

# Transfer Learning for Identification of Rare Bird Species

## I. Installing and Importing Libraries

```
# --- Installing and importing libraries---
```

```
# Installing libraries
#!pip install birdnetlib
#!pip install librosa==0.9.2
#!pip install opensoundscape
#!pip install git+https://github.com/kitzeslab/bioacoustics-model-zoo
#!pip install tensorflow tensorflow-hub
#!pip install pydub
#!pip install joblib
```

```
# Import packages
import os
import bioacoustics_model_zoo as bmz
import joblib
```

```
from pydub import AudioSegment
from pydub.utils import mediainfo
from google.colab import drive
from bioacoustics_model_zoo import BirdNET
from birdnetlib import Recording
from birdnetlib.analyzer import Analyzer
from datetime import datetime
from collections import defaultdict
from collections import Counter
```

```
from google.colab import drive
```

```
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import classification_report, accuracy_score, f1_score
from sklearn.model_selection import train_test_split
from sklearn.model_selection import StratifiedGroupKFold
from sklearn.model_selection import StratifiedKFold
from sklearn.neighbors import KNeighborsClassifier
from sklearn.neural_network import MLPClassifier
from sklearn.preprocessing import StandardScaler
from sklearn.metrics.pairwise import cosine_similarity
```

```
↳ <frozen importlib._bootstrap>:1047: ImportWarning: _PyDrive2ImportHook.find_spec()
  <frozen importlib._bootstrap>:1047: ImportWarning: _PyDriveImportHook.find_spec()
  <frozen importlib._bootstrap>:1047: ImportWarning: _BokehImportHook.find_spec() no
  <frozen importlib._bootstrap>:1047: ImportWarning: _PyDrive2ImportHook.find_spec()
  . . . . .
```

```

<frozen importlib._bootstrap>:1047: ImportWarning: _PyDriveImportHook.find_spec()
<frozen importlib._bootstrap>:1047: ImportWarning: _BokehImportHook.find_spec() no
<frozen importlib._bootstrap>:1047: ImportWarning: _PyDrive2ImportHook.find_spec()
<frozen importlib._bootstrap>:1047: ImportWarning: _PyDriveImportHook.find_spec()
<frozen importlib._bootstrap>:1047: ImportWarning: _BokehImportHook.find_spec() no

```

## ▼ II. Pre-Processing Audio Files

```

# --- Transform audio to standard WAV and padding to 3 seconds ---

# Mount Google Drive
drive.mount('/content/drive')

# Input folders (original audio)
input_folders = {
    ... "NEG": "/content/drive/My Drive/AlternativeBirds/",
    ... "POS": "/content/drive/My Drive/OnlyAudioDoliornis/",
    ... "OPOS": "/content/drive/My Drive/OnlyAudioHapalopsittaca/"
}

# Output folders (converted WAV files)
output_folders = {
    ... "NEG": "/content/drive/My Drive/AlternativeBirds_wav/",
    ... "POS": "/content/drive/My Drive/OnlyAudioDoliornis_wav/",
    ... "OPOS": "/content/drive/My Drive/OnlyAudioHapalopsittaca_wav/"
}

# Function to convert and pad audio
def convert_and_pad_to_3s(input_path, output_path):
    ... try:
    ...     audio = AudioSegment.from_file(input_path)
    ...     duration_ms = 3000 # 3 seconds

    ...     if len(audio) < duration_ms:
    ...         silence = AudioSegment.silent(duration=duration_ms - len(audio))
    ...         audio = audio + silence

    ...     # Let pydub handle the writing
    ...     audio.export(output_path, format="wav")

    ...     print(f"Processed and saved: {output_path}")
    ... except Exception as e:
    ...     print(f"Error processing {input_path}: {e}")

# Process each folder
for key in input_folders:
    ... input_dir = input_folders[key]
    ... output_dir = output_folders[key]
    ... os.makedirs(output_dir, exist_ok=True)

    ... audio_files = [
    ...     os.path.join(input_dir, f)
    ...     for f in os.listdir(input_dir)

```

```

    .....if f.lower().endswith(('.mp3', '.wav', '.flac', '.ogg'))
    ....]

    ....for input_path in audio_files:
    .....filename = os.path.splitext(os.path.basename(input_path))[0] + ".wav"
    .....output_path = os.path.join(output_dir, filename)
    .....convert_and_pad_to_3s(input_path, output_path)

```

```

Drive already mounted at /content/drive; to attempt to forcibly remount, call driv
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC187 - Grey-bre
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC2069 - Plum-cr
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC3470 - Blue-ba
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC2019 - Plum-cr
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC2012 - Jameson
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC1598 - Blue-ba
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC20706 - Blue-t
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC14584 - Grey-b
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC62895 - Black
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC178 - Jameson_
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/AlternativeBirds_wav/XC2915 - Andean
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/OnlyAudioDoliornis_wav/XC10781 - Bay-
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/OnlyAudioDoliornis_wav/XC20615 - Bay-
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/OnlyAudioDoliornis_wav/XC41676 - Bay-
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/OnlyAudioDoliornis_wav/XC65640 - Bay-
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
    audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/OnlyAudioDoliornis_wav/XC142403 - Bay

