

# New Plant Disease Detection

## CSML1020 Course Project

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### ABSTRACT

This paper will explore the classification of plant images to identify new plant diseases using a machine learning model.

The dataset was obtained from Kaggle and consists of over 87,000 rgb images of healthy and diseased crop leaves labeled by plant and disease type in 38 different classes.

### CCS CONCEPTS

• Artificial Intelligence • Machine Learning • Image Classification

## 1 Introduction

The problem we will examine is a supervised multi-class image classification problem. The goal is to investigate which supervised machine learning models will give the best results in classifying the images from our dataset in the predefined categories.

## 2 Existing Work

|   |   |
|---|---|
| Plant Disease Classification-VGG16<br><a href="https://www.kaggle.com/wiwidsetiawan/plant-disease-classification-vgg16">https://www.kaggle.com/wiwidsetiawan/plant-disease-classification-vgg16</a>   | Example of classification using the VGG16 pre trained model               |
| Fork of Plant Diseases Classification Using inception3<br><a href="https://www.kaggle.com/vimaladit/fork-of-plant-diseases-classification-using-inception3">https://www.kaggle.com/vimaladit/fork-of-plant-diseases-classification-using-inception3</a> | Example of classification using the Inception Version 3 pre trained model |
|   |   |
|   |   |

## 3 Methodology

### 3.1 Data Preparation

The dataset was downloaded from the Kaggle website:  
<https://www.kaggle.com/vipooool/new-plant-diseases-dataset>

The dataset did not need any manipulation as it was previously divided into a useable directory structure for training and validation as well as several test images.

**Table 1: Dataset Parsed from Category Folder Names**

|    | plant                   | condition                            | count | status    | disease  |
|----|-------------------------|--------------------------------------|-------|-----------|----------|
| 0  | Tomato                  | Target_Spot                          | 50    | unhealthy | spot     |
| 1  | Tomato                  | Early_blight                         | 50    | unhealthy | blight   |
| 2  | Apple                   | healthy                              | 50    | healthy   | healthy  |
| 3  | Tomato                  | healthy                              | 50    | healthy   | healthy  |
| 4  | Blueberry               | healthy                              | 50    | healthy   | healthy  |
| 5  | Grape                   | healthy                              | 50    | healthy   | healthy  |
| 6  | Peach                   | healthy                              | 50    | healthy   | healthy  |
| 7  | Cherry_(including_sour) | Powdery_mildew                       | 50    | unhealthy | mildew   |
| 8  | Tomato                  | Leaf_Mold                            | 50    | unhealthy | mold     |
| 9  | Apple                   | Black_rot                            | 50    | unhealthy | rot      |
| 10 | Squash                  | Powdery_mildew                       | 50    | unhealthy | mildew   |
| 11 | Corn_(maize)            | Northern_Leaf_Blight                 | 50    | unhealthy | blight   |
| 12 | Strawberry              | Leaf_scorch                          | 50    | unhealthy | scorch   |
| 13 | Pepper_bell             | healthy                              | 50    | healthy   | healthy  |
| 14 | Orange                  | Huanglongbing_(Citrus_greening)      | 50    | unhealthy | greening |
| 15 | Potato                  | Late_blight                          | 50    | unhealthy | blight   |
| 16 | Tomato                  | Late_blight                          | 50    | unhealthy | blight   |
| 17 | Strawberry              | healthy                              | 50    | healthy   | healthy  |
| 18 | Tomato                  | Tomato_Yellow_Leaf_Curl_Virus        | 50    | unhealthy | virus    |
| 19 | Corn_(maize)            | Common_rust                          | 50    | unhealthy | rust     |
| 20 | Raspberry               | healthy                              | 50    | healthy   | healthy  |
| 21 | Tomato                  | Tomato_mosaic_virus                  | 50    | unhealthy | virus    |
| 22 | Pepper_bell             | Bacterial_spot                       | 50    | unhealthy | spot     |
| 23 | Cherry_(including_sour) | healthy                              | 50    | healthy   | healthy  |
| 24 | Tomato                  | Septoria_leaf_spot                   | 50    | unhealthy | spot     |
| 25 | Peach                   | Bacterial_spot                       | 50    | unhealthy | spot     |
| 26 | Apple                   | Cedar_apple_rust                     | 50    | unhealthy | rust     |
| 27 | Tomato                  | Bacterial_spot                       | 50    | unhealthy | spot     |
| 28 | Grape                   | Esca_(Black_Measles)                 | 50    | unhealthy | measles  |
| 29 | Grape                   | Leaf_blight_(Isariopsis_Leaf_Spot)   | 50    | unhealthy | spot     |
| 30 | Corn_(maize)            | Cercospora_leaf_spot_Gray_leaf_spot  | 50    | unhealthy | spot     |
| 31 | Apple                   | Apple_scab                           | 50    | unhealthy | scab     |
| 32 | Grape                   | Black_rot                            | 50    | unhealthy | rot      |
| 33 | Potato                  | healthy                              | 50    | healthy   | healthy  |
| 34 | Corn_(maize)            | healthy                              | 50    | healthy   | healthy  |
| 35 | Potato                  | Early_blight                         | 50    | unhealthy | blight   |
| 36 | Soybean                 | healthy                              | 50    | healthy   | healthy  |
| 37 | Tomato                  | Spider_mites_Two-spotted_spider_mite | 50    | unhealthy | mite     |

