

CNN With Custom Images: Malaria Cell Images Dataset

The dataset contains 2 folders - Infected - Uninfected

And a total of 27,558 images.

This Dataset was taken from the official NIH Website: <https://ceb.nlm.nih.gov/repositories/malaria-datasets/>

```
In [2]: import os
import pandas as pd
import tensorflow as tf
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
from matplotlib.image import imread
```

```
In [4]: my_data= '/Users/usameturker/Desktop/DS/DL/cell_images/'
```

```
In [5]: os.listdir(my_data)
```

```
Out[5]: ['.DS_Store', 'test', 'train']
```

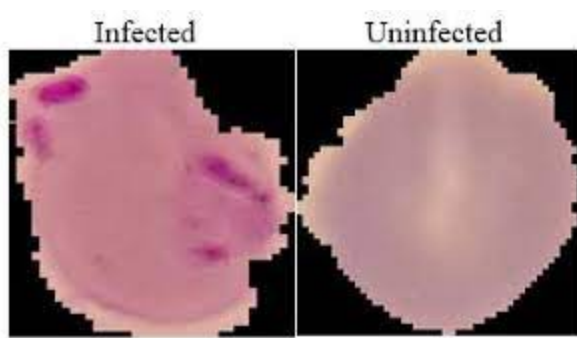
```
In [6]: train_path = my_data+'train/'
test_path = my_data+'test/'
```

```
In [7]: os.listdir(train_path)
```

```
Out[7]: ['.DS_Store', 'parasitized', 'uninfected']
```

```
In [8]: os.listdir(test_path)
```

```
Out[8]: ['.DS_Store', 'parasitized', 'uninfected']
```



```
In [9]: os.listdir(train_path+'parasitized')[:30]
```

```
Out[9]: ['C189P150ThinF_IMG_20151203_142224_cell_84.png',
'C91P52ThinF_IMG_20150821_123116_cell_189.png',
'C84P45ThinF_IMG_20150818_101226_cell_98.png',
'C144P105ThinF_IMG_20151015_163432_cell_310.png',
'C136P97ThinF_IMG_20151005_141803_cell_131.png',
'C60P21thinF_IMG_20150804_105034_cell_105.png',
'C176P137NThinF_IMG_20151201_122708_cell_126.png',
'C136P97ThinF_IMG_20151005_142437_cell_119.png',
'C173P134NThinF_IMG_20151130_115733_cell_227.png',
'C85P46ThinF_IMG_20150820_105154_cell_221.png',
'C101P62ThinF_IMG_20150918_151507_cell_49.png',
'C116P77ThinF_IMG_20150930_171844_cell_100.png',
'C48P9thinF_IMG_20150721_160406_cell_247.png',
'C184P145ThinF_IMG_20151203_103114_cell_149.png',
'C99P60ThinF_IMG_20150918_141620_cell_56.png',
'C132P93ThinF_IMG_20151004_152045_cell_111.png',
'C68P29N_ThinF_IMG_20150819_134112_cell_153.png',
'C116P77ThinF_IMG_20150930_171844_cell_114.png',
'C171P132ThinF_IMG_20151119_153150_cell_222.png',
'C176P137NThinF_IMG_20151201_122708_cell_132.png',
'C175P136NThinF_IMG_20151127_141325_cell_219.png',
'C132P93ThinF_IMG_20151004_151733_cell_140.png',
'C173P134NThinF_IMG_20151130_125501_cell_255.png',
'C60P21thinF_IMG_20150804_105034_cell_111.png',
'C118P79ThinF_IMG_20151002_105018_cell_144.png',
'C51AP12thinF_IMG_20150724_153313_cell_109.png',
'C189P150ThinF_IMG_20151203_142224_cell_90.png',
'C144P105ThinF_IMG_20151015_163432_cell_304.png',
'C136P97ThinF_IMG_20151005_141803_cell_125.png',
'C65P26N_ThinF_IMG_20150818_154010_cell_200.png']
```

```
In [10]: os.listdir(train_path+'parasitized')[17]
```

```
Out[10]: 'C116P77ThinF_IMG_20150930_171844_cell_114.png'
```

```
In [11]: para_cell = train_path+'parasitized'+'/C100P61ThinF_IMG_20150918_144823_cell_158.png'
```

```
In [12]: imread(para_cell)
```

```
Out[12]: array([[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]],

               [[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]],

               [[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]],

               ...,

               [[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]],

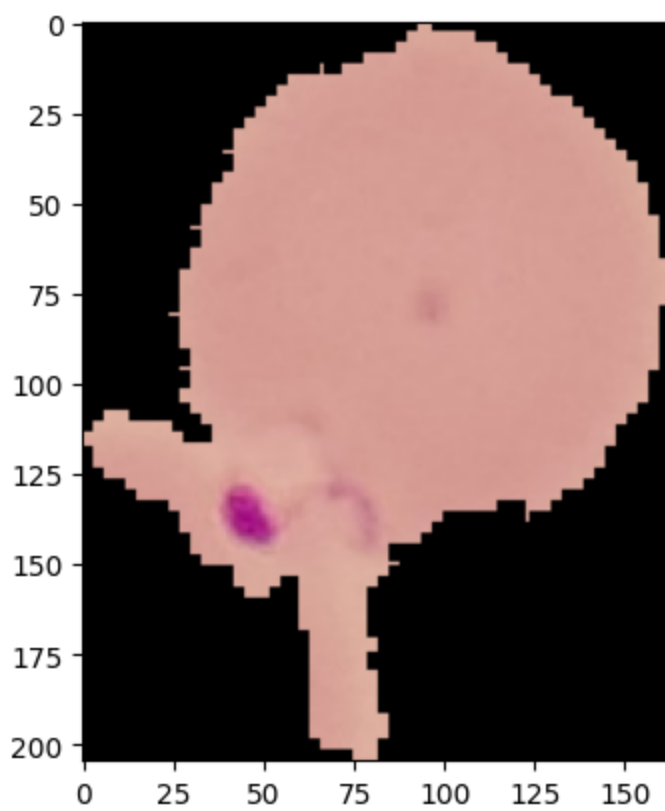
               [[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]],

               [[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]]], dtype=float32)
```

```
In [13]: para_img= imread(para_cell)
```

```
In [14]: plt.imshow(para_img)
```

```
Out[14]: <matplotlib.image.AxesImage at 0x282da2890>
```



```
In [15]: para_img.shape
```

```
Out[15]: (205, 163, 3)
```

```
In [16]: os.listdir(train_path+'uninfected')[:20]
```

```
Out[16]: ['C13NThinF_IMG_20150614_131318_cell_179.png',
'C205ThinF_IMG_20151106_152003_cell_21.png',
'C230ThinF_IMG_20151112_150647_cell_1.png',
'C134P95ThinF_IMG_20151005_121834_cell_65.png',
'C141P102ThinF_IMG_20151005_214836_cell_132.png',
'C182P143NThinF_IMG_20151201_171950_cell_93.png',
'C181P142ThinF_IMG_20151127_160616_cell_54.png',
'C142P103ThinF_IMG_20151005_223257_cell_18.png',
'C80P41ThinF_IMG_20150817_110608_cell_62.png',
'C144P105ThinF_IMG_20151015_163142_cell_66.png',
'C12NThinF_IMG_20150614_124212_cell_187.png',
'C223ThinF_IMG_20151112_104708_cell_190.png',
'C143P104ThinF_IMG_20151005_225746_cell_133.png',
'C84P45ThinF_IMG_20150818_101056_cell_87.png',
'C222ThinF_IMG_20151115_150925_cell_127.png',
'C1_thinF_IMG_20150604_104919_cell_82.png',
'C150P111ThinF_IMG_20151115_115950_cell_53.png',
'C170P131ThinF_IMG_20151119_120233_cell_183.png',
'C13NThinF_IMG_20150614_131529_cell_168.png',
'C1_thinF_IMG_20150604_104919_cell_96.png']
```

```
In [17]: uninfected_cell_path = train_path+'uninfected/'+os.listdir(train_path+'uninfected')[7]
```

```
In [18]: uninfected_cell_path
```

```
Out[18]: '/Users/usameturker/Desktop/DS/DL/cell_images//train/uninfected/C142P103ThinF_IMG_20151005_223257_cell_18.png'
```

```
In [19]: imread(uninfected_cell_path)
```

```

Out[19]: array([[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]],

               [[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]],

               [[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]],

               ...,

               [[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]],

               [[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]],

               [[0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.],
                ...,
                [0., 0., 0.],
                [0., 0., 0.],
                [0., 0., 0.]]], dtype=float32)