```

```

<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/OnlyAudioDoliornis_wav/XC222460 - Bay
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/OnlyAudioDoliornis_wav/XC222458 - Bay
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
audio.export(output_path, format="wav")
Processed and saved: /content/drive/My Drive/OnlyAudioDoliornis_wav/XC222459 - Bay

```

### ✓ III. Verifying if BirdNET Identifies our Rare Species

```
# --- Check if BirdNET identifies our rare species ---
```

```

def analyze_audio_folder(folder_path):
    analyzer = Analyzer() # Load BirdNET model

    audio_files = [
        f for f in os.listdir(folder_path)
        if f.lower().endswith((".mp3", ".wav"))
    ]

    for filename in audio_files:
        file_path = os.path.join(folder_path, filename)
        print(f"\nAnalyzing:")

        try:
            recording = Recording(analyzer, file_path)
            recording.analyze()

            detections = recording.detections

            if detections:
                print("Detections:")
                for d in detections:
                    print(f" - {d['common_name']} ({d['confidence']:.2f})")
            else:
                print("No detections found.")

        except Exception as e:
            print(f"Error analyzing {filename}: {e}")

# For Doliornis
print("Identification results corresponding to Doliornis:")
analyze_audio_folder(input_folders["POS"])

# For Hapalopsittaca
print("\nIdentification results corresponding to Hapalopsittaca:")
analyze_audio_folder(input_folders["OPOS"])

```

```

Identification results corresponding to Doliornis:
Labels loaded.
, , , ,

```

```
load model true
Model loaded.
Labels loaded.
load_species_list_model
Meta model loaded.
```

Analyzing:

```
read_audio_data
```

```
<frozen importlib._bootstrap>:1047: ImportWarning: _PyDrive2ImportHook.find_spec()
<frozen importlib._bootstrap>:1047: ImportWarning: _PyDriveImportHook.find_spec()
<frozen importlib._bootstrap>:1047: ImportWarning: _BokehImportHook.find_spec() no
<frozen importlib._bootstrap>:1047: ImportWarning: _PyDrive2ImportHook.find_spec()
<frozen importlib._bootstrap>:1047: ImportWarning: _PyDriveImportHook.find_spec()
<frozen importlib._bootstrap>:1047: ImportWarning: _BokehImportHook.find_spec() no
```

```
read_audio_data: complete, read 1 chunks.
```

```
analyze_recording XC10781 - Bay-vented Cotinga - Doliornis sclateri.mp3
```

```
No detections found.
```

Analyzing:

```
read_audio_data
```

```
read_audio_data: complete, read 9 chunks.
```

```
analyze_recording XC20615 - Bay-vented Cotinga - Doliornis sclateri.mp3
```

```
Detections:
```

- Dot-winged Antwren (0.50)
- Bertoni's Antbird (0.19)
- White-bellied Antbird (0.21)
- Human non-vocal (0.34)
- American Bullfrog (0.16)

Analyzing:

```
read_audio_data
```

```
read_audio_data: complete, read 40 chunks.
```

```
analyze_recording XC41676 - Bay-vented Cotinga - Doliornis sclateri.mp3
```

```
Detections:
```

- Western Rock Nuthatch (0.13)
- Gray Flycatcher (0.12)
- Turquoise Jay (0.26)
- White-throated Tyrannulet (0.11)
- Red-necked Phalarope (0.11)
- White-throated Tyrannulet (0.51)
- White-throated Tyrannulet (0.22)
- Barred Fruiteater (0.48)
- Turquoise Jay (0.20)
- Tschudi's Tapaculo (0.46)
- Tschudi's Tapaculo (0.64)
- Unicolored Tapaculo (0.42)
- Citrine Warbler (0.21)
- European Goldfinch (0.18)
- White-throated Tyrannulet (0.49)
- White-throated Tyrannulet (0.86)
- White-throated Tyrannulet (0.12)

Analyzing:

```
read_audio_data
```

```
read_audio_data: complete, read 41 chunks.
```

```
analyze_recording XC65640 - Bay-vented Cotinga - Doliornis sclateri.mp3
```

RESULT: BirdNET does not identify our rare species (Doliornis sclateri or Hapalopsittaca melanotis)

```
melanotis)
```