### 3.2 Data Exploration

The dataset consists of images of plant leaves in various conditions. The following graphs, show the distribution of this data.

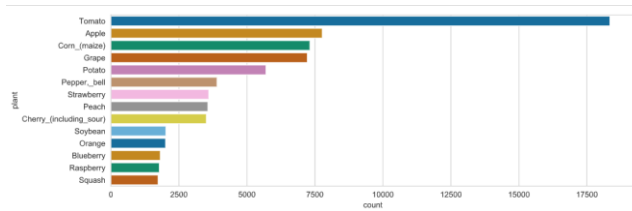


Figure 1: Number of images by plant

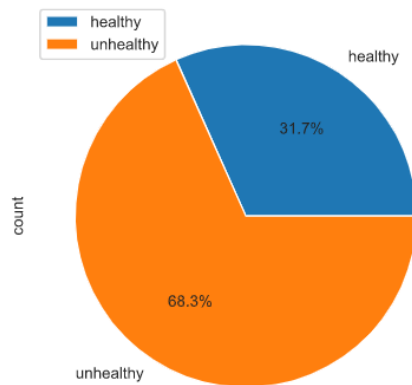


Figure 2: Relative image percentages by health status

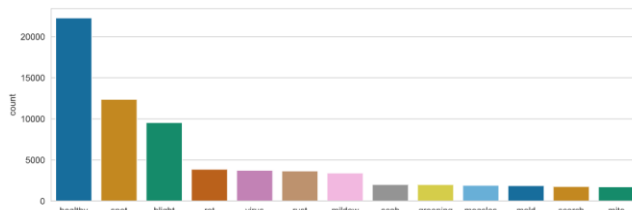


Figure 3: Number of images by disease

### 3.3 Data Preprocessing

The following data preprocessing methods will be evaluated to determine the best data augmentation for our input layer.

#### Data Augmentation

- Random Horizontal Shift
- Random Vertical Shift
- Random Horizontal Flip
- Random Vertical Flip
- Random Rotation
- Random Brightness
- Random Zoom

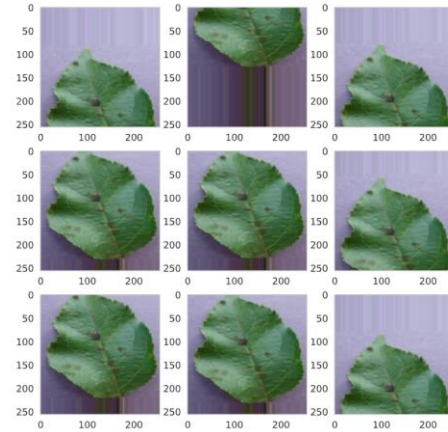


Figure 4: Data Augmentation Visualization for Random Vertical Shift

#### Data Pre-Processing

```
train_datagen = ImageDataGenerator(rescale=1./255,
                                   shear_range=0.2,
                                   zoom_range=0.2,
                                   width_shift_range=0.2,
                                   height_shift_range=0.2,
                                   fill_mode='nearest')

valid_datagen = ImageDataGenerator(rescale=1./255)

# batch_size = 128
batch_size = 48
training_set = train_datagen.flow_from_directory(dataDirTrain,
                                                  # target_size=(112, 112),
                                                  target_size=(224, 224),
                                                  batch_size=batch_size,
                                                  class_mode='categorical')

valid_set = valid_datagen.flow_from_directory(dataDirValidate,
                                              target_size=(224, 224),
                                              batch_size=batch_size,
                                              class_mode='categorical')

train_num = training_set.samples
valid_num = valid_set.samples
```