```

```

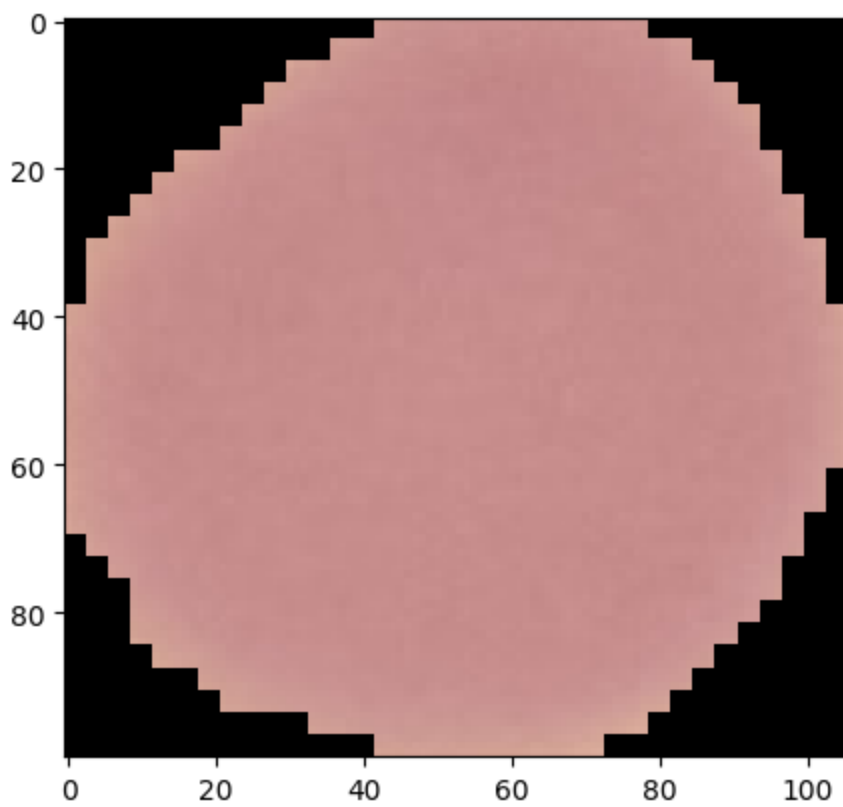
In [20]: uninfected_cell = imread(uninfected_cell_path)
         plt.imshow(uninfected_cell)

