

```
In [ ]: import matplotlib.pyplot as plt
import numpy as np
import os
import PIL
import tensorflow as tf

from tensorflow import keras
from tensorflow.keras import layers
from tensorflow.keras.models import Sequential
```

```
In [ ]: import pathlib
dataset_url = "https://mo.columbari.us/static/images.tgz"
data_dir = tf.keras.utils.get_file(' ', origin=dataset_url, untar=True)
data_dir = pathlib.Path(data_dir)
```

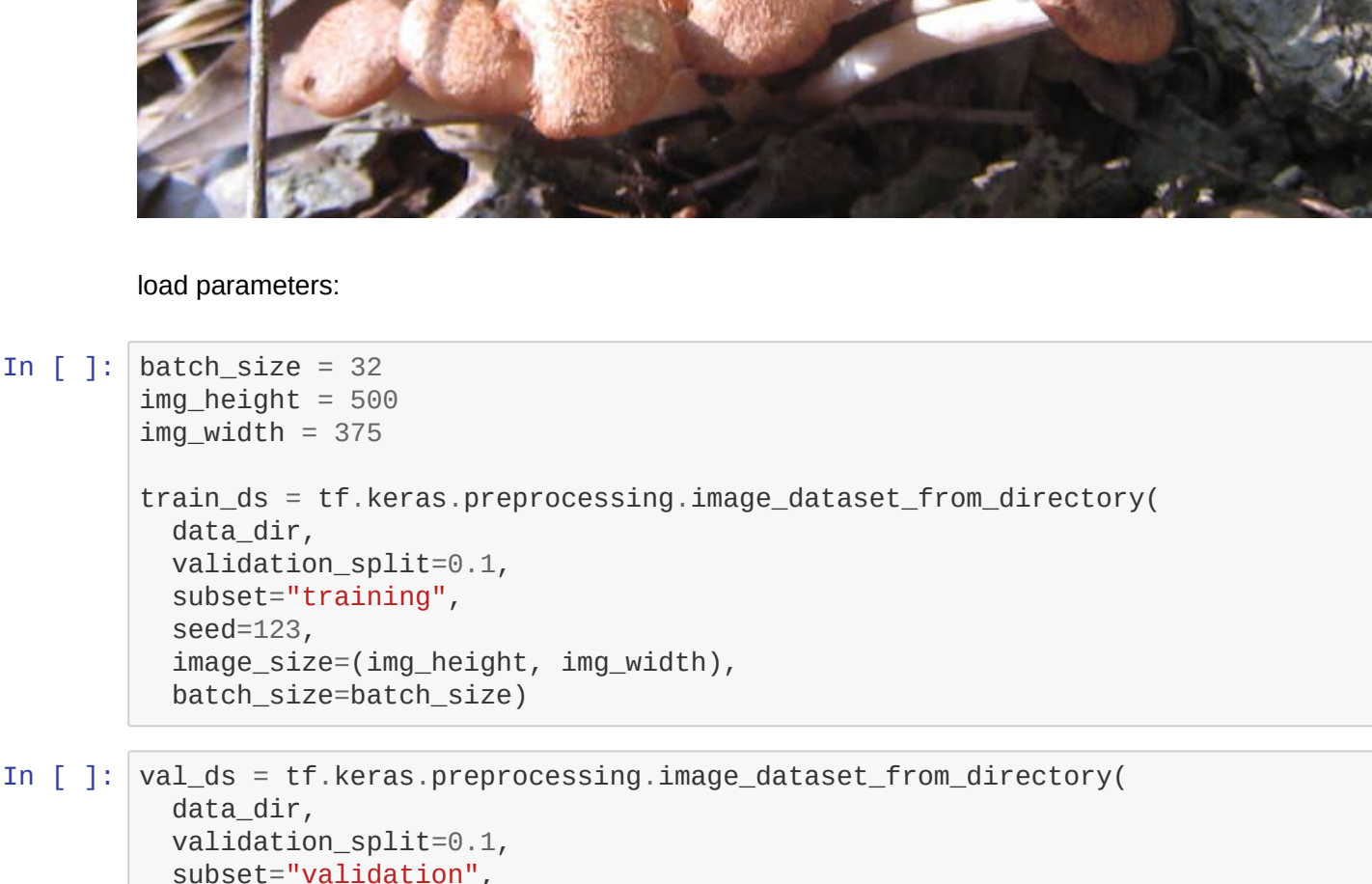
```
In [ ]: image_count = len(list(data_dir.glob('/**/*.jpg')))
print(image_count)

500
```

```
In [ ]: armillaria_tabescens = list(data_dir.glob('armillaria_tabescens/*'))
PIL.Image.open(str(armillaria_tabescens[0]))
```



```
In [ ]: PIL.Image.open(str(armillaria_tabescens[1]))
```