Found 1900 images belonging to 38 classes.  
Found 1900 images belonging to 38 classes.

Figure 5: Code Snippet For Data Preprocessing Steps

Table 2: Preprocessing Results Based on Baseline VGG16 Model with Transfer Learning

| Preprocessing Steps   | accuracy |
|---|----------|
| 0 rescale=1./255  | 0.81     |
| 1 rescale=1./255, shear_range=0.2   | 0.74     |
| 2 rescale=1./255, zoom_range=0.2  | 0.75     |
| 3 rescale=1./255, width_shift_range=0.2   | 0.74     |
| 4 rescale=1./255, width_shift_range=0.2   | 0.77     |
| 5 rescale=1./255, shear_range=0.2, zoom_range=0.2, width_shift_range=0.2, height_shift_range=0.2                                  | 0.71     |
| 6 rescale=1./255, horizontal_flip=True, vertical_flip=True, rotation_range=90, brightness_range=[0.2, 1.0], zoom_range=[0.5, 1.0] | 0.41     |

### 3.5 Model Evaluation & Selection

For the model evaluations, we will be running baselines (From scratch, with transfer learning, ...), Improving

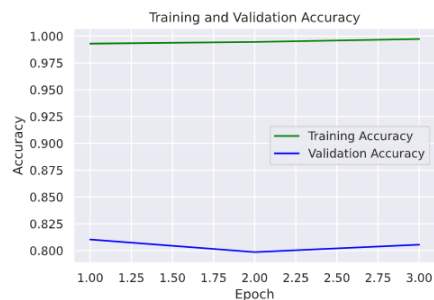
#### 3.5.1 Baseline Models

Several base models with transfer learning were run, using a sample of the complete dataset. The accuracy and loss were evaluated for each of the three base models: VGG16; ResNet50 and InceptionV3.

##### 3.5.1.1 VGG16 Base Model with Transfer Learning

| Model: "sequential"              |                   |          |
|----------------------------------|-------------------|----------|
| Layer (type)                     | Output Shape      | Param #  |
| vgg16 (Model)                    | (None, 7, 7, 512) | 14714688 |
| flatten (Flatten)                | (None, 25088)     | 0        |
| dense (Dense)                    | (None, 38)        | 953382   |
| Total params: 15,668,070         |                   |          |
| Trainable params: 953,382        |                   |          |
| Non-trainable params: 14,714,688 |                   |          |

| Preprocessing Steps accuracy |                                 |      |
|------------------------------|---------------------------------|------|
| 0                            | rescale=1./255                  | 0.81 |
| 1                            | rescale=1./255, shear_range=0.2 | 0.83 |
| 2                            | rescale=1./255, zoom_range=0.2  | 0.81 |



##### 3.5.1.2 ResNet50 Base Model with Transfer Learning

| Model: "sequential_1"            |                    |          |
|----------------------------------|--------------------|----------|
| Layer (type)                     | Output Shape       | Param #  |
| resnet50 (Model)                 | (None, 7, 7, 2048) | 23587712 |
| flatten_1 (Flatten)              | (None, 108352)     | 0        |
| dense_1 (Dense)                  | (None, 38)         | 3813414  |
| Total params: 27,481,126         |                    |          |
| Trainable params: 3,813,414      |                    |          |
| Non-trainable params: 23,587,712 |                    |          |

