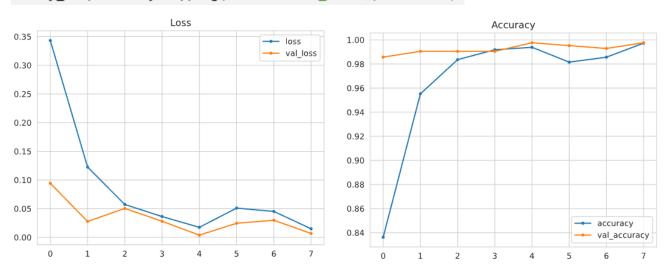
```
def create_tf_model():
   model = Sequential()
   model.add(Conv2D(filters=32, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
   model.add(MaxPooling2D(pool_size=(2, 2)))
   model.add(Conv2D(filters=64, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
   model.add(MaxPooling2D(pool_size=(2, 2)))
   model.add(Conv2D(filters=64, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
   model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Flatten())
   model.add(Dense(128, activation = 'relu'))
   model.add(Dropout(0.5))
   model.add(Dense(1, activation = 'sigmoid'))
   model.compile(loss='binary_crossentropy',
                  optimizer='adam',
                  metrics=['accuracy'])
   return model
```

early\_stop = EarlyStopping(monitor='val\_loss',patience=3)



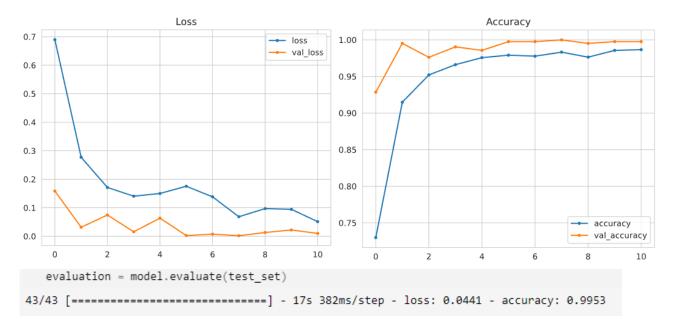
```
evaluation = model.evaluate(test_set)
43/43 [===========] - 16s 364ms/step - loss: 0.0039 - accuracy: 0.9976
```



0.9999896963781794 healthy

Model accuracy plot suggests slight overfitting may occur if further epochs were run as the accuracy and val\_accuracy are the same at the end of the curve.

```
def create_tf_model():
    model = Sequential()
    model.add(Conv2D(filters=32, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=64, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=64, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Flatten())
    model.add(Dense(128, activation = 'relu'))
   model.add(Dropout(0.5))
    model.add(Dense(1, activation = 'sigmoid'))
    model.compile(loss='binary_crossentropy',
                 optimizer='rmsprop',
                  metrics=['accuracy'])
    return model
```

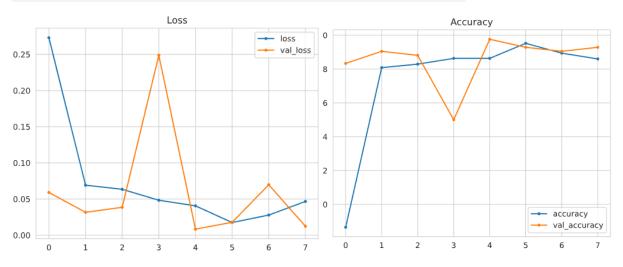




0.9999999817082621 healthy

Model accuracy plot suggests a normal fit. The loss models plot suggests that better learning is achievable.

```
def create_tf_model():
    model = Sequential()
   model.add(Conv2D(filters=32, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
   model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=64, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
   model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=64, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
   model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Flatten())
   model.add(Dense(128, activation = 'relu'))
    model.add(Dropout(0.5))
    model.add(Dense(2, activation = 'softmax'))
    model.compile(loss='categorical_crossentropy',
                  optimizer='adam',
                  metrics=['accuracy'])
    return model
```

