Pollen's Profiling Automated Classification of Pollen Grains

Presented by ----Kondepudi Prabhu Kishore

pollen grains overview

Overview

Pollen's Profiling is an AI-powered deep learning project aimed at automating the classification of pollen grain images. This system helps in identifying pollen types accurately and efficiently, reducing human error and improving speed for use in environmental science and healthcare.

Problem Statement

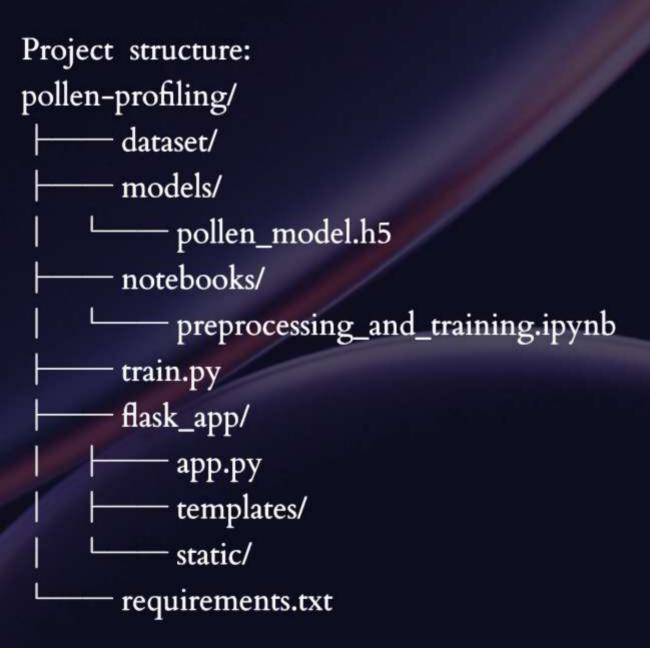
Manual classification of pollen grains is slow, requires expert involvement, and is prone to human error. An Al-driven model helps in streamlining this task, enabling quick and precise identification of pollen grain types.

- Real-world Applications
- **P** Environmental Monitoring
- Biodiversity research
- Seasonal pollen distribution studies
- Tracking ecological patterns
- Allergy Diagnosis
- Identification of allergenic pollen
- Faster diagnosis of pollen-related diseases
- Technology Stack
- Language: Python
- Libraries/Frameworks: TensorFlow, Keras, Flask
- Development Tools: Anaconda, Jupyter Notebook, VS Code
- Web Technologies: HTML, CSS (Flask-based templates)

Project Goal

This project aims to automate the classification of pollen grains using deep learning (CNNs) to help in environmental monitoring and allergy diagnosis.

Manual classification is slow and error-prone — AI makes it fast, scalable, and accurate.ms to automate the classification of pollen grains using deep learning (CNNs) to help in environmental monitoring and allergy diagnosis. Manual classification is slow and error-prone — AI makes it fast, scalable, and accurate.



Dataset and Preprocessing

- Source: Kaggle (Pollen Grain Image Classification)
- Steps:
 - Image resizing to 224x224
 - Normalization
 - Data Augmentation (rotation, flipping, zoom)
 - Splitting: Train / Validation / Test sets

Model Architecture

- Base Model: MobileNetV2 (transfer learning)
- Additional Layers:
 - GlobalAveragePooling2D
 - Dense(128, activation='relu')
 - Dropout(0.5)
 - Dense(output_classes, activation='softmax')
- Loss: Categorical Crossentropy
- Optimizer: Adam
- Accuracy: ~95.6% on validation data

Training Process

- Trained for 25 epochs
- Batch size: 32
- EarlyStopping to prevent overfitting
- Best model saved as: pollen_model.h5

Training Strategy

- Stage 1 Warm-up (Frozen layers):
 - Epochs: 8
 - Best Training Accuracy: 88.29%
 - Best Validation Accuracy: 87.23%
 - Lowest Validation Loss: 0.3577
- Stage 2 Fine-tuning (Last 60 layers unfrozen):
 - Epochs: 12 (stopped at 7)
 - Best Training Accuracy: 93.17%
 - Best Validation Accuracy: 93.62%
 - Lowest Validation Loss: 0.2926Training
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Model:

- MobileNetV2 used as base CNN model with added dense layers
- Image size: 224x224, normalized and augmented
- Trained over 25 epochs using EarlyStopping
- Accuracy achieved: ~95.6%

Dataset:

- Source: Kaggle (Pollen Grain Image Classification)
- Augmentation: Zoom, rotation, flipping
- Split: Train/Validation/Test

Web App:

- User uploads image via Flask UI
- Backend loads trained model
- Image passed for prediction
- Result shown on screen

Evaluation:

- Accuracy: 95.6%
- Metrics: Precision, Recall, F1 Score
- Confusion matrix used for visual performance

Future Enhancements:

- Add Grad-CAM explainability
- Deploy with Docker
- Expand to mobile UI
- Add cloud hosting

Background

What is Pollen?