load parameters:

```
In [ ]: batch_size = 32
img_height = 500
img_width = 375

train_ds = tf.keras.preprocessing.image_dataset_from_directory(
    data_dir,
    validation_split=0.1,
    subset="training",
    seed=123,
    image_size=(img_height, img_width),
    batch_size=batch_size)
```

```
In [ ]: val_ds = tf.keras.preprocessing.image_dataset_from_directory(
    data_dir,
    validation_split=0.1,
    subset="validation",
    seed=123,
    image_size=(img_height, img_width),
    batch_size=batch_size)
```

Found 500 files belonging to 36 classes.
Using 50 files for validation.

```
In [ ]: class_names = train_ds.class_names
print(class_names)

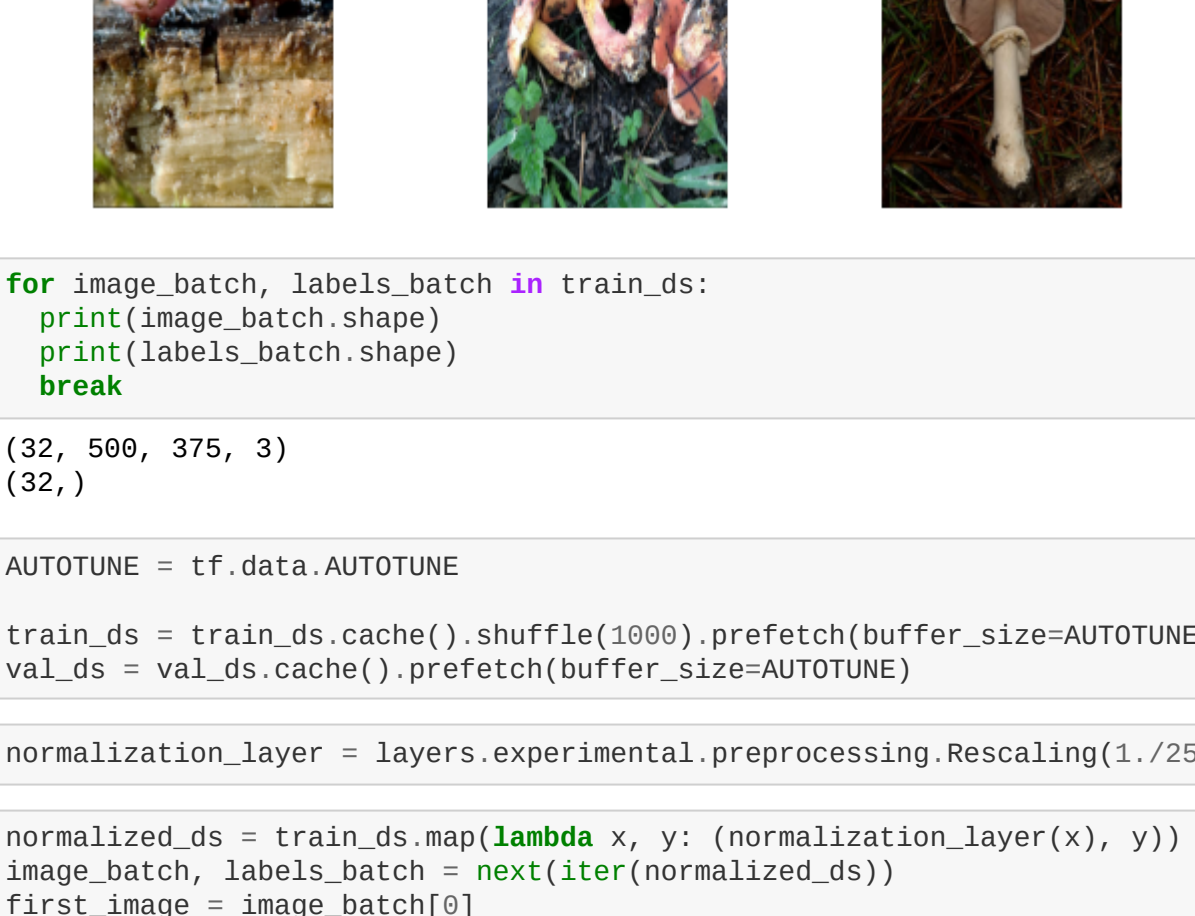
['agaricus_hondensis', 'amanita_volvata', 'armillaria_tabescens', 'ascocoryne_cylichnium', 'auriscalpium_andinum', 'biatora_chrysanthia', 'bisporella_resinicola', 'butyriboletus_floridanus', 'chlorosplenium_chloria', 'colpoma_chrysanthum', 'cortinarius_burlinghamiae', 'ductifera_pululahuana', 'exidiopsis_calcea', 'flammulaster_muricatus', 'gamundia_striatula', 'ganoderma_testaceum', 'gloeophyllum', 'hygrophorus_persoonii', 'hypocrepopsis_rhododendri', 'hypogymnia_duplicata', 'hypogymnia_schizidiata', 'lachnellula_calyciformis', 'lentinelus_cochleatus', 'leptogium_corticola', 'leptoporus_mollis', 'lobaria_lobata', 'muellerella_lichenicola', 'peltula_euploca', 'phaeotremella_foliacea', 'pholiota_chococensis', 'pholiota_tuberculosa', 'pluteus_longistriatus', 'rhizomarasmius_pyrrhocephalus', 'russula_moderata', 'thamnia_subuliformis', 'tuber_oligospermum']
```