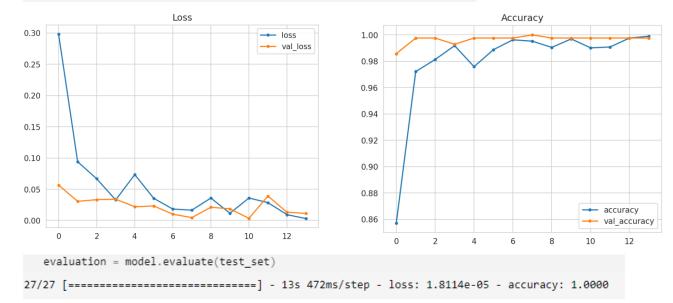




0.99985063 powdery mildew

The model plots suggest poor learning. The prediction for the image was reversed, indicating an issue with the code for evaluating the model.

```
def create_tf_model():
    model = Sequential()
    model.add(Conv2D(filters=16, kernel_size=(5,5),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=32, kernel_size=(4,4),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=64, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=32, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Flatten())
    model.add(Dense(64, activation = 'relu'))
    model.add(Dropout(0.3))
    model.add(Dense(1, activation = 'sigmoid'))
    model.compile(loss='binary_crossentropy',
                  optimizer='adam',
                  metrics=['accuracy'])
    return model
```





0.9999972517478 healthy

Model accuracy plot suggests slight overfitting as the loss and accuracy lines have crossed the corresponding val\_loss and val\_accuracy lines.

The first v6 created gave the following error during model evaluation:

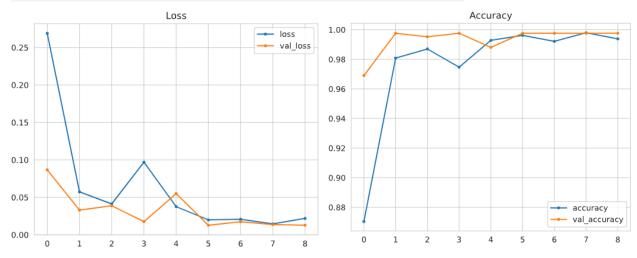
Load saved model

```
from keras.models import load_model
   model = load_model('outputs/v6/mildew_detector_model.h5')
                                                    Traceback (most recent call last)
Cell In[3], line 2
       1 from keras.models import load model
----> 2 model = load_model('outputs/v6/
File ~/.local/lib/python3.8/site-packages/keras/saving/save.py:205, in load model(filepath, custom objects, compile, options)
           filepath = path_to_string(filepath)
if isinstance(filepath, str):
    return saved_model_load.load(filepath, compile, options)
     204
 --> 205
     207 raise IOError(
           'Unable to load model. Filepath is not an hdf5 file (or h5py is not 'available) or SavedModel.')
     208
File ~/.local/lib/python3.8/site-packages/keras/saving/saved_model/load.py:108, in load(path, compile, options)
     103 # TODO(kathywu): Add saving/loading of optimizer, compiled losses and metrics.
104 # TODO(kathywu): Add code to load from objects that contain all endpoints
     105
     106 # Look for metadata file or parse the SavedModel
     107 metadata = saved_metadata pb2.SavedMetadata()
--> 108 meta_graph_def = tf.__internal__.saved_model.parse_saved_model(path).meta_graphs[0]
109 object_graph_def = meta_graph_def.object_graph_def
     110 path_to_metadata_pb = os.path.join(path, constants.SAVED_METADATA_PATH)
File ~/.local/lib/python3.8/site-packages/tensorflow/python/saved_model/loader_impl.py:118, in parse_saved_model(export_dir)
116 raise IOError("Cannot parse file %s: %s." % (path_to_pbtxt, str(e)))
     117 else:
--> 118 raise IOError(
119 "SavedMode
                  "SavedModel file does not exist at: %s%s{%s|%s}" %
             "SavedModel file does not exist at: %s%s{%s|%s}" % (export_dir, os.path.sep, constants.SAVED_MODEL_FILENAME_PBTXT,
     120
     121
                   {\tt constants.SAVED\_MODEL\_FILENAME\_PB))}
OSError: SavedModel file does not exist at: outputs/v6/mildew_detector_model.h5/{saved_model.pbtxt|saved_model.pb}
```