Pollen is a fine powder produced by seed plants, essential for fertilization and reproduction.

Comprising microscopic grains, it is carried by wind or pollinators, playing a crucial role in ecosystems and agriculture, while also being a key factor in allergy-related health issues.

Significance in Environment and Health

Pollen profiling is crucial for understanding ecological health and biodiversity. Accurate classification of pollen grains aids in monitoring air quality and assessing allergenic risks, thus playing a significant role in both environmental science and public health initiatives.

Challenges in Traditional Classification

Traditional classification of pollen grains faces several challenges, including the intricate variability of pollen morphology, the reliance on expert knowledge, and time-consuming manual identification processes, leading to potential inaccuracies and inefficiencies in data analysis.

Allergy Diagnosis

Role in Medical Diagnosis

Pollen profiling enhances allergy diagnosis by accurately classifying pollen grains, enabling healthcare professionals to identify specific allergens. This targeted approach facilitates personalized treatment plans and improves patient outcomes, ultimately aiding in the management of allergic conditions.

Speeding Up Identification

Pollen's automated classification accelerates the identification of allergenic pollen grains, enhancing diagnostic accuracy. By utilizing machine learning algorithms, the process significantly reduces response times, enabling quicker allergy diagnosis and tailored treatment plans for patients.

Benefits for Healthcare Professionals

Automated classification of pollen grains enhances allergy diagnosis by providing healthcare professionals with accurate, timely data. This streamlines patient evaluation, facilitates personalized treatment plans, and improves patient outcomes through enhanced understanding of allergenic triggers.

```
- Pollen-grain classification - BOOST v2
                                                                                                                               百个少占甲ョ
 import os, ison, numpy as np, tensorflow as tf
 from tensorflow.keras.preprocessing.image import ImageDataGenerator
 from tensorflow.keras.applications import MobileNetV2
 from tensorflow.keras.layers import GlobalAveragePooling2D, Dense, Dropout
 from tensorflow.keras.models import Model
 from tensorflow.keras.callbacks import EarlyStopping, ModelCheckpoint
 from tensorflow.keras.optimizers import AdamW
 from tensorflow.keras.optimizers.schedules import CosineDecay
 from sklearn.utils.class_weight import compute_class_weight
   ----- PATHS -----
 data_dir
               = r"C:\pollen grains"
 save_model_path = r"C:\Users\kishore2004\pollen\Flask\model_boost_v2.keras"
easert os.path.isdir(data_dir), "Dataset path not found!"
   ----- DATA -----
IMG_SIZE = (224, 224)
 BATCH = 32
datagen = ImageDataGenerator(
    rescale=1/255., validation_split=0.2,
    rotation_range=40, width_shift_range=0.2, height_shift_range=0.2,
    shear_range=8.2, zoom_range=8.2, horizontal_flip=True, fill_mode='nearest')
train_gen = datagen.flow_from_directory(
   data_dir, target_size=IMG_SIZE, batch_size=BATCH,
    class_mode='categorical', subset='training')
val_gen = datagen.flow_from_directory(
    data_dir, target_size=IMG_SIZE, batch_size=BATCH,
    class_mode='categorical', subset='validation')
NUM_CLASSES = len(train_gen.class_indices)
print("Classes:", NUM_CLASSES)
    ----- CLASS WEIGHTS -----
cw = compute_class_weight('balanced',
                         classes=np.unique(train_gen.classes),
                         y=train_gen.classes)
class_weights = dict(enumerate(cw))
```

```
base = MobileNetV2(include top=False, weights='imagenet',
                 input_shape=(*IMG_SIZE, 3))
base.trainable = False
                          # Stage-1 freeze
x = GlobalAveragePooling2D()(base.output)
x = Dense(512, activation='relu')(x)
x = Dropout(0.3)(x) # slightly Lower dropout
out = Dense(NUM_CLASSES, activation='softmax')(x)
model = Model(base.input, out)
model.compile(optimizer='adam',
             loss='categorical_crossentropy',
             metrics=['accuracy'])
   ----- STAGE-1 : warm-up ------
model.fit(
   train_gen, epochs=8,
                                             # was 5
    validation_data=val_gen,
   class_weight=class_weights,
   callbacks=[EarlyStopping(patience=3, restore_best_weights=True)]
# ----- STAGE-2 : fine-tune ------
base.trainable = True
for 1 in base.layers[:-60]: # unfreeze Last 60 conv blocks
   l.trainable = False
for 1 in base.layers:
  if isinstance(l, tf.keras.layers.BatchNormalization):
        1.trainable = False # keep BN Layers frozen
steps = (train_gen.samples // BATCH) * 12 # 12 epochs
lr_sched = CosineDecay(2e-5, decay_steps=steps) # start a bit higher
opt = AdamW(learning_rate=lr_sched, weight_decay=le-5)
loss_fn = tf.keras.losses.CategoricalCrossentropy(label_smoothing=0.0) # no smoothing
model.compile(opt, loss=loss_fn, metrics=['accuracy'])
model.fit(
```