Visualize the data

first 9 images from the training dataset:

```
In [ ]: import matplotlib.pyplot as plt

plt.figure(figsize=(10, 10))
for images, labels in train_ds.take(1):
    for i in range(9):
        ax = plt.subplot(3, 3, i + 1)
        plt.imshow(images[i].numpy().astype('uint8'))
        plt.title(class_names[labels[i]])
        plt.axis("off")
```



```
In [ ]: for image_batch, labels_batch in train_ds:
print(image_batch.shape)
print(labels_batch.shape)
break

(32, 500, 375, 3)
(32,)
```

```
In [ ]: AUTOTUNE = tf.data.AUTOTUNE

train_ds = train_ds.cache().shuffle(1000).prefetch(buffer_size=AUTOTUNE)
val_ds = val_ds.cache().prefetch(buffer_size=AUTOTUNE)
```

```
In [ ]: normalization_layer = layers.experimental.preprocessing.Rescaling(1./255)
```

```
In [ ]: normalized_ds = train_ds.map(lambda x, y: (normalization_layer(x), y))
image_batch, labels_batch = next(iter(normalized_ds))
first_image = image_batch[0]
# Notice the pixels values are now in [0,1].
print(np.min(first_image), np.max(first_image))

0.10051952 0.9915769
```

```
In [ ]: num_classes = 36

model = Sequential([
    layers.experimental.preprocessing.Rescaling(1./255, input_shape=(img_height, img_width, 3)),
    layers.Conv2D(16, 3, padding='same', activation='relu'),
    layers.MaxPooling2D(),
    layers.Conv2D(32, 3, padding='same', activation='relu'),
    layers.MaxPooling2D(),
    layers.Conv2D(64, 3, padding='same', activation='relu'),
    layers.MaxPooling2D(),
    layers.Flatten(),
    layers.Dense(128, activation='relu'),
    layers.Dense(num_classes)
])
```

Compile

```
In [ ]: model.compile(optimizer='adam',
                    loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),
                    metrics=['accuracy'])
```

```
In [ ]: Model: "sequential_1"

Layer (type) Output Shape Param #
=====
rescaling_3 (Rescaling) (None, 500, 375, 3) 0
conv2d_3 (Conv2D) (None, 500, 375, 16) 448
max_pooling2d_3 (MaxPooling2D) (None, 250, 187, 16) 0
conv2d_4 (Conv2D) (None, 250, 187, 32) 4640
max_pooling2d_4 (MaxPooling2D) (None, 125, 93, 32) 0
conv2d_5 (Conv2D) (None, 125, 93, 64) 18496
max_pooling2d_5 (MaxPooling2D) (None, 62, 46, 64) 0
flatten_1 (Flatten) (None, 182528) 0
dense_2 (Dense) (None, 128) 23363712
dense_3 (Dense) (None, 36) 4644
=====
Total params: 23,391,940
Trainable params: 23,391,940
Non-trainable params: 0
```

```
In [ ]: data_augmentation = keras.Sequential(
    [
        layers.experimental.preprocessing.RandomFlip("horizontal",
                                                    input_shape=(img_height,
                                                                img_width,
                                                                3)),
        layers.experimental.preprocessing.RandomRotation(0.1),
        layers.experimental.preprocessing.RandomZoom(0.1),
    ]
)
```

```
In [ ]: model = Sequential([
    data_augmentation,
    layers.experimental.preprocessing.Rescaling(1./255),
    layers.Conv2D(16, 3, padding='same', activation='relu'),
    layers.MaxPooling2D(),
    layers.Conv2D(32, 3, padding='same', activation='relu'),
    layers.MaxPooling2D(),
    layers.Conv2D(64, 3, padding='same', activation='relu'),
    layers.MaxPooling2D(),
    layers.Dropout(0.2),
    layers.Flatten(),
    layers.Dense(128, activation='relu'),
    layers.Dense(num_classes)
])
```

