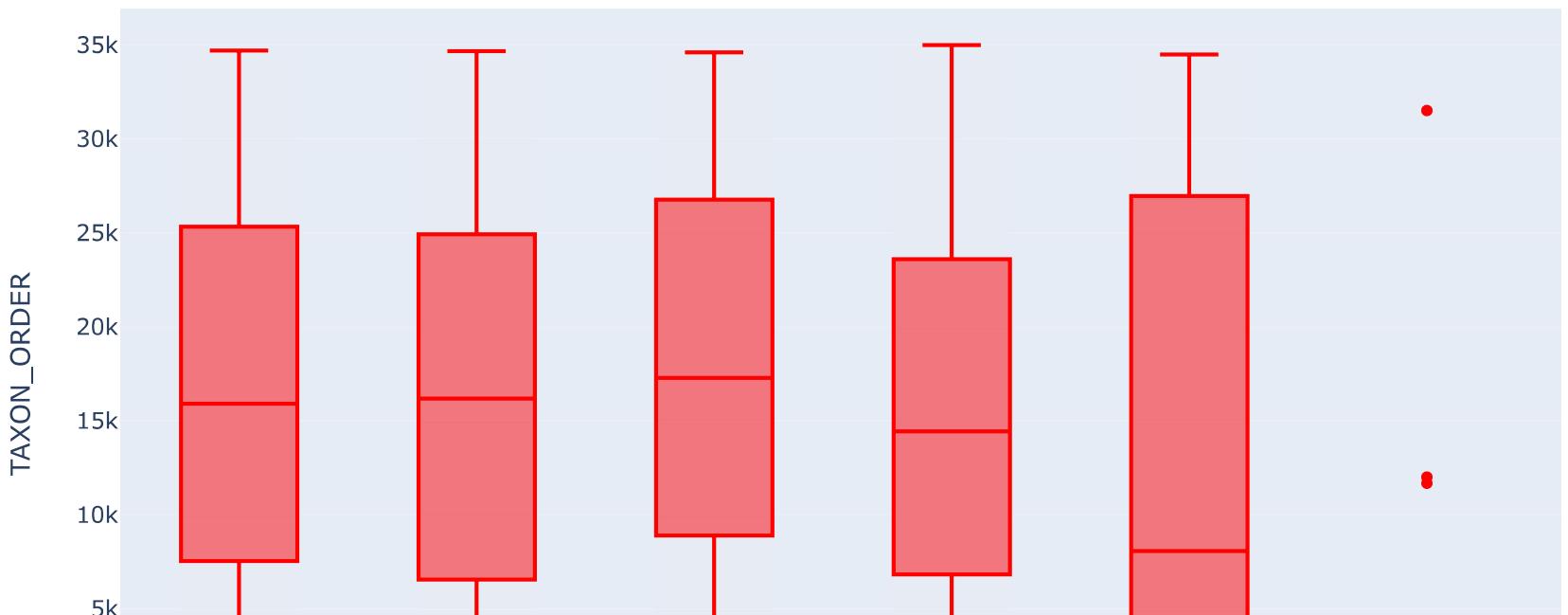
# Scatter plot of the taxonomic order counts by family
fig4 = px.scatter(train_species_df, x="FAMILY", y="TAXON_ORDER")
fig4.update_layout(title_text="Taxonomic Order Distribution by Family")