## ▼ IV. Obtaining Embeddings from BirdNET for Transfer Learning

```
# --- Get the embeddings of our rare species from BirdNET for transfer learning ---

# Initialize BirdNET model
model = bmz.BirdNET()

# Create function for extracting embeddings
import numpy as np
def extract_embeddings(audio_folder, output_folder):
    ... os.makedirs(output_folder, exist_ok=True)

    ... audio_files = [
    ...     os.path.join(audio_folder, f)
    ...     for f in os.listdir(audio_folder)
    ...     if f.lower().endswith(('.mp3', '.wav'))
    ... ]

    ... if not audio_files:
    ...     print(f"No audio files found in {audio_folder}")
    ...     return

    ... print(f"Embedding {len(audio_files)} files from: {audio_folder}")

    ... # Run embedding
    ... embeddings_df = model.embed(audio_files)

    ... # Group by filename (from MultiIndex)
    ... grouped = embeddings_df.groupby(level=0)

    ... for file_name, group in grouped:
    ...     emb_array = group.to_numpy() # shape: (n_chunks, 1024)
    ...     base_name = os.path.splitext(os.path.basename(file_name))[0]
    ...     save_path = os.path.join(output_folder, base_name + ".npy")
    ...     np.save(save_path, emb_array)
    ...     print(f"Saved: {save_path} with shape {emb_array.shape}")

# Run the function
audio_to_embedding_folders = {
    ... "/content/drive/My Drive/AlternativeBirds_wav/": "/content/drive/My Drive/Alter
    ... "/content/drive/My Drive/OnlyAudioDoliornis_wav/": "/content/drive/My Drive/Onl
    ... "/content/drive/My Drive/OnlyAudioHapalopsittaca_wav/": "/content/drive/My Driv
}

for audio_folder, embedding_folder in audio_to_embedding_folders.items():
    ... extract_embeddings(audio_folder, embedding_folder)
```

File BirdNET\_GLOBAL\_6K\_V2.4\_Labels\_af.txt already exists; skipping download.  
downloading model from URL...

File BirdNET\_GLOBAL\_6K\_V2.4\_Model\_FP16\_tflite already exists; skipping download

```

FILE D:\content\drive\My Drive\AlternativeBirds_wav/ already exists, skipping download.
Embedding 11 files from: /content/drive/My Drive/AlternativeBirds_wav/
/usr/local/lib/python3.11/dist-packages/opensoundscape/ml/cnn.py:599: UserWarning:
    This architecture is not listed in opensoundscape.ml.cnn_archi
    It will not be available for loading after saving the model wi
    To make it re-loadable, define a function that generates the a
    then use opensoundscape.ml.cnn_architectures.register_architec

```

The function can also set the returned object's .constructor\_n to avoid this warning and ensure it is reloaded correctly by o

See opensoundscape.ml.cnn\_architectures module for examples of

```

warnings.warn(
/usr/local/lib/python3.11/dist-packages/opensoundscape/ml/cnn.py:623: UserWarning:
    warnings.warn(
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC14584 - Grey-breasted Mounta
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC1598 - Blue-banded Toucanet
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC178 - Jameson_s Snipe - Gall
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC187 - Grey-breasted Mountain
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC2012 - Jameson_s Snipe - Gal
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC2019 - Plum-crowned Parrot -
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC2069 - Plum-crowned Parrot -
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC20706 - Blue-throated Piping
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC2915 - Andean Cock-of-the-ro
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC3470 - Blue-banded Toucanet
Saved: /content/drive/My Drive/AlternativeBirds_Emb/XC62895 - Black Tinamou - Tina
Embedding 12 files from: /content/drive/My Drive/OnlyAudioDoliornis_wav/
Saved: /content/drive/My Drive/OnlyAudioDoliornis_Emb/XC142403 - Bay-vented Cotinga
Saved: /content/drive/My Drive/OnlyAudioDoliornis_Emb/XC20615 - Bay-vented Cotinga
Saved: /content/drive/My Drive/OnlyAudioDoliornis_Emb/XC222458 - Bay-vented Cotinga
Saved: /content/drive/My Drive/OnlyAudioDoliornis_Emb/XC222459 - Bay-vented Cotinga
Saved: /content/drive/My Drive/OnlyAudioDoliornis_Emb/XC222460 - Bay-vented Cotinga
Saved: /content/drive/My Drive/OnlyAudioDoliornis_Emb/XC296816 - Bay-vented Cotinga
Saved: /content/drive/My Drive/OnlyAudioDoliornis_Emb/XC296818 - Bay-vented Cotinga
Saved: /content/drive/My Drive/OnlyAudioDoliornis_Emb/XC296819 - Bay-vented Cotinga
Saved: /content/drive/My Drive/OnlyAudioDoliornis_Emb/XC41676 - Bay-vented Cotinga
Saved: /content/drive/My Drive/OnlyAudioDoliornis_Emb/XC65640 - Bay-vented Cotinga
Embedding 11 files from: /content/drive/My Drive/OnlyAudioHapalopsittaca_wav/
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC151165 - Black-winged
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC151172 - Black-winged
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC16129 - Black-winged
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC2004 - Black-winged P
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC2005 - Black-winged P
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC2070 - Black-winged P
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC349146 - Black-winged
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC350403 - Black-winged
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC354158 - Black-winged
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC354160 - Black-winged
Saved: /content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/XC821443 - Black-winged