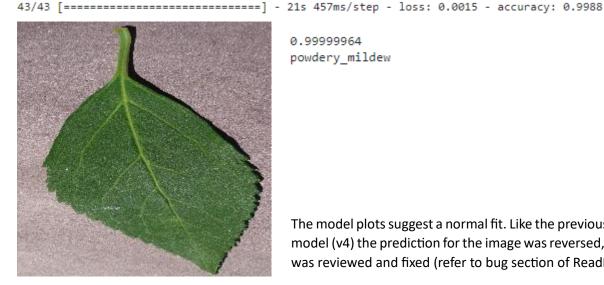
To test if the model was created satisfactorily, the model was tested on the dashboard which also gave an error.

Due to the error, this version was overwritten with the next model created.

```
def create_tf_model():
   model = Sequential()
   model.add(Conv2D(filters=32, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
   model.add(MaxPooling2D(pool_size=(2, 2)))
   model.add(Conv2D(filters=16, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
   model.add(MaxPooling2D(pool_size=(2, 2)))
   model.add(Conv2D(filters=8, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
   model.add(MaxPooling2D(pool_size=(2, 2)))
   model.add(Flatten())
   model.add(Dense(64, activation = 'relu'))
   model.add(Dropout(0.3))
   model.add(Dense(2, activation = 'softmax'))
   model.compile(loss='categorical_crossentropy',
                  optimizer='adam',
                  metrics=['accuracy'])
    return model
```



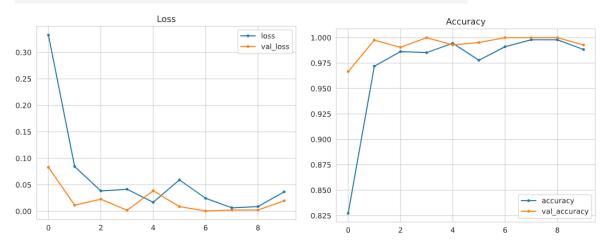
```
evaluation = model.evaluate(test_set)
```



0.99999964 powdery mildew

The model plots suggest a normal fit. Like the previous softmax model (v4) the prediction for the image was reversed, the code was reviewed and fixed (refer to bug section of ReadMe).

```
def create_tf_model():
    model = Sequential()
    model.add(Conv2D(filters=32, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=16, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=16, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Flatten())
    model.add(Dense(64, activation = 'relu'))
    model.add(Dropout(0.3))
    model.add(Dense(2, activation = 'softmax'))
    model.compile(loss='categorical_crossentropy',
                 optimizer='adam',
                  metrics=['accuracy'])
    return model
```

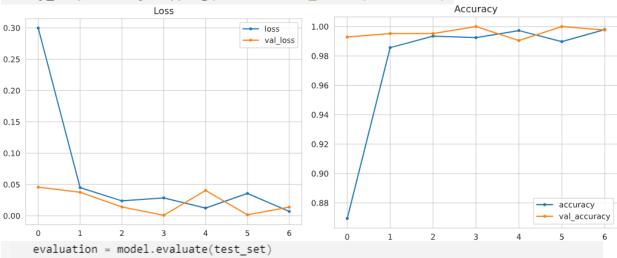




0.9999974 healthy

The model plots suggest slight overfitting.

```
def create_tf_model():
    model = Sequential()
    model.add(Conv2D(filters=16, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=16, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Conv2D(filters=16, kernel_size=(3,3),input_shape=image_shape, activation='relu',))
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Flatten())
    model.add(Dense(128, activation = 'relu'))
    model.add(Dropout(0.25))
    model.add(Dense(2, activation = 'softmax'))
    model.compile(loss='categorical_crossentropy',
                 optimizer='adam',
                  metrics=['accuracy'])
    return model
```





0.9737485386431217 Infected

The model plots suggest slight overfitting.