# Show the plots
fig1.show()
fig3.show()
fig4.show()
```

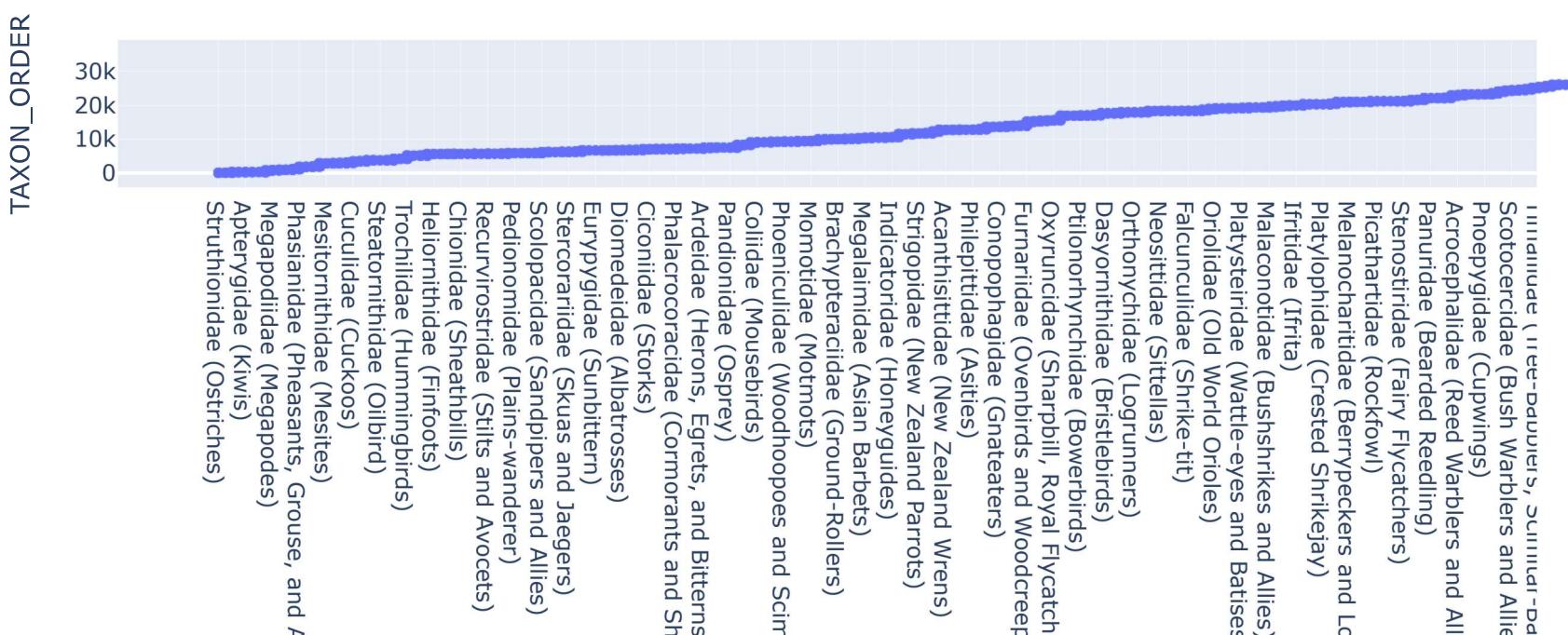
Distribution of Taxonomic Orders



Taxonomic Order Distribution by Category



Taxonomic Order Distribution by Family



AUDIO EXPLORATION

Audio Files : An audio file is a type of digital file format that stores recorded sound or music. It can be played back through speakers or headphones and is commonly used in a variety of applications, such as music, film, television, radio, and other forms of media. Audio files come in many different formats, including **MP3, WAV, OGG, AAC, and FLAC**.

How to Visualize Audio Files ?

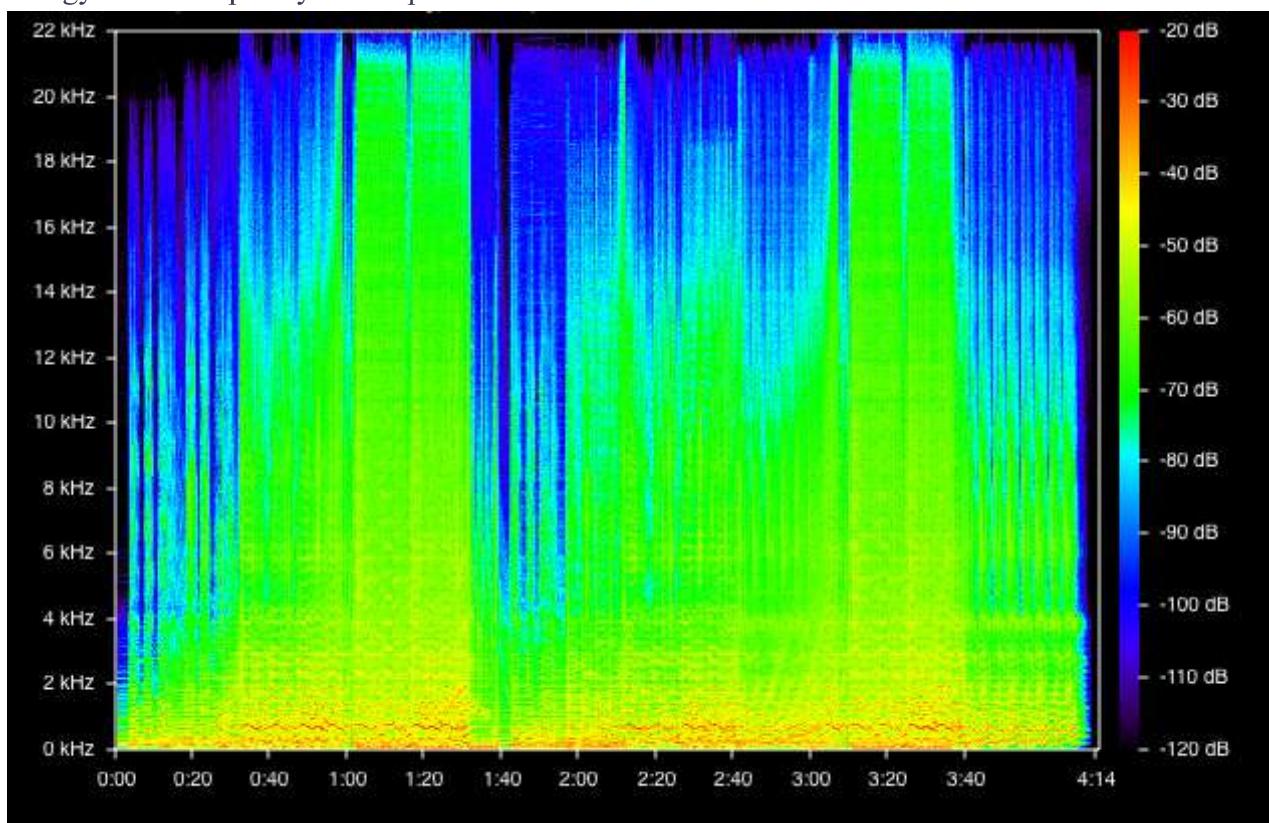
There are many ways in which we can view audio in 2D like :

- 1. Waveforms :** In audio processing, a waveform is a graphical representation of a sound signal that shows how the signal varies over time. It is a plot of the amplitude of the sound wave on the y-axis versus time on the x-axis. Waveforms can be

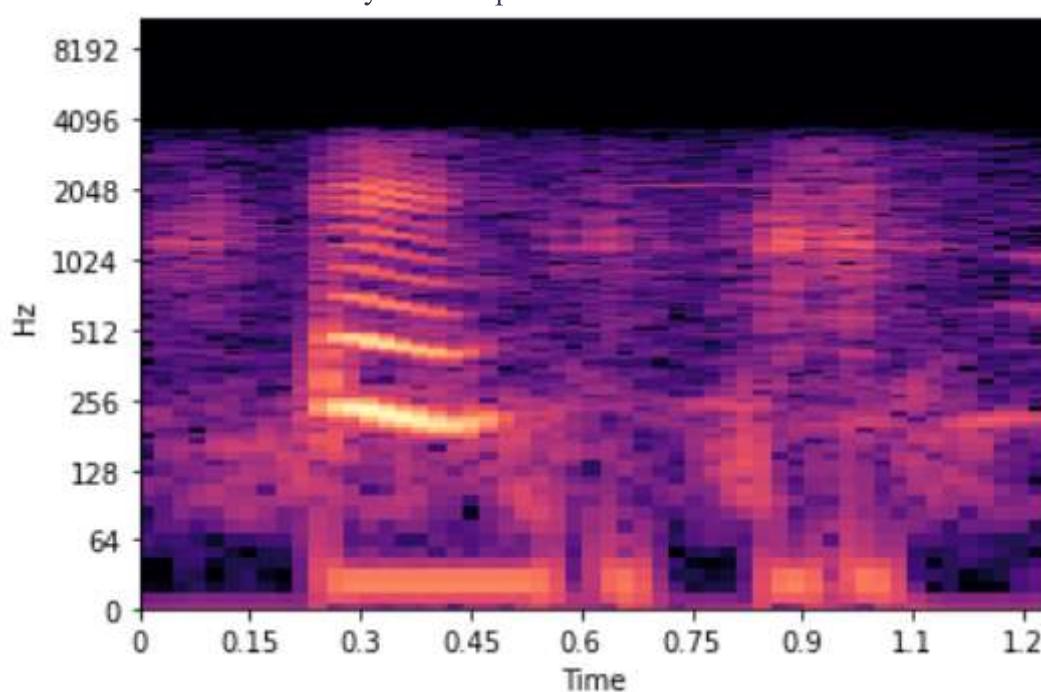
used to visualize and analyze the properties of audio signals, such as frequency, amplitude, phase, and duration.



2. Spectograms : A spectrogram is a visual representation of the spectrum of frequencies of a sound or other signal as it varies with time. It is a way of analyzing the sound signal to understand how the signal changes over time and what frequencies are present in the signal at any given point in time. The x-axis of a spectrogram represents time, while the y-axis represents frequency. The intensity of the color or brightness of each point in the spectrogram represents the amplitude or energy of the frequency at that point in time.

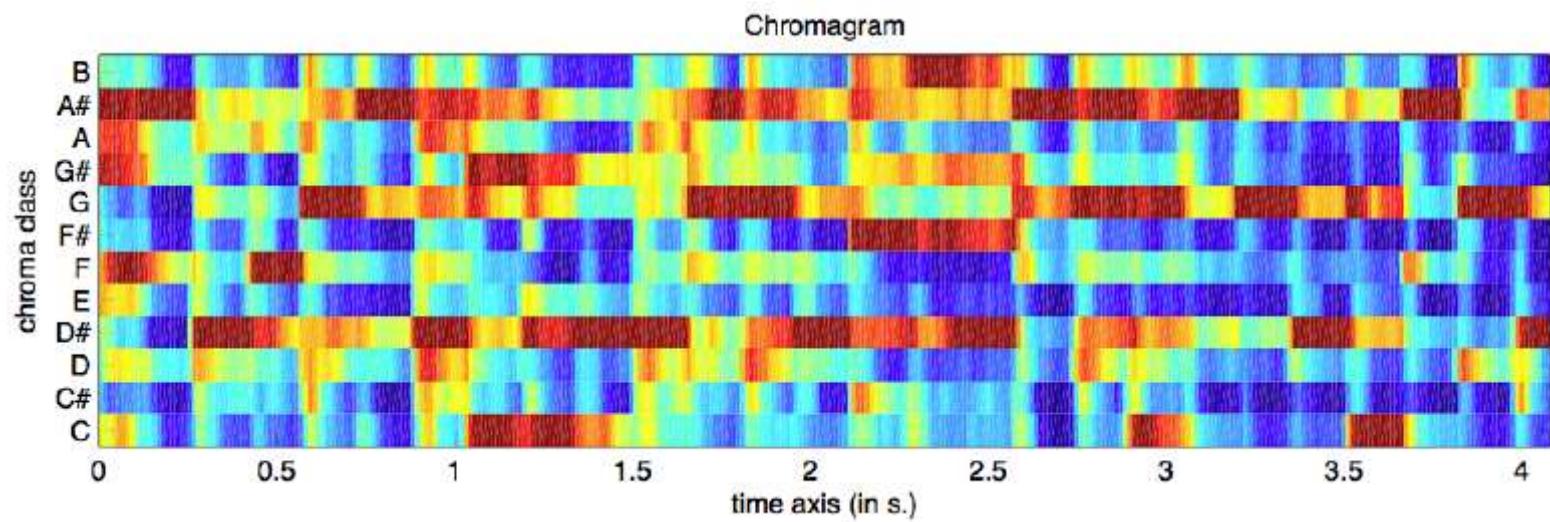


3. Mel Spectograms : Mel spectrograms are a type of spectrogram that use the Mel scale to determine the spacing of frequency bins. The Mel scale is a perceptual scale of pitches judged by listeners to be equal in distance from one another. By using the Mel scale, the frequency bins are spaced closer together at lower frequencies and farther apart at higher frequencies, which better matches the way humans perceive sound.