```

```
# --- Check number of embeddings per folder---
```

```
# Folder dictionary
```

```

folders = {
    0: "/content/drive/My Drive/AlternativeBirds_Emb/",
    1: "/content/drive/My Drive/OnlyAudioDoliornis_Emb/",

```

```

1: "/content/drive/My Drive/OnlyAudioDoliornis_Emb/",
2: "/content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/"
}

# Count embeddings
embedding_counts = {}

for label, folder_path in folders.items():
    total_embeddings = 0
    for file in os.listdir(folder_path):
        if file.endswith('.npy'):
            filepath = os.path.join(folder_path, file)
            data = np.load(filepath)

            # If 1D, count as 1 embedding; otherwise, use number of rows
            if data.ndim == 1:
                total_embeddings += 1
            else:
                total_embeddings += data.shape[0]

    embedding_counts[label] = total_embeddings

# Print results
for label, count in embedding_counts.items():
    print(f"Class {label}: {count} embeddings")

Class 0: 102 embeddings
Class 1: 154 embeddings
Class 2: 74 embeddings

# --- Check the number of embeddings in one recording ---
# Embeddings are of 3 seconds, therefore, some recordings may have more than one embed

# Choose a folder and file
folder = "/content/drive/My Drive/AlternativeBirds_Emb/"
files = [f for f in os.listdir(folder) if f.endswith('.npy')]

# Load one file (e.g., the first one)
file_path = os.path.join(folder, files[0])
data = np.load(file_path)

# Show info
print(f"File: {files[0]}")
print("Shape:", data.shape)
print("Rows:\n", data)

```

```

File: XC14584 - Grey-breasted Mountain Toucan - Andigena hypoglauca.npy
Shape: (4, 1024)
Rows:
[[1.00081    0.05319615 0.35048535 ... 0.82700044 0.6854153  0.01185115]
 [0.85154    0.08075234 0.6724665  ... 1.2943727  1.0188434  0.04159411]
 [0.14932956 0.05190347 0.4674158  ... 0.9484196  1.5497198  0.0411665 ]
 [0.5318307  0.         0.5636771  ... 1.0350096  1.2452976  0.00568366]]

```



## ✓ V. Evaluation of Models for Transfer Learning

### ✓ 5.1 Data Pre-Processing for Model Evaluation

```
# --- Data preprocessing for model evaluation ---

# Load embeddings and labels
X = []
y = []
file_ids = []

folders = {
    0: "/content/drive/My Drive/AlternativeBirds_Emb/",
    1: "/content/drive/My Drive/OnlyAudioDoliornis_Emb/",
    2: "/content/drive/My Drive/OnlyAudioHapalopsittaca_Emb/"
}

for label, folder_path in folders.items():
    for file in os.listdir(folder_path):
        if file.endswith('.npy'):
            filepath = os.path.join(folder_path, file)
            data = np.load(filepath)
            for row in data:
                X.append(row)
                y.append(label)
                file_ids.append(file) # Track filename

X = np.array(X)
y = np.array(y)
file_ids = np.array(file_ids)

X_all_data = X.copy()
y_all_data = y.copy()

# Split based on unique files to avoid data leakage.
# Reason: Embeddings from the same audio file are correlated.
# Including some for training and others for testing can inflate performance.

unique_files = np.unique(file_ids)
file_labels = [y[file_ids == f][0] for f in unique_files]

# Train and test split
train_files, test_files = train_test_split(
    unique_files, test_size=0.2, stratify=file_labels, random_state=42
)

train_mask = np.isin(file_ids, train_files)
test_mask = np.isin(file_ids, test_files)

X_train = X[train_mask]
```

```

x_train = x[train_mask]
y_train = y[train_mask]
x_test = x[test_mask]
y_test = y[test_mask]

# Normalize embeddings using StandardScaler
scaler = StandardScaler()
X_train = scaler.fit_transform(X_train)
X_test = scaler.transform(X_test)

# Check results
print("X_train shape:", X_train.shape)
print("y_train shape:", y_train.shape)
print("X_test shape:", X_test.shape)
print("y_test shape:", y_test.shape)

```

```

X_train shape: (249, 1024)
y_train shape: (249,)
X_test shape: (81, 1024)
y_test shape: (81,)

```

## ✓ 5.2 Evaluation of Multinomial Logistic Regression (Standard)

```

# --- Evaluation of multinomial logistic regression for prediction of rare species -

# Train the model
clf = LogisticRegression(solver='lbfgs', max_iter=500) #multinomial
clf.fit(X_train, y_train)

# Evaluate
y_pred = clf.predict(X_test)
print(classification_report(y_test, y_pred))

acc_mnl = round(accuracy_score(y_test, y_pred), 4)
print("\nAccuracy:", acc_mnl)