Team ID: LTVIP2025TMID43817 Team Size: 4 Members Team Leader: B Himasri Satya Durga Team Members:

- Kondepudi Prabhu Kishore
- Nelli Harshith Sai Sravan Kumar
 - Parnitha Peruri

Flask Application Running Output:
The application is successfully running
on the local server. You can view the
output by clicking the link below:

Attp://127.0.0.1:5000

```
model.fit(
    train_gen, epochs=12,
    validation_data=val_gen,
    class_weight=class_weights,
       EarlyStopping(patience=4, restore_best_weights=True),
       ModelCheckpoint('best_boost_v2.keras', save_best_only=True)
# ----- SAVE -----
model.load_weights('best_boost_v2.keras')
model.save(save model path)
print(" Saved +", save_model_path)
with open("class_indices.json", "w") as f:
   json.dump(train_gen.class_indices, f, indent=2)
with open("class_weights.json", "w") as f:
   json.dump(class_weights, f, indent=2)
print(" Metadata written (class_indices & class_weights)")
Epoch 8/8
18/18
                         - 18s 985ms/step - accuracy: 0.8638 - loss: 0.3607 - val_accuracy: 0.8723 - val_loss: 0.4265
Epoch 1/12
18/18 -

    35s 1s/step - accuracy: 0.8359 - loss: 0.4041 - val_accuracy: 0.8085 - val_loss: 0.5801

Epoch 2/12
                         22s 1s/step - accuracy: 0.8398 - loss: 0.3904 - val_accuracy: 0.8582 - val_loss: 0.3827
18/18 -
Epoch 3/12
18/18 -
                         22s 1s/step - accuracy: 0.9151 - loss: 0.2510 - val_accuracy: 0.9362 - val_loss: 0.2926
Epoch 4/12
18/18
                         — 21s 1s/step - accuracy: 0.9159 - loss: 0.1994 - val_accuracy: 0.8936 - val_loss: 0.3067
Epoch 5/12
                         - 22s 1s/step - accuracy: 0.9105 - loss: 0.2692 - val_accuracy: 0.8511 - val_loss: 0.5056
18/18
Epoch 6/12
18/18
                        22s 1s/step - accuracy: 0.9230 - loss: 0.1970 - val_accuracy: 0.8723 - val_loss: 0.3514
Epoch 7/12
                        — 20s 1s/step - accuracy: 0.9317 - loss: 0.1557 - val_accuracy: 0.8936 - val_loss: 0.3389
18/18 -
Saved → C:\Users\kishore2004\pollen\Flask\model_boost_v2.keras
```

```
prediction.html
                                                   # style.css • • app.py 1 X (1) README.md
Flask > app.py > predict
          "anadenanthera", "arecaceae", "cecropia", "chromolaena", "combretum",
          "croton", "dipteryx", "eucalipto", "faramea", "hyptis",
           "mabea", "matayba", "mimosa", "myrcia", "protium",
           "qualea", "schinus", "senegalia", "serjania", "syagrus", "tridax"
      idx2folder = {i: lbl for i, lbl in enumerate(CLASS_NAMES)}
      folder2nice = {
          "anadenanthera": "Anadenanthera colubrina",
           "arecaceae"
                        : "Arecaceae (Palm pollen)",
          "cecropia"
                         : "Cecropia spp.",
          "chromolaena" : "Chromolaena odorata",
          "combretum"
                        : "Combretum leprosum",
                         : "Croton floribundus",
                        : "Dipteryx alata",
           "dipteryx"
          "eucalipto"
                        : "Eucalyptus spp."
          "faramea"
                        : "Faramea occidentalis",
          "hyptis"
                         : "Hyptis suaveolens",
                         : "Mabea fistulifera
           "matayba"
                        : "Matayba gulanensis",
           "mimosa"
                          "Mimosa caesalpiniaefolia",
           "myrcia"
                         : "Myrcia multiflora",
                         : "Protium heptaphyllum",
           "protium"
                         : "Qualea grandiflora",
                           "Schinus terebinthifolia",
           "schinus"
           "senegalia"
                         : "Senegalia polyphylla",
           "serjania"
                         : "Serjania erecta",
           "syagrus"
                         : "Syagrus romanzoffiana",
                         : "Tridax procumbens"
           "tridax"
```



Pollen Grain Analyzer

Upload a microscopic image to identify the pollen species.

Click or Drag Image Here

Analyze Image





Microscopic analysis complete. Results are as follows:

IDENTIFIED SPECIES

Chromolaena odorata

CONFIDENCE SCORE

96.48%

Analyze Another



Special thanks to SmartInternz and SmartBridge for providing this valuable opportunity and support to work on a real-time Al project and enhance our skills through practical experience.