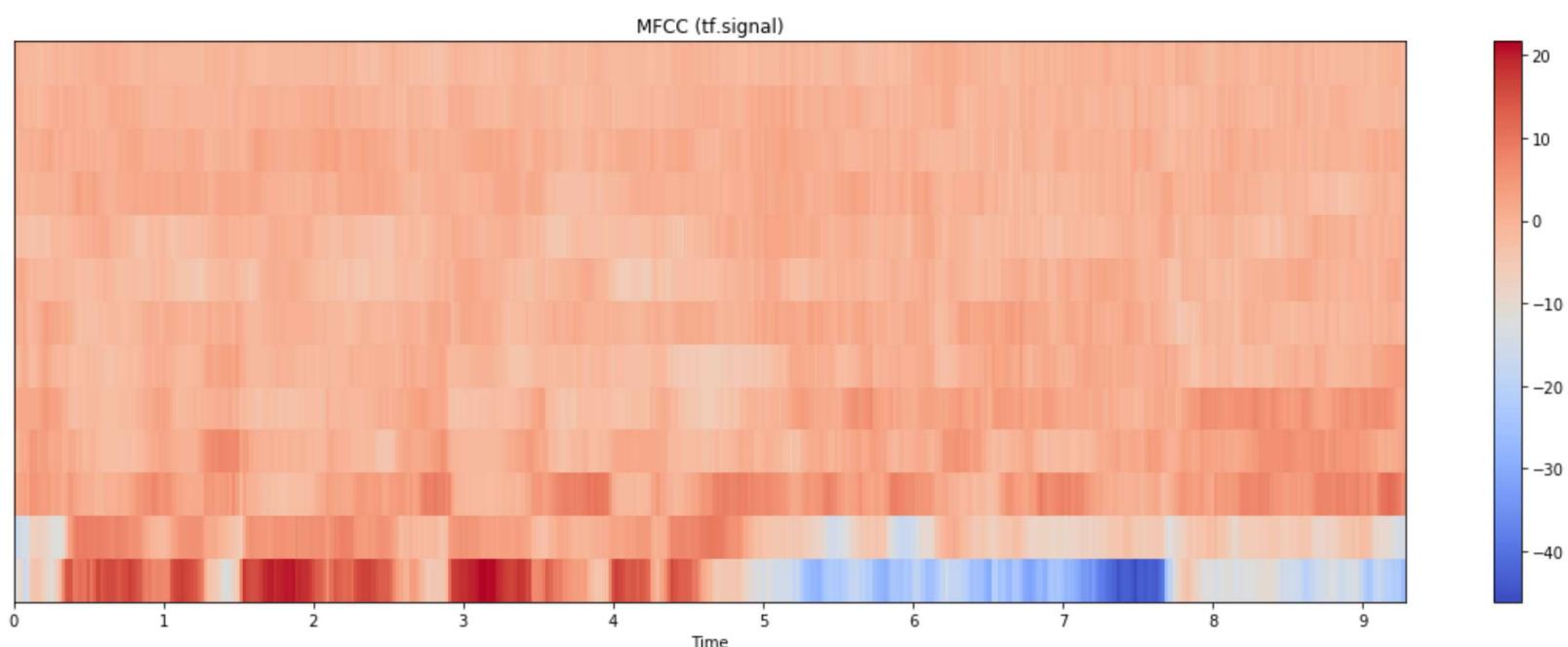


4. Chromagram : A chromagram is a visual representation of the pitch content of an audio signal, where each column corresponds to a particular frequency band or pitch class. It is a 2D representation of the distribution of energy in a sound, plotted as a function of time and pitch class. The chromagram is calculated by taking the short-time Fourier transform (STFT) of the audio signal and projecting the resulting spectrogram onto a pitch-class basis, typically using a bank of 12 log-spaced

filters to represent the 12 pitch classes of the Western musical scale



5. **MFCC** : MFCC stands for Mel Frequency Cepstral Coefficients, which is a feature extraction technique commonly used in speech processing, music information retrieval, and other related fields. MFCCs are derived from a short-time Fourier transform (STFT) of the audio signal, which involves breaking the signal down into small, overlapping segments and performing a Fourier transform on each segment to obtain its frequency spectrum. The Mel scale is then applied to the frequency spectrum to transform it into a logarithmic scale that approximates the human auditory system's perception of sound.



Helper Function

```
In [19]: def audio_eda(audio_path):

    # Load an audio file
    samples, sample_rate = librosa.load(audio_path)

    # Visualize the waveform
    plt.figure(figsize=(14, 5))
    librosa.display.waveform(samples, sr=sample_rate)
    plt.title('Waveform')

    # Compute the spectrogram
    spectrogram = librosa.stft(samples)
    spectrogram_db = librosa.amplitude_to_db(abs(spectrogram))

    # Visualize the spectrogram
    plt.figure(figsize=(14, 5))
    librosa.display.specshow(spectrogram_db, sr=sample_rate, x_axis='time', y_axis='log')
    plt.colorbar(format='%.2f dB')
    plt.title('Spectrogram (dB)')