```

	precision	recall	f1-score	support
0	1.00	0.84	0.91	25
1	0.78	1.00	0.88	42
2	0.67	0.29	0.40	14
accuracy			0.83	81
macro avg	0.81	0.71	0.73	81
weighted avg	0.83	0.83	0.80	81

Accuracy: 0.8272

## ✓ 5.3 Evaluation of Multinomial Logistic Regression with Cross Validation

```

# --- Evaluation of multinomial logistic regression with CV for prediction of rare

### Cross-validation setup
X = []
y = []
groups = [] # file-level grouping

for label, folder_path in folders.items():
    for file in os.listdir(folder_path):
        if file.endswith('.npy'):
            filepath = os.path.join(folder_path, file)
            data = np.load(filepath)
            for row in data:
                X.append(row)
                y.append(label)
            groups.append(file) # group by filename

X = np.array(X)
y = np.array(y)
groups = np.array(groups)

cv = StratifiedGroupKFold(n_splits=5)

# Evaluate model with cross validation
accuracies = []
fold = 1
for train_idx, test_idx in cv.split(X, y, groups):
    print(f"\n=== Fold {fold} ===")

    X_train, X_test = X[train_idx], X[test_idx]
    y_train, y_test = y[train_idx], y[test_idx]

    clf = LogisticRegression(solver='lbfgs', max_iter=500) #multinomial
    clf.fit(X_train, y_train)
    y_pred = clf.predict(X_test)

    acc = accuracy_score(y_test, y_pred)
    accuracies.append(acc)

    print(classification_report(y_test, y_pred))
    fold += 1

# Print average accuracy
avg_acc_mnl = round(np.mean(accuracies), 4)
print("Average Accuracy:", avg_acc_mnl)

```

```
=== Fold 1 ===
```

	precision	recall	f1-score	support
0	0.80	0.21	0.33	19
1	0.68	1.00	0.81	40
2	1.00	0.64	0.78	14

accuracy			0.73	73
macro avg	0.83	0.62	0.64	73
weighted avg	0.77	0.73	0.68	73

=== Fold 2 ===

	precision	recall	f1-score	support
0	0.94	1.00	0.97	17
1	1.00	1.00	1.00	40
2	1.00	0.93	0.96	14

accuracy			0.99	71
macro avg	0.98	0.98	0.98	71
weighted avg	0.99	0.99	0.99	71

=== Fold 3 ===

	precision	recall	f1-score	support
0	0.78	1.00	0.88	29
1	1.00	0.83	0.91	24
2	0.90	0.64	0.75	14

accuracy			0.87	67
macro avg	0.89	0.83	0.85	67
weighted avg	0.89	0.87	0.86	67

=== Fold 4 ===

	precision	recall	f1-score	support
0	1.00	0.85	0.92	20
1	0.69	1.00	0.82	25
2	0.67	0.29	0.40	14

accuracy			0.78	59
macro avg	0.79	0.71	0.71	59
weighted avg	0.79	0.78	0.75	59

=== Fold 5 ===

	precision	recall	f1-score	support
0	1.00	0.94	0.97	17
1	0.96	1.00	0.98	25
2	0.94	0.94	0.94	18

accuracy			0.97	60
macro avg	0.97	0.96	0.96	60

## ▼ 5.4 Evaluation of KNN model

```
# --- Evaluation of KNN for prediction of rare species ---
```

```
# Train the model (K=3)
```

```

knn = KNeighborsClassifier(n_neighbors=3)
knn.fit(X_train, y_train)

# Evaluate
y_pred_knn = knn.predict(X_test)
print(classification_report(y_test, y_pred_knn))

acc_knn = round(accuracy_score(y_test, y_pred), 4)
print("\nAccuracy:", acc_knn)

```

	precision	recall	f1-score	support
0	0.93	0.76	0.84	17
1	0.58	1.00	0.74	25
2	0.67	0.11	0.19	18
accuracy			0.67	60
macro avg	0.73	0.63	0.59	60
weighted avg	0.71	0.67	0.60	60

Accuracy: 0.9667

## ✓ 5.4 Evaluation of KNN model with Cross Validation

```

# --- Evaluation of KNN with CV for prediction of rare species ---

# Set up Stratified Group K-Fold CV
cv = StratifiedGroupKFold(n_splits=5)

accuracies = []
fold = 1
for train_idx, test_idx in cv.split(X, y, groups):
    ... print(f"\n=== Fold {fold} ===")

    ... X_train, X_test = X[train_idx], X[test_idx]
    ... y_train, y_test = y[train_idx], y[test_idx]

    ... # Train and evaluate KNN
    ... knn = KNeighborsClassifier(n_neighbors=3)
    ... knn.fit(X_train, y_train)
    ... y_pred_knn = knn.predict(X_test)

    ... acc = accuracy_score(y_test, y_pred_knn)
    ... accuracies.append(acc)

    ... print(classification_report(y_test, y_pred_knn, zero_division=0))
    ... fold += 1

# Print average accuracy
avg_acc_knn = round(np.mean(accuracies), 4)
print("Average Accuracy:", avg_acc_knn)