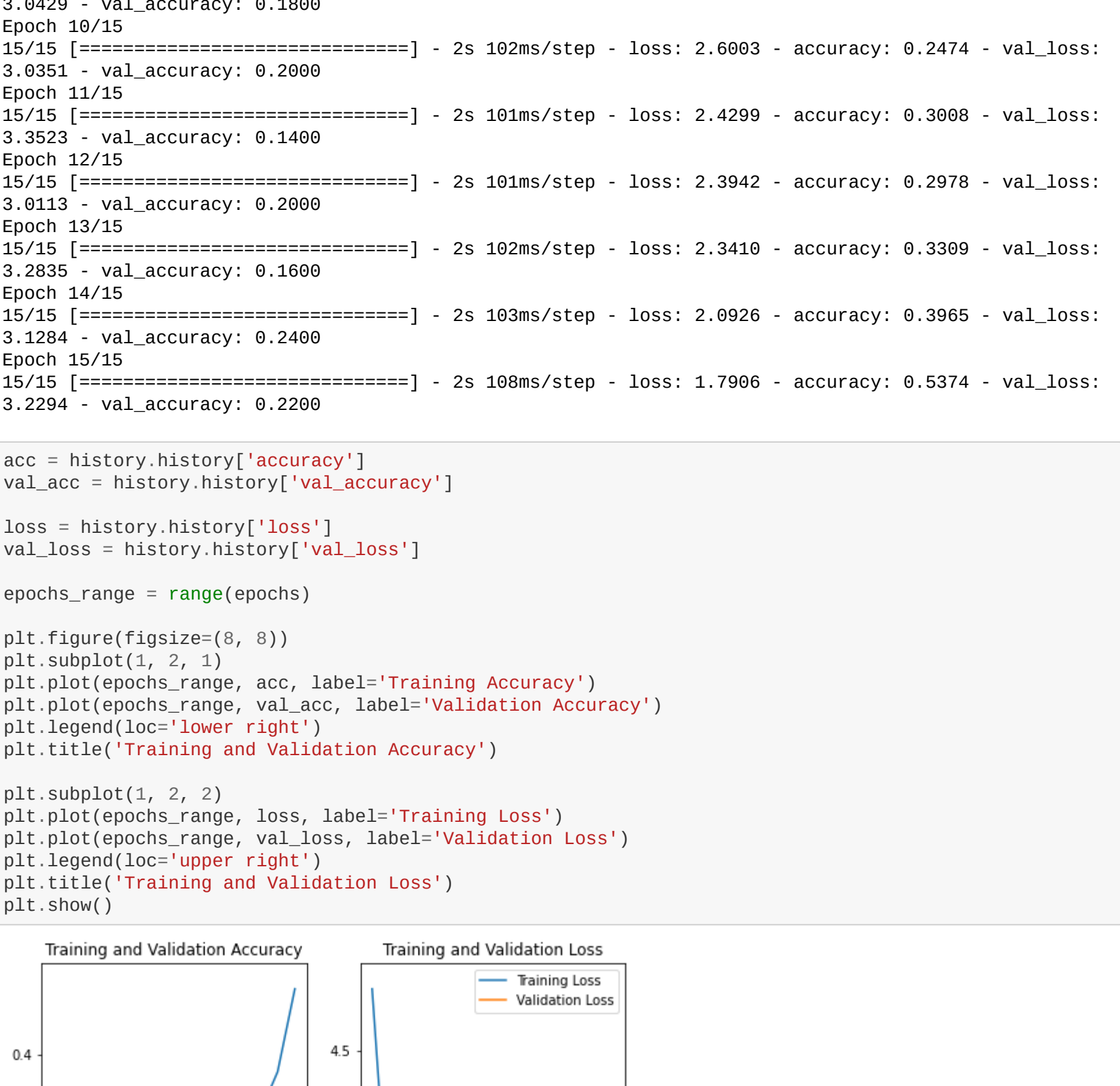
```
In [ ]: model.compile(optimizer='adam',
                    loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),
                    metrics=['accuracy'])
```

```
In [ ]: model.summary()

Model: "sequential_3"

Layer (type) Output Shape Param #
=====
sequential_2 (Sequential) (None, 500, 375, 3) 0
rescaling_4 (Rescaling) (None, 500, 375, 3) 0
conv2d_6 (Conv2D) (None, 500, 375, 16) 448
max_pooling2d_6 (MaxPooling2D) (None, 250, 187, 16) 0
conv2d_7 (Conv2D) (None, 250, 187, 32) 4640
max_pooling2d_7 (MaxPooling2D) (None, 125, 93, 32) 0
conv2d_8 (Conv2D) (None, 125, 93, 64) 18496
max_pooling2d_8 (MaxPooling2D) (None, 62, 46, 64) 0
dropout (Dropout) (None, 62, 46, 64) 0
flatten_2 (Flatten) (None, 182528) 0
dense_4 (Dense) (None, 128) 23363712
dense_5 (Dense) (None, 36) 4644
=====
Total params: 23,391,940
Trainable params: 23,391,940
Non-trainable params: 0
```

```
In [ ]: epochs = 15
history = model.fit(
    train_ds,
    validation_data=val_ds,
    epochs=epochs
)
```