    # Compute the mel spectrogram

    # Visualize the mel spectrogram
    S = librosa.feature.melspectrogram(y=samples, sr=sample_rate)

    # Visualize mel spectrogram
    plt.figure(figsize=(10, 4))
    librosa.display.specshow(librosa.power_to_db(S, ref=np.max), y_axis='mel', fmax=8000, x_axis='time')
    plt.colorbar(format='%.2f dB')
    plt.title('Mel spectrogram')
    plt.tight_layout()

    # Compute the chromagram

    chromagram = librosa.feature.chroma_stft(y = samples , sr = sample_rate)

    # Visualize the chromagram
    plt.figure(figsize=(14, 5))
    librosa.display.specshow(chromagram, sr=sample_rate, x_axis='time', y_axis='chroma')
    plt.colorbar()
    plt.title('Chromagram')
```

```

# Compute the MFCCs
mfccs = librosa.feature.mfcc(y=samples, sr=sample_rate, n_mfcc=13)

# Visualize the MFCCs
plt.figure(figsize=(14, 5))
librosa.display.specshow(mfccs, sr=sample_rate, x_axis='time')
plt.colorbar()
plt.title('MFCCs')

# Show the plots
display(Audio(samples, rate=sample_rate))
plt.show()

```

Audio Exploration

Black-and-white Mannikin

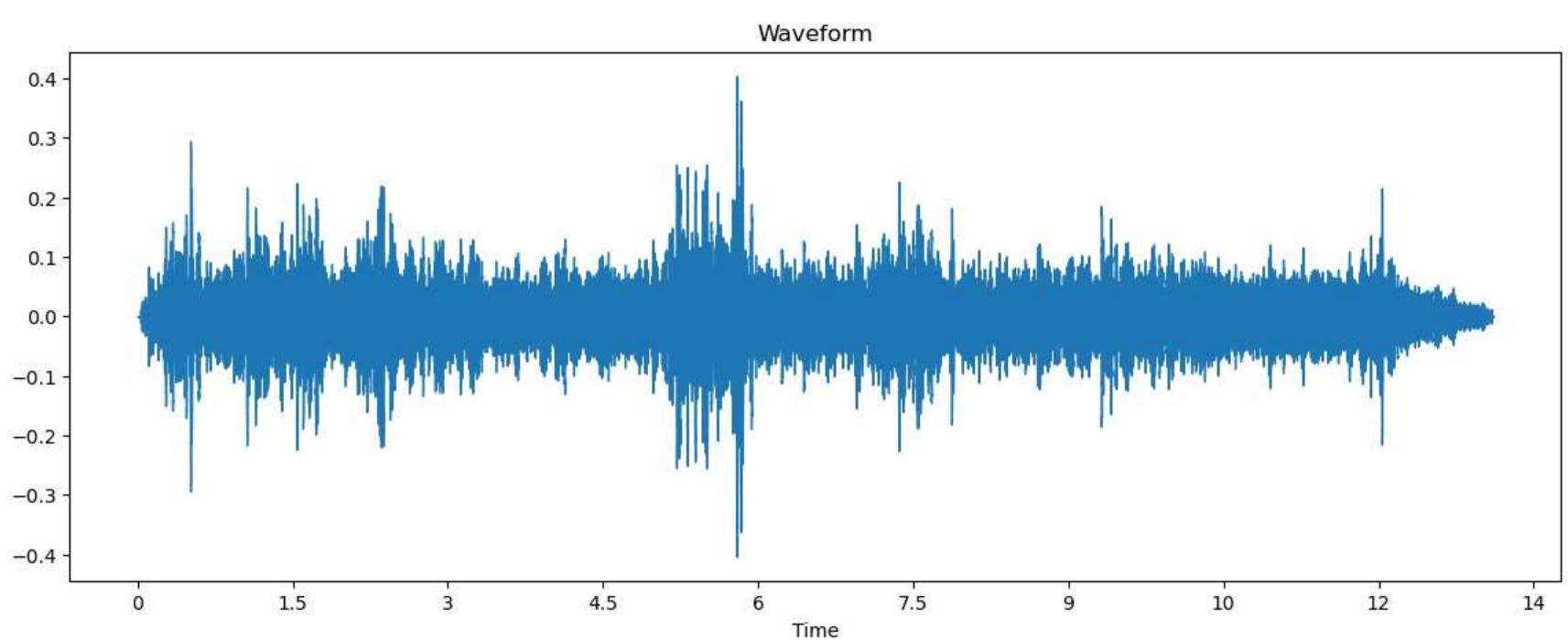


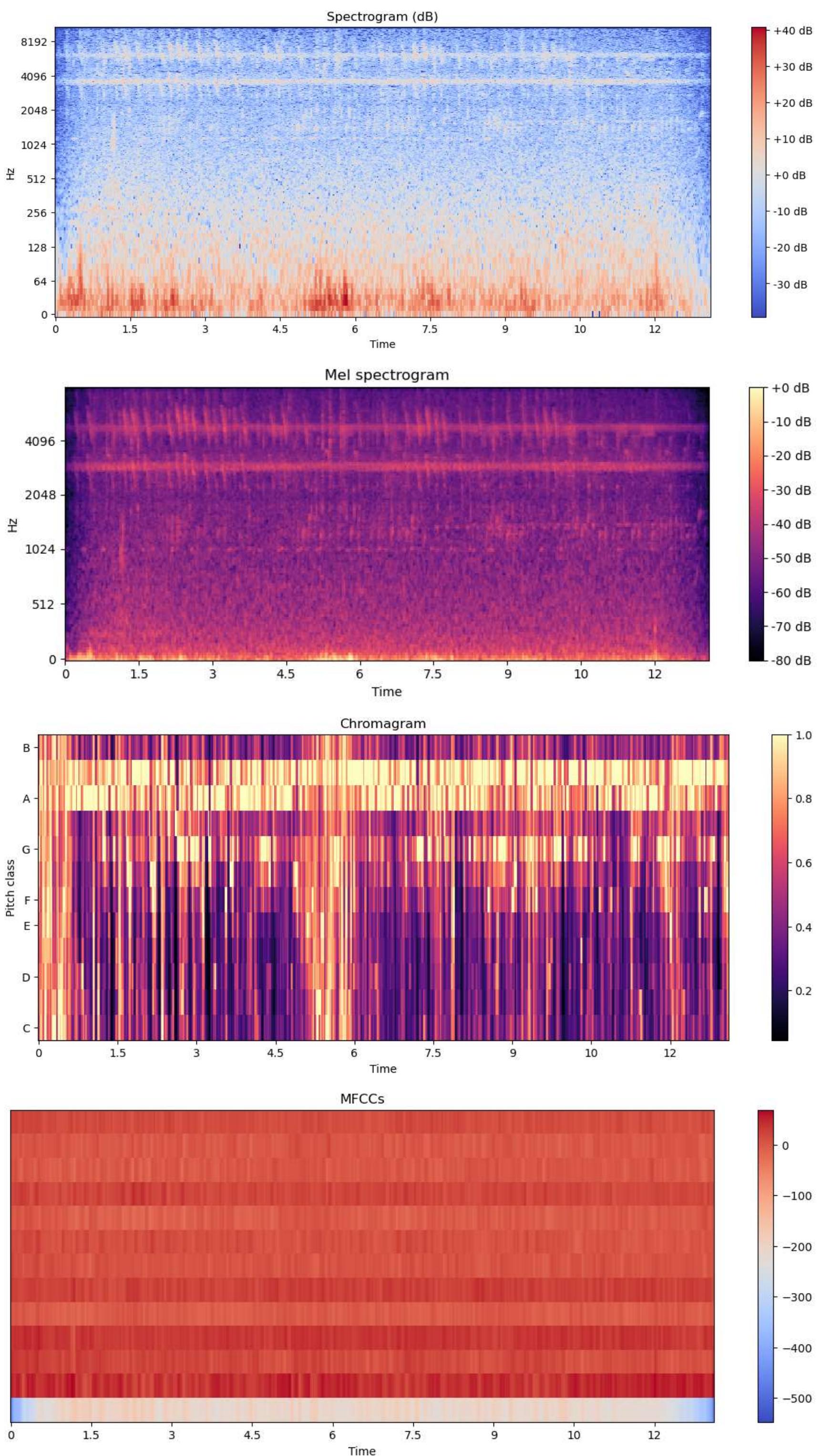
Primary Label : bawman1

Scientific Name : Spermestes bicolor

In [20]: `audio_eda("/input/birdclef-2023/train_audio/bawman1/XC115075.ogg")`

▶ 0:00 / 0:13 ━ ━ ━





African Black-headed Oriole

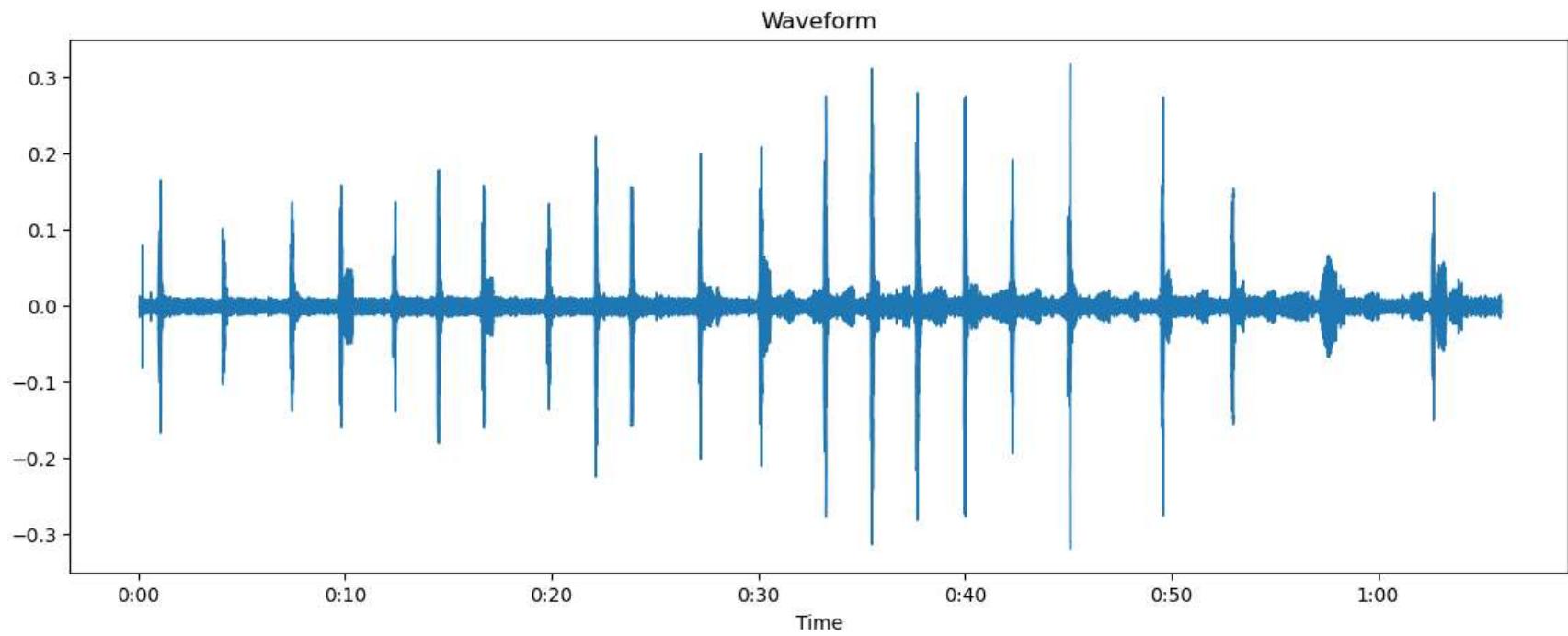


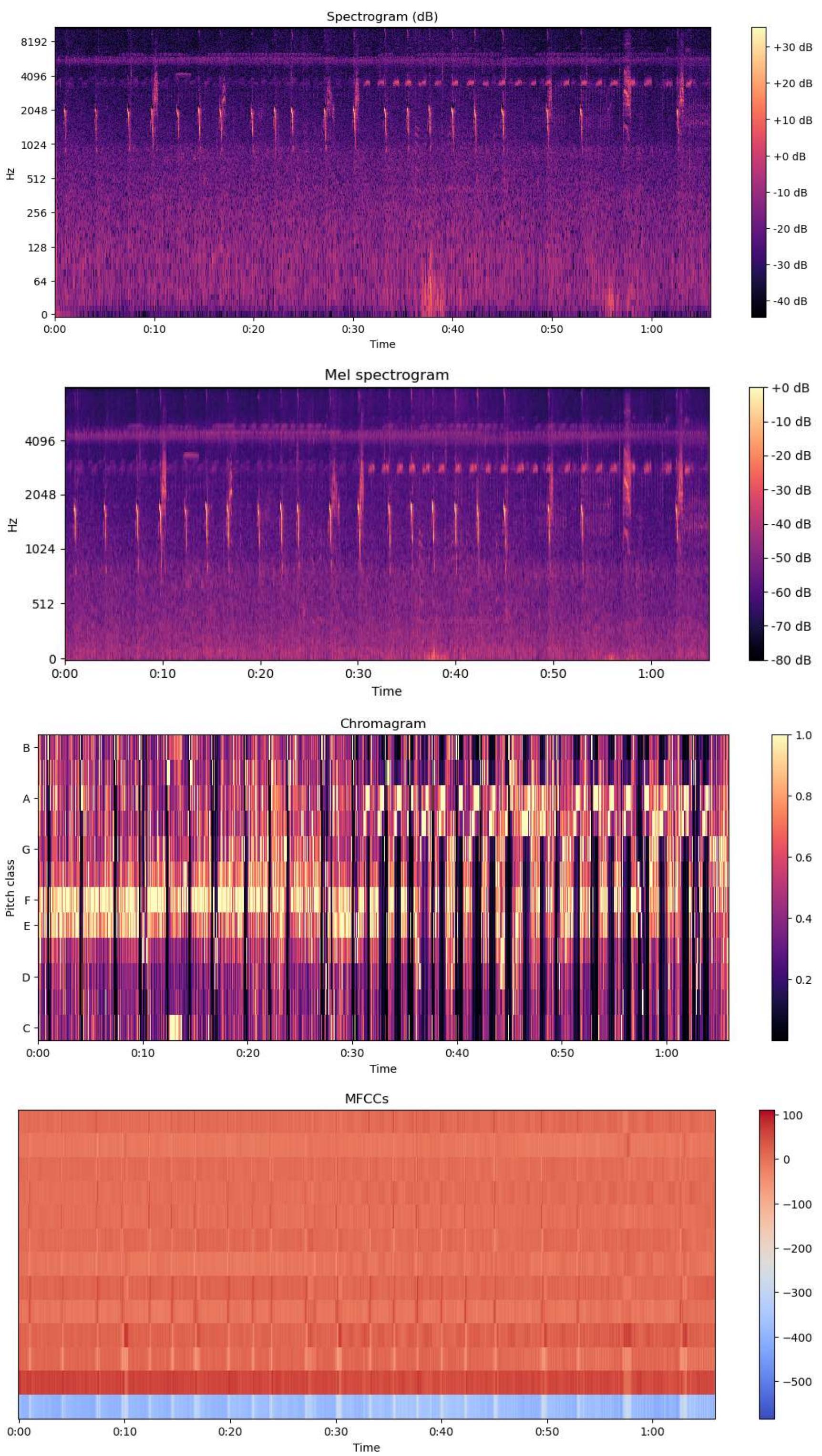
Primary Label : abhori1

Scientific Name : Oriolus larvatus