```

=== Fold 1 ===

	precision	recall	f1-score	support
0	1.00	0.05	0.10	19
1	0.62	1.00	0.77	40
2	1.00	0.57	0.73	14
accuracy			0.67	73
macro avg	0.88	0.54	0.53	73
weighted avg	0.79	0.67	0.59	73

=== Fold 2 ===

	precision	recall	f1-score	support
0	1.00	0.94	0.97	17
1	0.91	1.00	0.95	40
2	1.00	0.79	0.88	14
accuracy			0.94	71
macro avg	0.97	0.91	0.93	71
weighted avg	0.95	0.94	0.94	71

=== Fold 3 ===

	precision	recall	f1-score	support
0	0.93	0.97	0.95	29
1	0.70	0.96	0.81	24
2	1.00	0.29	0.44	14
accuracy			0.82	67
macro avg	0.88	0.74	0.73	67
weighted avg	0.86	0.82	0.79	67

=== Fold 4 ===

	precision	recall	f1-score	support
0	1.00	0.55	0.71	20
1	0.57	1.00	0.72	25
2	1.00	0.29	0.44	14
accuracy			0.68	59
macro avg	0.86	0.61	0.63	59
weighted avg	0.82	0.68	0.65	59

=== Fold 5 ===

	precision	recall	f1-score	support
0	0.93	0.76	0.84	17
1	0.58	1.00	0.74	25
2	0.67	0.11	0.19	18
accuracy			0.67	60
macro avg	0.73	0.62	0.59	60
weighted avg	0.67	0.62	0.59	60

macro avg	0.73	0.63	0.59	60
-----------	------	------	------	----

## 5.5 Evaluation of Few Shot Learning

```
# --- Evaluation of few-shot learning for prediction of rare species ---
# The simplest version using nearest neighbor with cosine similarity.
```

```
def few_shot_predict(X_train, y_train, X_query, k=3):
    ... similarities = cosine_similarity(X_query, X_train)
    ... predictions = []
    ... for sim_row in similarities:
    ...     top_k_indices = sim_row.argsort()[-k:][::-1]
    ...     top_k_labels = y_train[top_k_indices]
    ...     predicted_label = Counter(top_k_labels).most_common(1)[0][0]
    ...     predictions.append(predicted_label)
    ... return predictions
```

```
y_pred_fewshot = few_shot_predict(X_train, y_train, X_test, k=3)
print(classification_report(y_test, y_pred_fewshot))
```

```
acc_fewshot = round(accuracy_score(y_test, y_pred), 4)
print("\nAccuracy:", acc_fewshot)
```

	precision	recall	f1-score	support
0	0.94	0.94	0.94	17
1	0.83	0.96	0.89	25
2	0.93	0.72	0.81	18
accuracy			0.88	60
macro avg	0.90	0.87	0.88	60
weighted avg	0.89	0.88	0.88	60

Accuracy: 0.9667

## 5.6 Evaluation of Few Shot Learning with Cross Validation

```
# --- Evaluation of few-shot learning with CV for prediction of rare species ---
```

```
cv = StratifiedGroupKFold(n_splits=5)
accuracies = []
fold = 1
for train_idx, test_idx in cv.split(X, y, groups):
    ... print(f"\n=== Fold {fold} ===")
    ... X_train, X_test = X[train_idx], X[test_idx]
    ... y_train, y_test = y[train_idx], y[test_idx]

    ... y_pred = few_shot_predict(X_train, y_train, X_test, k=3)
    ... acc = accuracy_score(y_test, y_pred)
    ... accuracies.append(acc)
```

```
...print(classification_report(y_test, y_pred, zero_division=0))
...fold += 1

# Print average accuracy
avg_acc_fewshot = round(np.mean(accuracies), 4)
print("Average Accuracy:", avg_acc_fewshot)
```

=== Fold 1 ===

	precision	recall	f1-score	support
0	0.00	0.00	0.00	19
1	0.63	1.00	0.78	40
2	1.00	0.71	0.83	14
accuracy			0.68	73
macro avg	0.54	0.57	0.54	73
weighted avg	0.54	0.68	0.59	73

=== Fold 2 ===

	precision	recall	f1-score	support
0	1.00	1.00	1.00	17
1	0.98	1.00	0.99	40
2	1.00	0.93	0.96	14
accuracy			0.99	71
macro avg	0.99	0.98	0.98	71
weighted avg	0.99	0.99	0.99	71