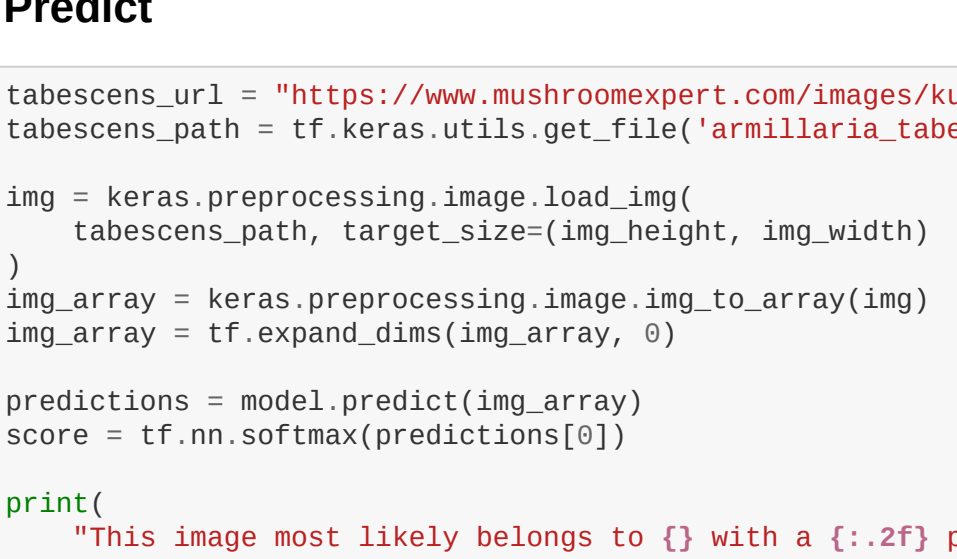
```
In [ ]: acc = history.history['accuracy']
val_acc = history.history['val_accuracy']

loss = history.history['loss']
val_loss = history.history['val_loss']

epochs_range = range(epochs)

plt.figure(figsize=(8, 8))
plt.subplot(1, 2, 1)
plt.plot(epochs_range, acc, label='Training Accuracy')
plt.plot(epochs_range, val_acc, label='Validation Accuracy')
plt.legend(loc='lower right')
plt.title('Training and Validation Accuracy')

plt.subplot(1, 2, 2)
plt.plot(epochs_range, loss, label='Training Loss')
plt.plot(epochs_range, val_loss, label='Validation Loss')
plt.legend(loc='upper right')
plt.title('Training and Validation Loss')
plt.show()
```



Predict

```
In [ ]: tabescens_url = "https://www.mushroomexpert.com/images/kuo6/armillaria_tabescens_06.jpg"
tabescens_path = tf.keras.utils.get_file('armillaria_tabescens_06', origin=tabescens_url)

img = keras.preprocessing.image.load_img(
    tabescens_path, target_size=(img_height, img_width)
)
img_array = keras.preprocessing.image.img_to_array(img)
img_array = tf.expand_dims(img_array, 0)

predictions = model.predict(img_array)
score = tf.nn.softmax(predictions[0])

print(
    "This image most likely belongs to {} with a {:.2f} percent confidence."
    .format(class_names[np.argmax(score)], 100 * np.max(score))
)
```

Downloading data from https://www.mushroomexpert.com/images/kuo6/armillaria_tabescens_06.jpg
90112/85752 [=====] - 0s 1us/step
This image most likely belongs to armillaria_tabescens with a 51.50 percent confidence.