```
In [21]: audio_eda("/input/birdclef-2023/train_audio/abhori1/XC120251.ogg")
```

▶ 0:00 / 1:05 ━ ━ ━ ━ ━





African Bare-eyed Thrush

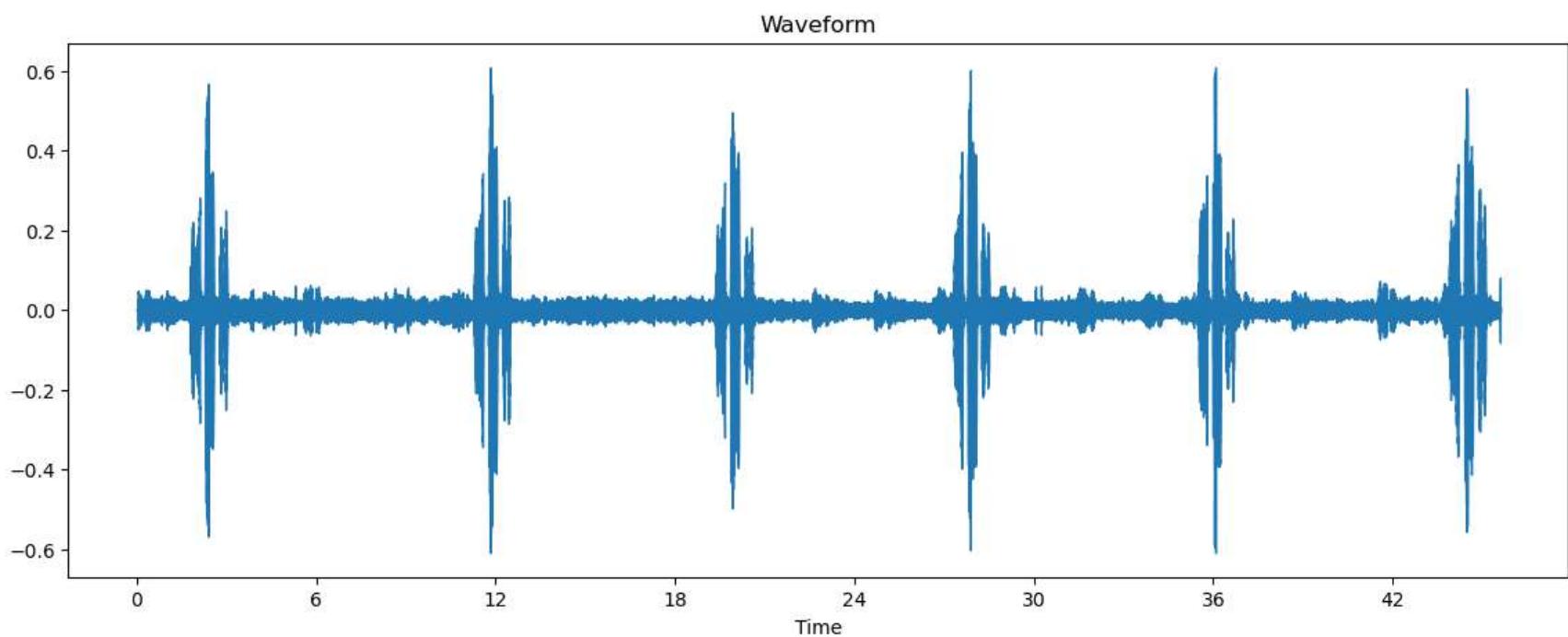


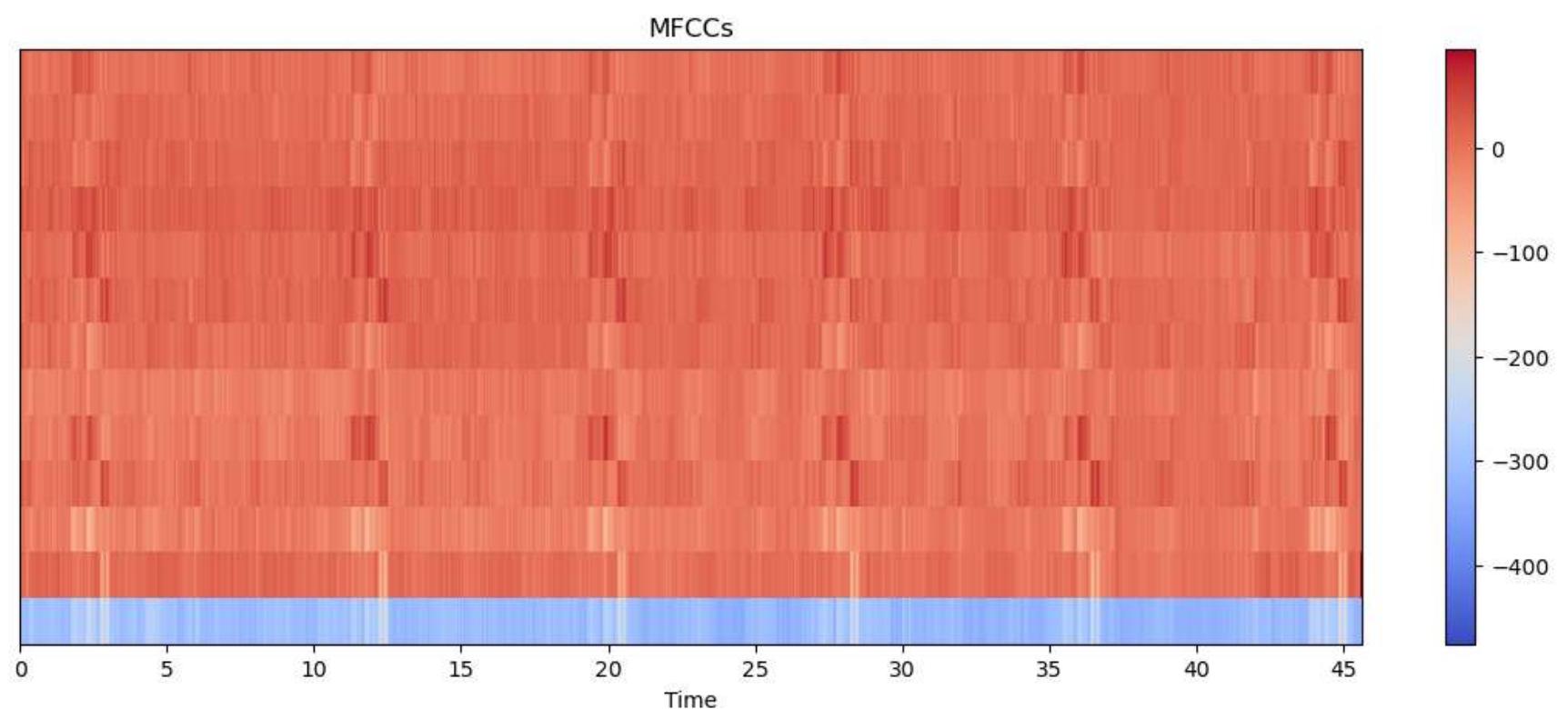
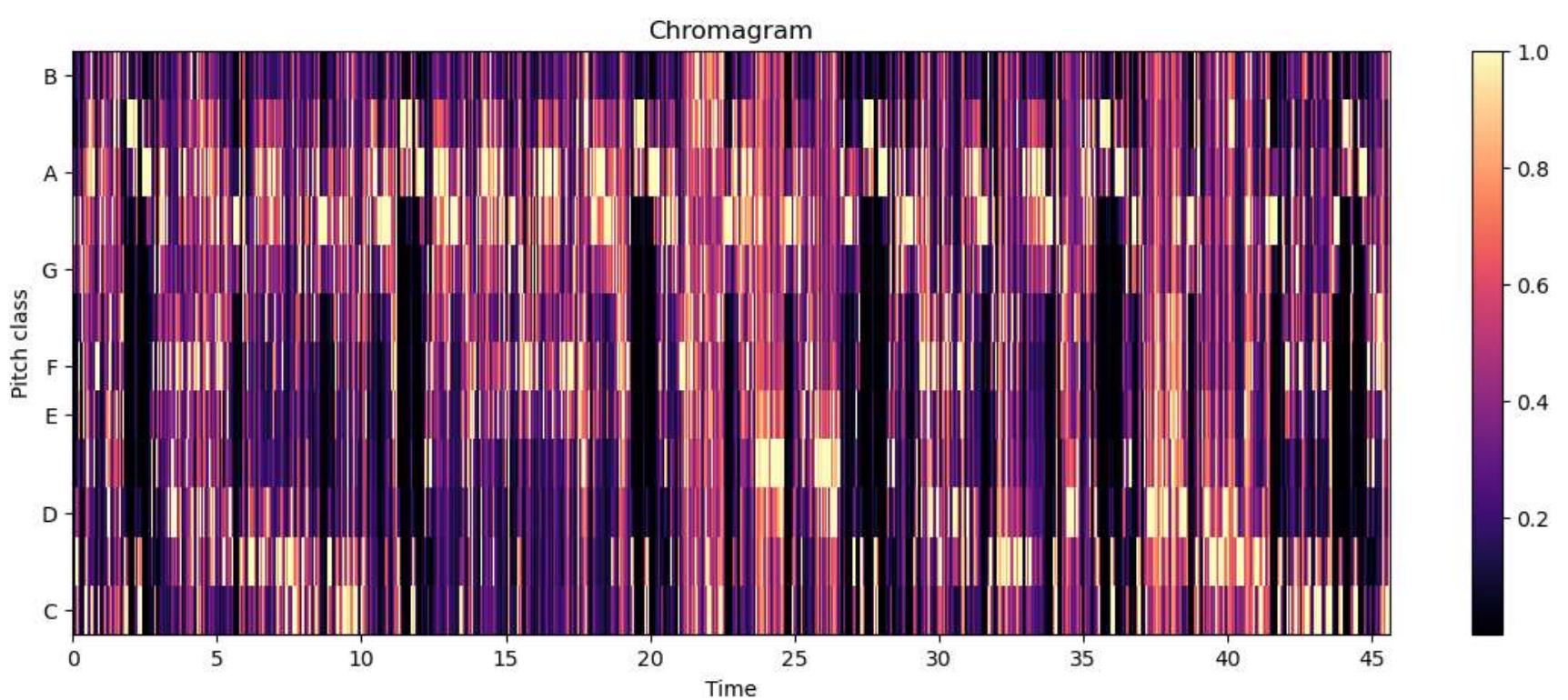
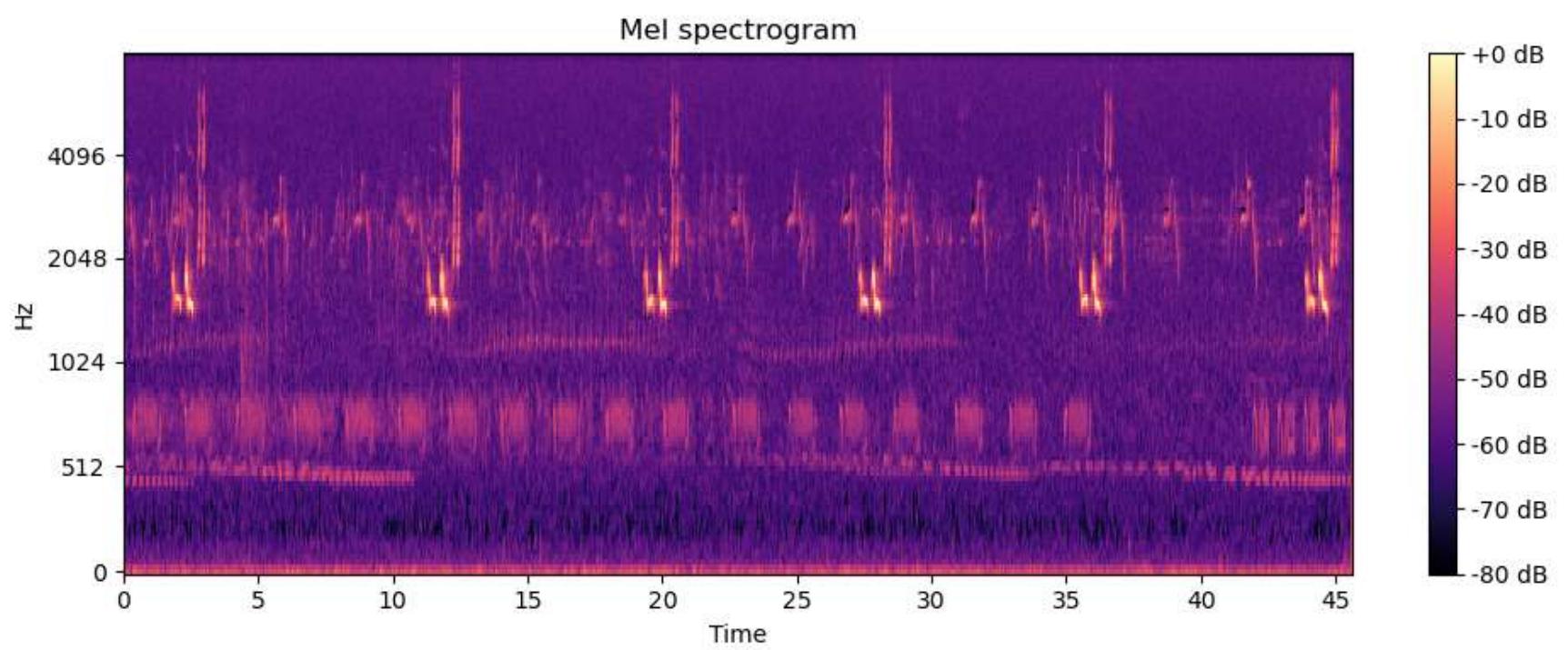
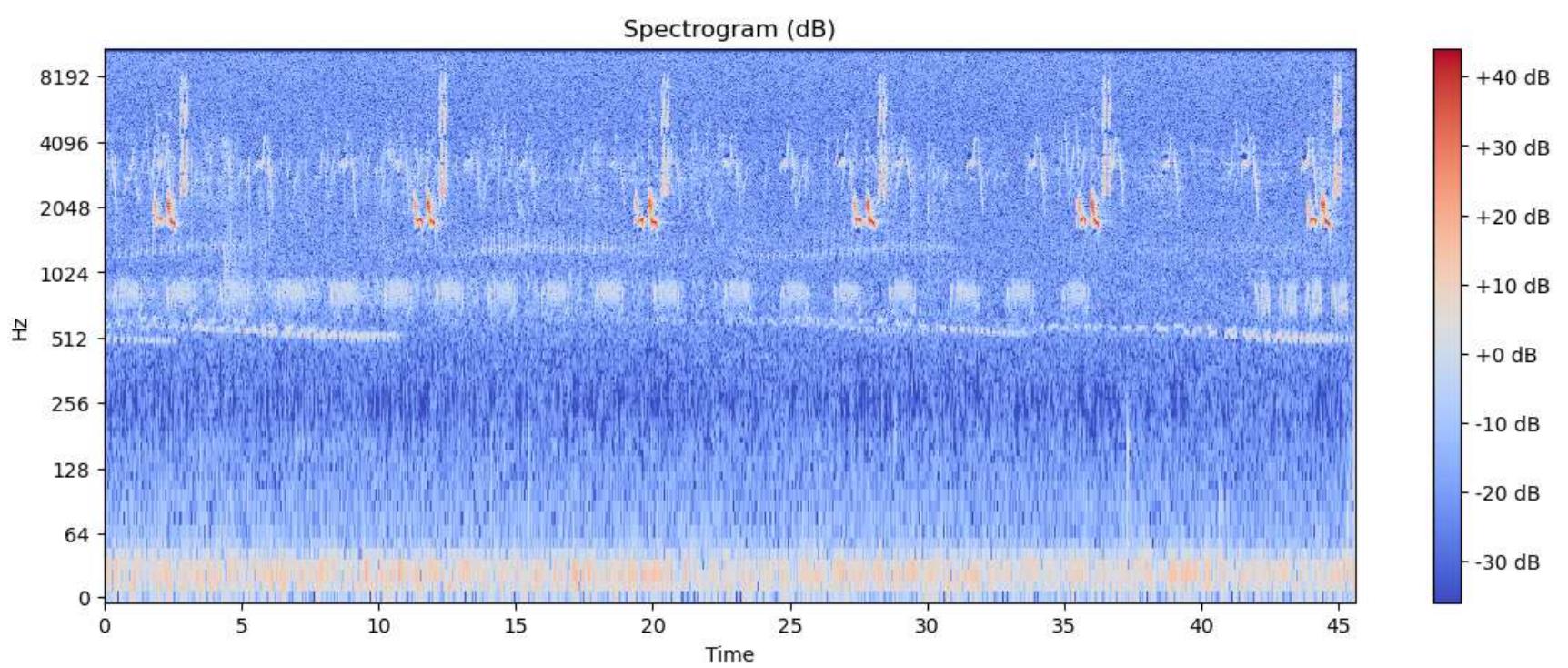
Primary Label : abethr1

Scientific Name : *Turdus tephronotus*