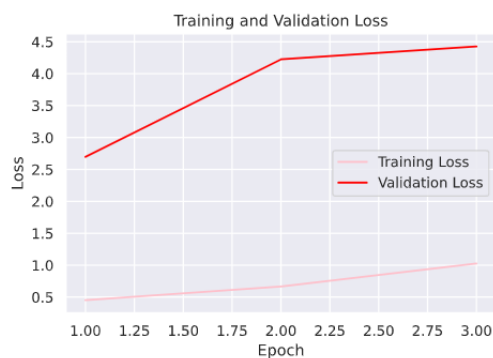
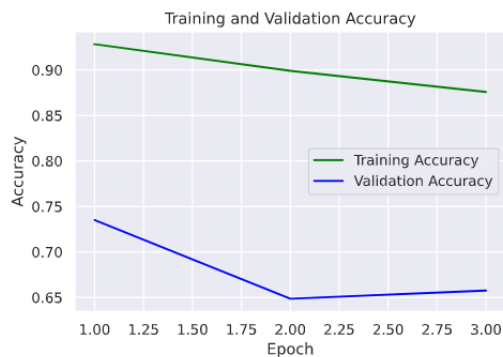
| Preprocessing Steps accuracy |                                 |      |
|------------------------------|---------------------------------|------|
| 0                            | rescale=1./255                  | 0.25 |
| 1                            | rescale=1./255, shear_range=0.2 | 0.29 |
| 2                            | rescale=1./255, zoom_range=0.2  | 0.29 |



### 3.5.1.3 InceptionV3 Base Model with Transfer Learning

| Model: "sequential_2"            |                    |          |
|----------------------------------|--------------------|----------|
| Layer (type)                     | Output Shape       | Param #  |
| inception_v3 (Model)             | (None, 5, 5, 2048) | 21882784 |
| flatten_2 (Flatten)              | (None, 51200)      | 0        |
| dense_2 (Dense)                  | (None, 38)         | 1945638  |
| Total params: 23,748,422         |                    |          |
| Trainable params: 1,945,638      |                    |          |
| Non-trainable params: 21,802,784 |                    |          |

|   | Preprocessing Steps             | accuracy |
|---|---------------------------------|----------|
| 0 | rescale=1./255                  | 0.74     |
| 1 | rescale=1./255, shear_range=0.2 | 0.74     |
| 2 | rescale=1./255, zoom_range=0.2  | 0.66     |



### 3.5.2 Improved Models

#### Hyperparameter Tuning

```
from keras.wrappers.scikit_learn import KerasRegressor
from sklearn.model_selection import RandomizedSearchCV, KFold, cross_val_score
from keras.optimizers import Adam
from keras import regularizers

# Create a KerasClassifier
model = KerasRegressor(build_fn = create_model, verbose = 0)

# Define the parameters to try out
params = {'activation': ['softmax', 'tanh'],
          'batch_size': [16, 32, 48, 64],
          'epochs': [3, 5, 10],
          'learning_rate': [0.01, 0.001, 0.0001]}

# Create a random search cv object passing in the parameters to try
random_search = RandomizedSearchCV(model,
                                   param_distributions = params,
                                   cv = KFold(10))

X_train, y_train = next(training_set)

random_search_results = random_search.fit(X_train, y_train)

# Print best score and parameters
print("Best Score: ", random_search_results.best_score_, "and Best Params: ", random_search_results.best_params_)
```

Best Paramaters: {'learning\_rate': 0.0001, 'epochs': 3, 'batch\_size': 32, 'activation': 'softmax'}

### 3.5.3 Final Model and Predictions

#### To Complete

## 4 Results

#### To Complete

## 5 Discussion

#### To Complete

## 6 Conclusion

#### To Complete

#### ACM Reference format:

FirstName Surname, FirstName Surname and FirstName Surname. 2018. Insert Your Title Here: Insert Subtitle Here. In *Proceedings of ACM Woodstock conference (WOODSTOCK'18)*. ACM, New York, NY, USA, 2 pages. <https://doi.org/10.1145/1234567890>

## ACKNOWLEDGMENTS

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## REFERENCES

- [1] Wiwid Setiawan, 2020. T Plant Disease Classification-VGG16 <https://www.kaggle.com/wiwidsetiawan/plant-disease-classification-vgg16>
- [2] Vimal Adit. 2019. Fork of Plant Diseases Classification Using inception3 <https://www.kaggle.com/vimaladit/fork-of-plant-diseases-classification-using-inception3>
- [3]