=== Fold 3 ===

	precision	recall	f1-score	support
0	0.97	0.97	0.97	29
1	0.86	1.00	0.92	24
2	1.00	0.71	0.83	14
accuracy			0.93	67
macro avg	0.94	0.89	0.91	67
weighted avg	0.93	0.93	0.92	67

=== Fold 4 ===

	precision	recall	f1-score	support
0	1.00	0.90	0.95	20
1	0.68	1.00	0.81	25
2	1.00	0.29	0.44	14
accuracy			0.80	59
macro avg	0.89	0.73	0.73	59
weighted avg	0.86	0.80	0.77	59



=== Fold 5 ===

	precision	recall	f1-score	support
0	0.94	0.94	0.94	17
1	0.83	0.96	0.89	25
2	0.93	0.72	0.81	18
accuracy			0.88	60
macro avg	0.90	0.87	0.88	60

## ✓ 5.7 Evaluation of Neural Network (MLP)

# --- Evaluation of MLP (Multilayer Perceptron) for prediction of rare species ---  
 # A type of feedforward neural network suitable for small samples

# Train the model

```
clf = MLPClassifier(hidden_layer_sizes=(256, 128), max_iter=300, random_state=42)
clf.fit(X_train, y_train)
```

# Evaluate

```
print(classification_report(y_test, clf.predict(X_test)))
```

```
acc_mlp = round(accuracy_score(y_test, y_pred), 4)
print("\nAccuracy:", acc_mlp)
```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	17
1	0.83	1.00	0.91	25
2	1.00	0.72	0.84	18
accuracy			0.92	60
macro avg	0.94	0.91	0.92	60
weighted avg	0.93	0.92	0.91	60

Accuracy: 0.8833

## ✓ 5.8 Evaluation of Neural Network (MLP) with Cross Validation

# --- Evaluation of MLP (Multilayer Perceptron) with CV for prediction of rare spec  
 import pandas as pd

fold = 1

accuracies = []

```
for train_idx, test_idx in cv.split(X, y, groups):
    ... print(f"\n=== Fold {fold} ===")
```

```
    ... X_train, X_test = X[train_idx], X[test_idx]
```

```
    ... y_train, y_test = y[train_idx], y[test_idx]
```

```

...clf = MLPClassifier(hidden_layer_sizes=(128, 64), max_iter=300, random_state=42)
...clf.fit(X_train, y_train)
...y_pred = clf.predict(X_test)

...acc = accuracy_score(y_test, y_pred)
...accuracies.append(acc)

...print(pd.DataFrame(classification_report(y_test, y_pred, output_dict=True)).tra
...fold += 1

# Print average accuracy
avg_acc_mlp = round(np.mean(accuracies), 4)
print("\nAverage Accuracy:", avg_acc_mlp)

```

=== Fold 1 ===

```

<frozen importlib._bootstrap>:1047: ImportWarning: _PyDrive2ImportHook.find_spec()
<frozen importlib._bootstrap>:1047: ImportWarning: _PyDriveImportHook.find_spec()
<frozen importlib._bootstrap>:1047: ImportWarning: _BokehImportHook.find_spec() no

```

	precision	recall	f1-score	support
0	0.833333	0.263158	0.400000	19.000000
1	0.714286	1.000000	0.833333	40.000000
2	0.909091	0.714286	0.800000	14.000000
accuracy	0.753425	0.753425	0.753425	0.753425
macro avg	0.818903	0.659148	0.677778	73.000000
weighted avg	0.782631	0.753425	0.714155	73.000000

=== Fold 2 ===

	precision	recall	f1-score	support
0	0.944444	1.000000	0.971429	17.000000
1	1.000000	0.975000	0.987342	40.000000
2	0.928571	0.928571	0.928571	14.000000
accuracy	0.971831	0.971831	0.971831	0.971831
macro avg	0.957672	0.967857	0.962447	71.000000
weighted avg	0.972613	0.971831	0.971943	71.000000

=== Fold 3 ===

	precision	recall	f1-score	support
0	0.805556	1.000000	0.892308	29.000000
1	1.000000	0.875000	0.933333	24.000000
2	0.900000	0.642857	0.750000	14.000000
accuracy	0.880597	0.880597	0.880597	0.880597
macro avg	0.901852	0.839286	0.858547	67.000000
weighted avg	0.894942	0.880597	0.877268	67.000000

=== Fold 4 ===

	precision	recall	f1-score	support
0	1.000000	0.800000	0.888889	20.000000
1	0.714286	1.000000	0.833333	25.000000
2	0.500000	0.285714	0.363636	14.000000
accuracy	0.762712	0.762712	0.762712	0.762712
macro avg	0.738095	0.695238	0.695286	59.000000
weighted avg	0.760291	0.762712	0.740712	59.000000