```
In [22]: audio_eda("/input/birdclef-2023/train_audio/abethr1/XC128013.ogg")
```

▶ 0:00 / 0:45 ━ ━ ━





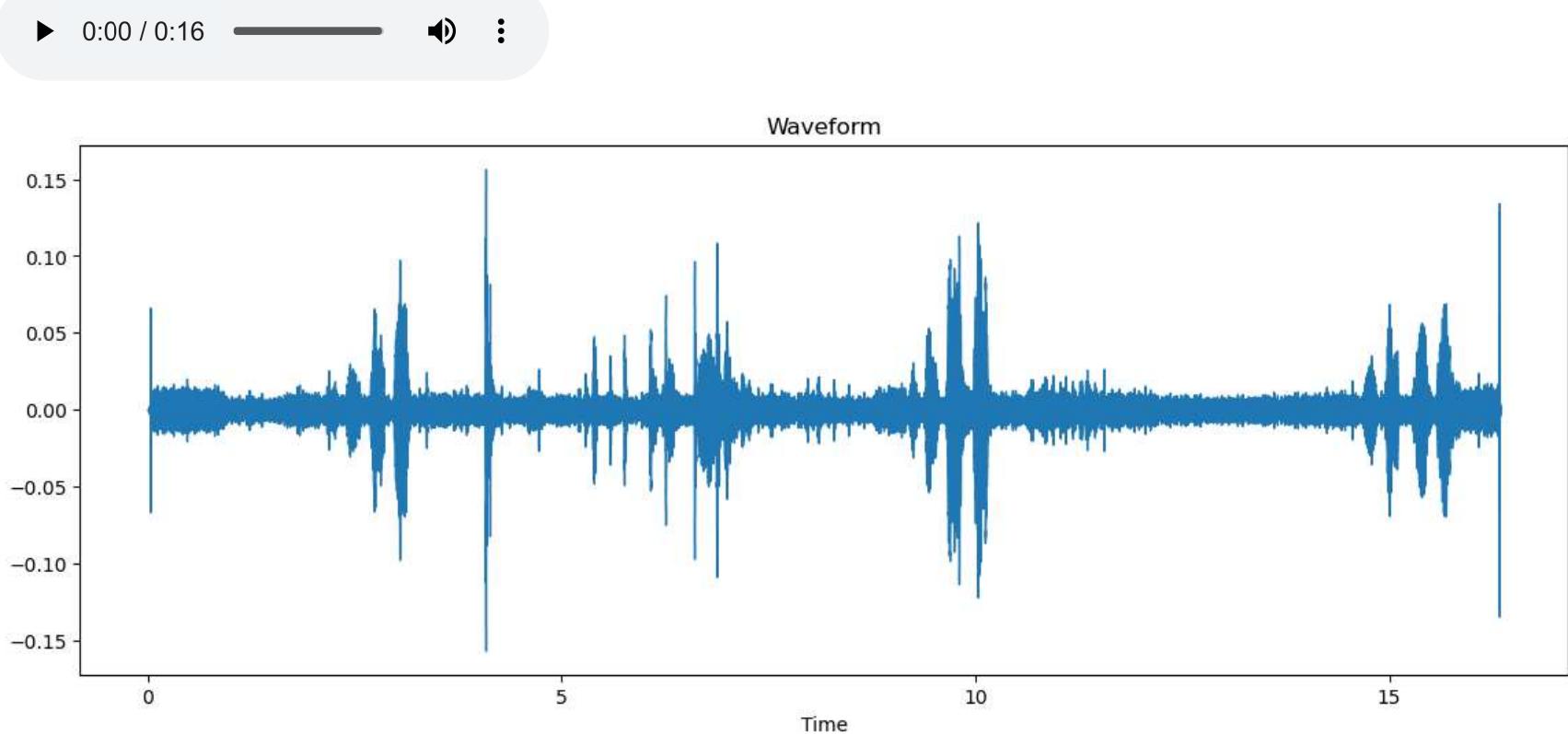
African Gray Flycatcher

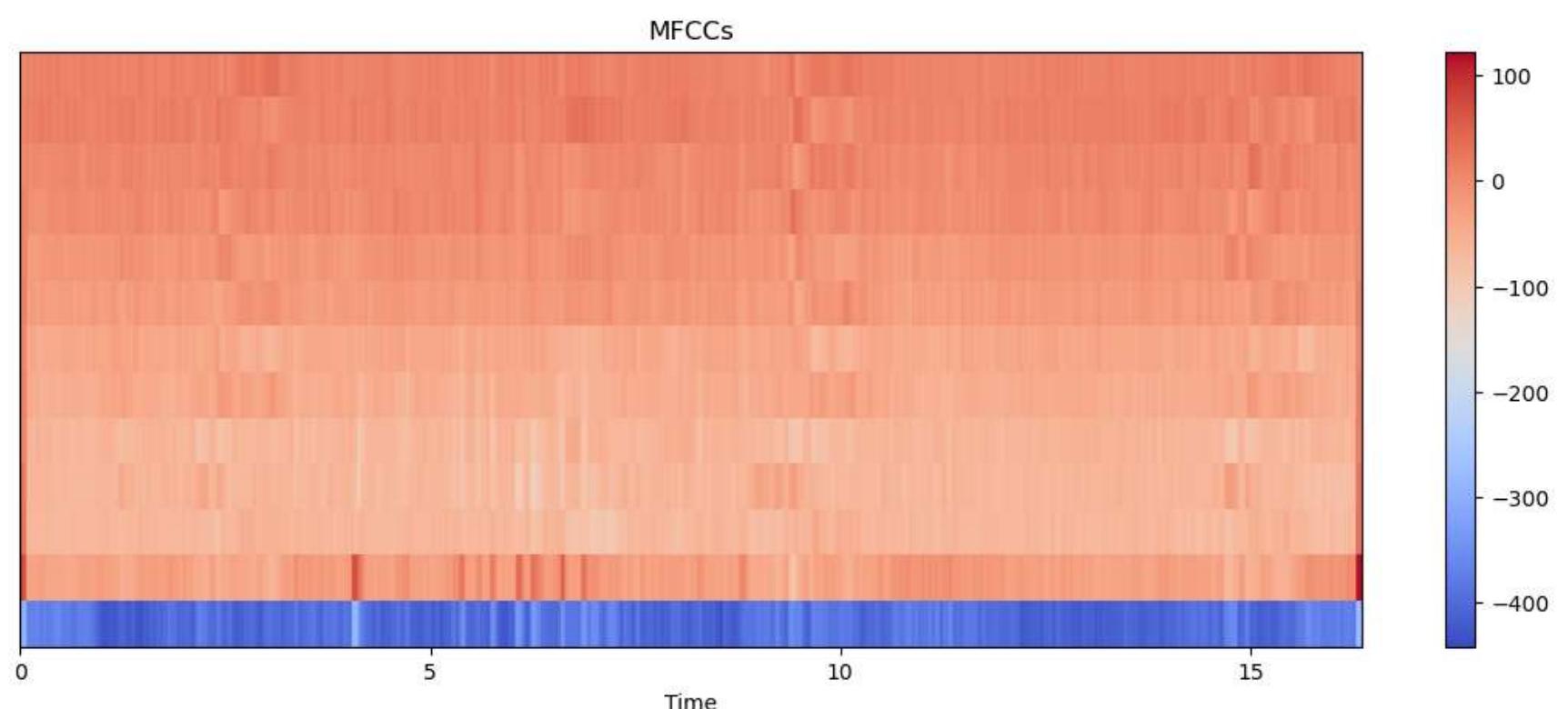
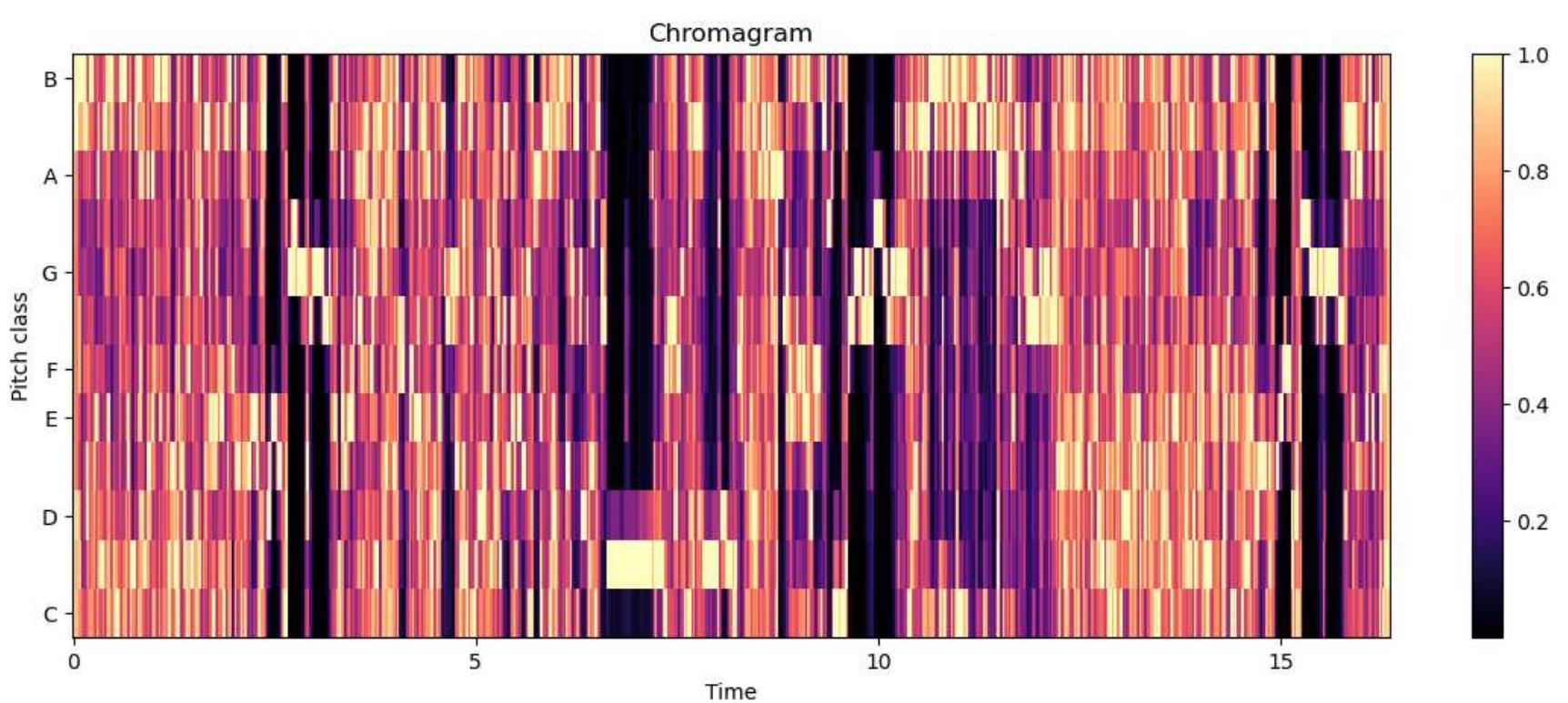
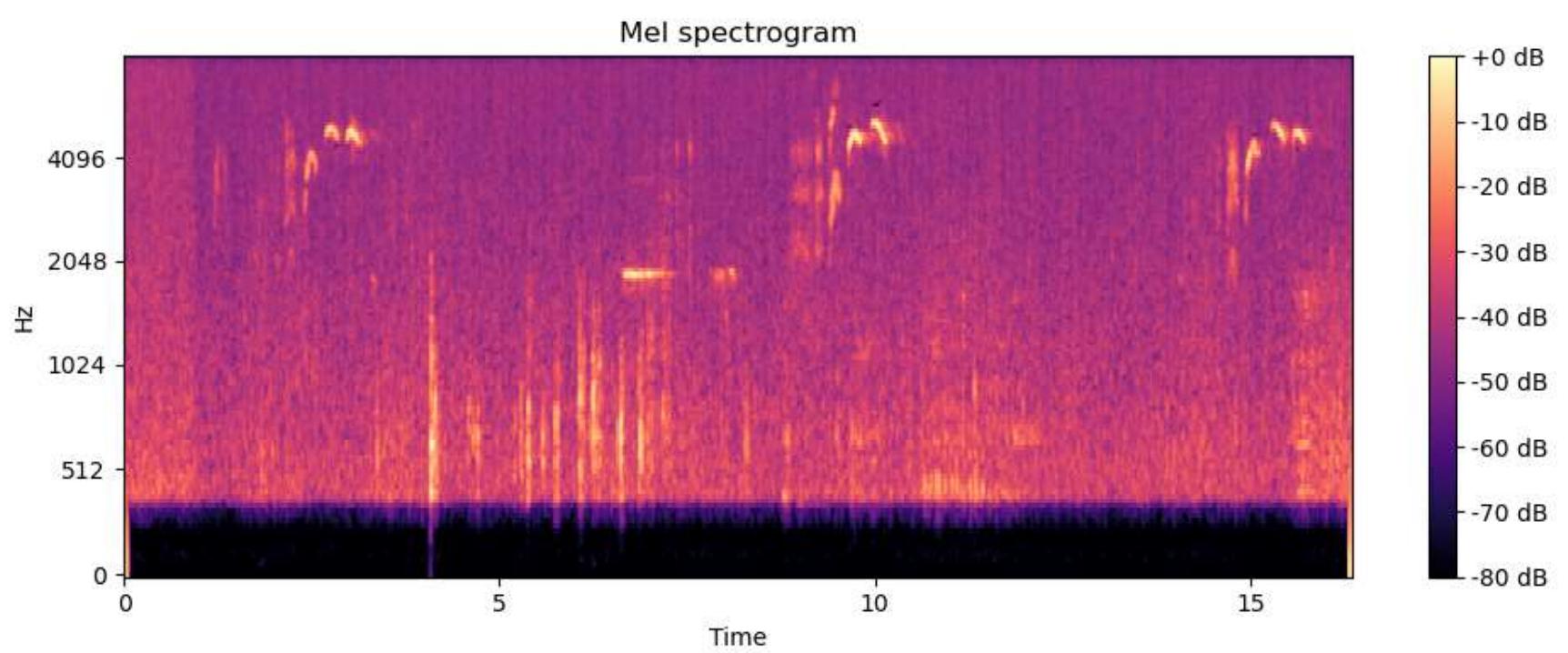
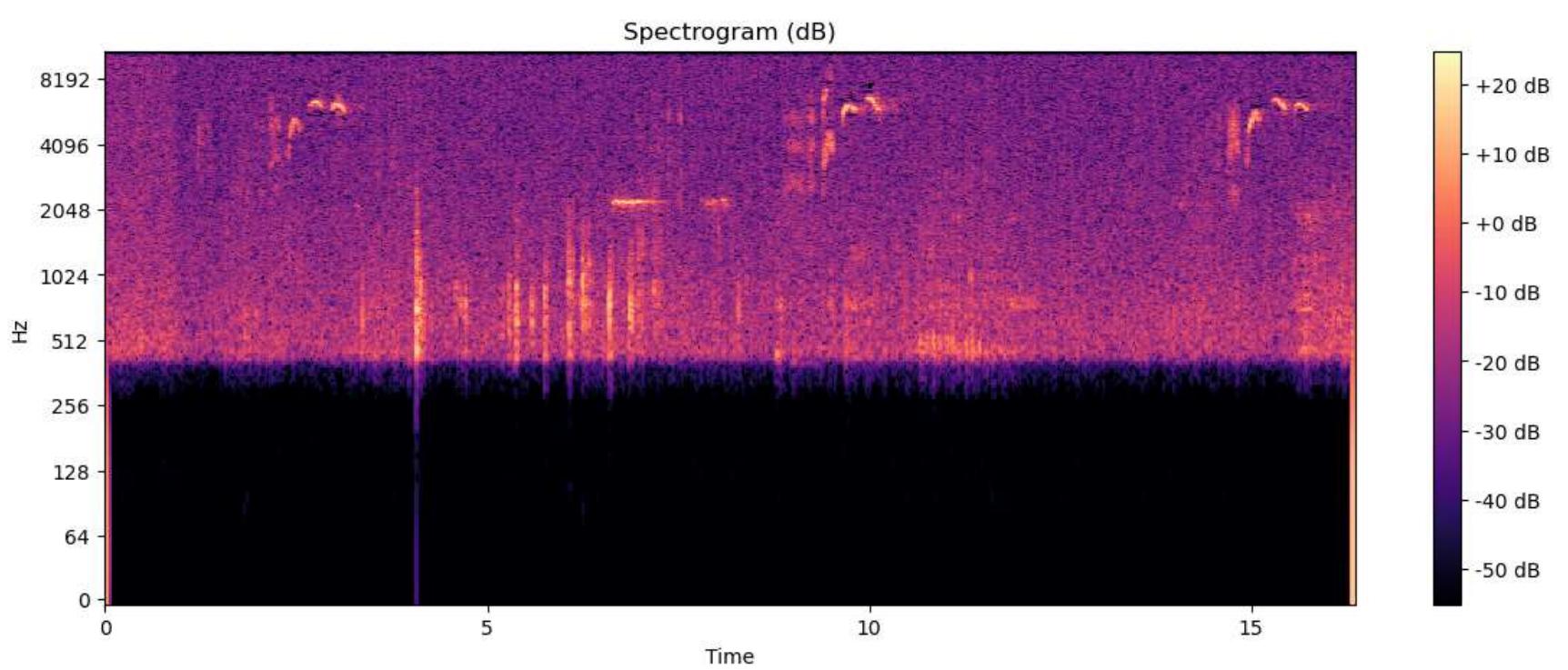


Primary Label : afgfly1

Scientific Name : Bradornis microrhynchus