=== Fold 5 ===

	precision	recall	f1-score	support
0	0.944444	1.000000	0.971429	17.000000

1	1.000000	0.960000	0.979592	25.000000
2	1.000000	1.000000	1.000000	18.000000
accuracy	0.983333	0.983333	0.983333	0.983333
macro avg	0.981481	0.986667	0.983673	60.000000
weighted avg	0.984259	0.983333	0.983401	60.000000

Average Accuracy: 0.8704

## ✓ VI. Summary of Model Evaluation Results

```
# Summary of accuracy results
results_table = pd.DataFrame({
    ... "Model (with Cross Validation)": ["Multinomial Logistic", "K-Nearest Neighbors",
    ... "Avg Accuracy": [avg_acc_mnl, avg_acc_knn, avg_acc_fewshot, avg_acc_mlp]
})

# Print results
print(results_table)
```

	Model (with Cross Validation)	Avg Accuracy
0	Multinomial Logistic	0.8648
1	K-Nearest Neighbors	0.7561
2	Few-Shot Learning	0.8552
3	Multilayer Perceptron	0.8704

## ✓ VII. Selection of the Best Model Using Transfer Learning for Rare Species Prediction

Based on cross-validated performance using BirdNET embeddings, the Multilayer Perceptron (MLP) achieved the highest average accuracy and is selected as the most reliable model for identifying our rare bird species.

```
# Train the final model on all data
final_mlp = MLPClassifier(hidden_layer_sizes=(128, 64), max_iter=300, random_state=
final_mlp.fit(X_all_data, y_all_data)

# Save the model
joblib.dump(final_mlp, "mlp_model_birdnet.pkl")
print("MLP model saved as 'mlp_model_birdnet.pkl'")
```

MLP model saved as 'mlp\_model\_birdnet.pkl'

## ✓ VIII. Example: Applying the Transfer Learning Model for Rare Bird Species Identification

```
# Prepare the recording
input_file = "XC20615 - Bay-vented Cotinga - Doliornis sclateri.mp3"
output_file = "XC20615 - Bay-vented Cotinga - Doliornis sclateri.wav"

# Pre-process the recording
convert_and_pad_to_3s(input_file, output_file)
```

```
Processed and saved: XC20615 - Bay-vented Cotinga - Doliornis sclateri.wav
<ipython-input-2-3223057341c3>:31: ResourceWarning: unclosed file <_io.BufferedRan
audio.export(output_path, format="wav")
```

```
# Get embeddings from BirdNET
model = bmz.BirdNET()
embedding1 = model.embed(output_file)
X_input = embedding1.reset_index(drop=True).values
print("\nThe embeddings are:\n", X_input)
```

```
File BirdNET_GLOBAL_6K_V2.4_Labels_af.txt already exists; skipping download.
downloading model from URL...
File BirdNET_GLOBAL_6K_V2.4_Model_FP16.tflite already exists; skipping download.
/usr/local/lib/python3.11/dist-packages/opensoundscape/ml/cnn.py:599: UserWarning:
    This architecture is not listed in opensoundscape.ml.cnn_archi
    It will not be available for loading after saving the model wi
    To make it re-loadable, define a function that generates the a
    then use opensoundscape.ml.cnn_architectures.register_architec

    The function can also set the returned object's .constructor_n
    to avoid this warning and ensure it is reloaded correctly by o

    See opensoundscape.ml.cnn_architectures module for examples of
```

```
warnings.warn(
/usr/local/lib/python3.11/dist-packages/opensoundscape/ml/cnn.py:623: UserWarning:
    warnings.warn(
```

```
The embeddings are:
[[0.03338824 0.8628469 0.5336458 ... 0.609214 0.07550486 0.37615904]
 [1.0899563 0.1301161 0.08624835 ... 0.32889202 0.61821526 0.38222626]
 [0.8048906 0.5017663 0. ... 0.16950338 0.43497276 0.18075454]
 ...
 [0. ... 0.9102369 0.27659604 ... 1.2303518 0.17949228 0. ... ]
 [0.18699229 0.32541603 0.18230295 ... 1.7507849 0. ... 0.01350991]
 [1.0755877 0.4611834 0.09565047 ... 0.15522702 0.07813891 0.19568026]]
```

```
# Load the saved model for prediction
model_transferlearning = joblib.load("mlp_model_birdnet.pkl")
y_pred = model_transferlearning.predict(X_input)
```

```
# Mapping of label to species name
label_to_species = {
    0: "No species identified",
    1: "Doliornis sclateri"
```

```
1: Doliornis sclateri ,
2: "Hapalopsittaca melanotis"
}

# Convert predictions to species names
species_pred = [label_to_species[label] for label in y_pred]

# Print or use the result
print("Predictions per audio segment:", species_pred)

# Most frequent prediction
most_common = Counter(species_pred).most_common(1)[0][0]
print("\nFinal predicted species:", most_common)
```

Predictions per audio segment: ['Doliornis sclateri', 'Doliornis sclateri', 'Dolio

Final predicted species: Doliornis sclateri