```
In [23]: audio_eda("/input/birdclef-2023/train_audio/afgfly1/XC134487.ogg")
```





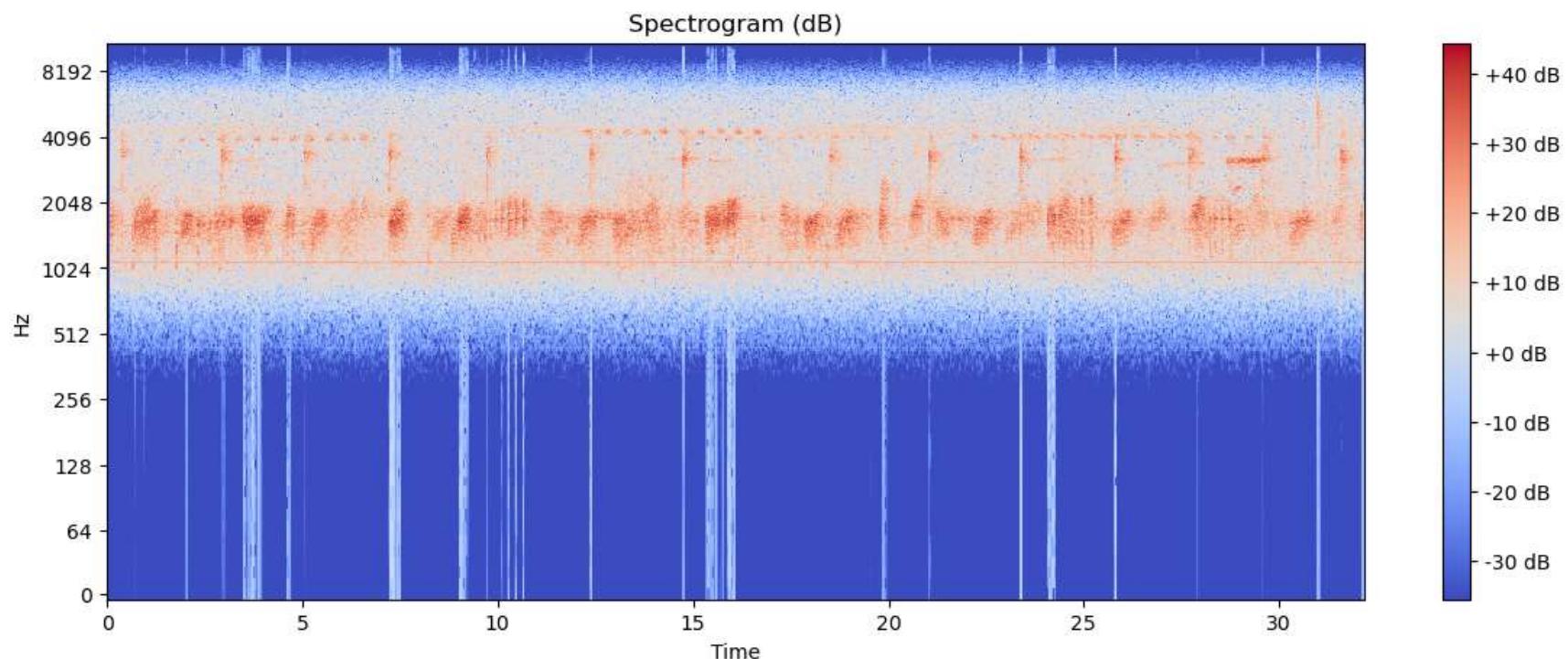
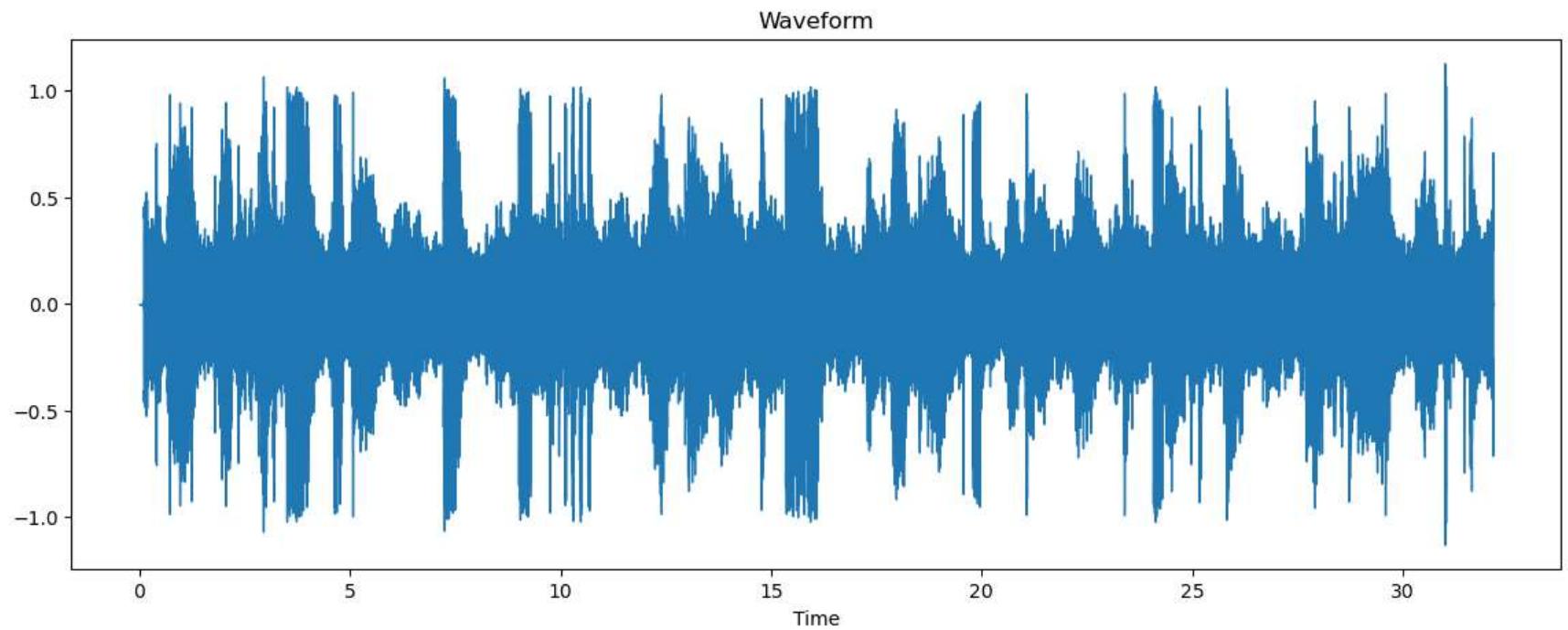
African Goshawk



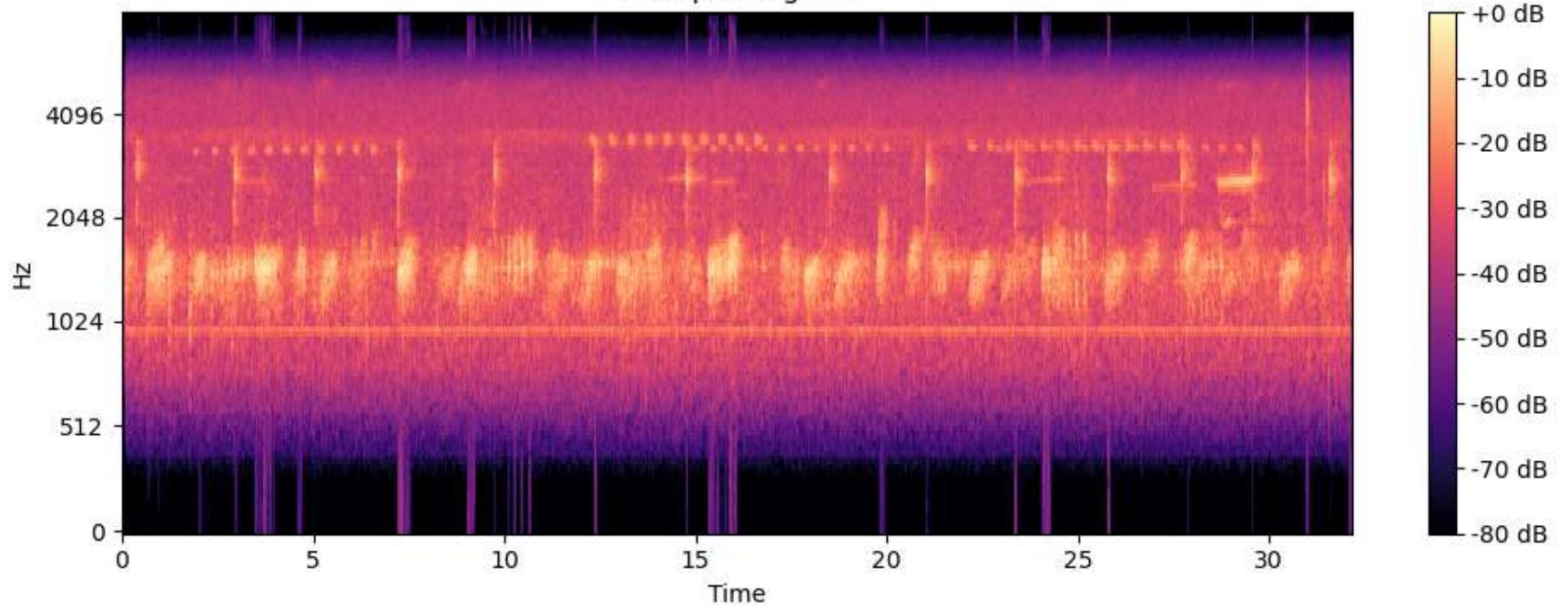
Primary Label : afrgos1

Scientific Name : Accipiter tachiro

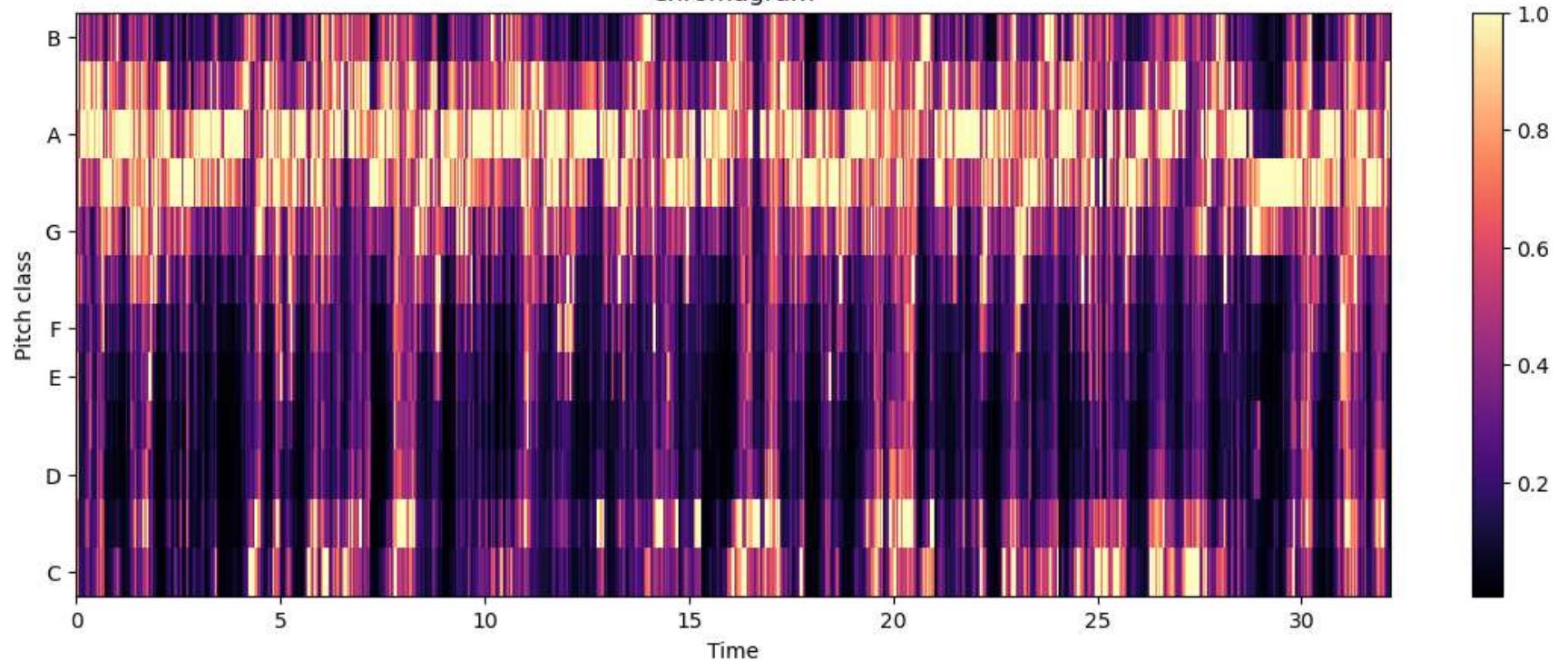
```
In [24]: audio_eda("/input/birdclef-2023/train_audio/afrgos1/XC115984.ogg")
```



Mel spectrogram



Chromagram



MFCCs