```

```

Out[20]: <matplotlib.image.AxesImage at 0x282e9b220>

```



```
In [21]: len(os.listdir(train_path+'parasitized'))
```

```
Out[21]: 12480
```

```
In [22]: len(os.listdir(train_path+'uninfected'))
```

```
Out[22]: 12480
```

```
In [23]: uninfected_cell.shape
```

```
Out[23]: (100, 106, 3)
```

```
In [24]: para_img.shape
```

```
Out[24]: (205, 163, 3)
```

```
In [25]: x = []  
y = []  
for image in os.listdir(test_path+'uninfected'):  
    img = imread(test_path+'uninfected/'+image)  
    d1,d2,_ = img.shape  
    x.append(d1)  
    y.append(d2)
```

```
In [26]: y[:20]
```

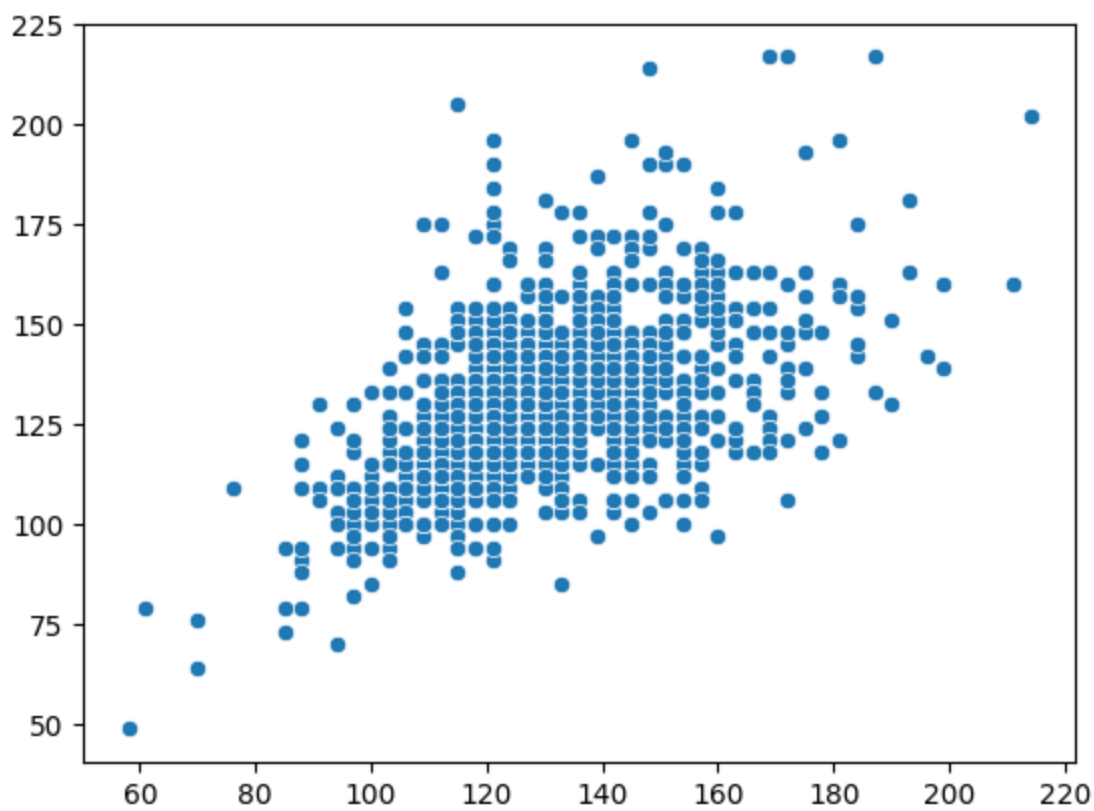
```
Out [26]: [127,  
          121,  
          151,  
          130,  
          124,  
          127,  
          136,  
          124,  
          124,  
          130,  
          127,  
          115,  
          154,  
          115,  
          100,  
          121,  
          151,  
          142,  
          118,  
          172]
```

```
In [27]: x[:20]
```

```
Out [27]: [127,  
          109,  
          127,  
          118,  
          127,  
          118,  
          133,  
          115,  
          133,  
          142,  
          139,  
          121,  
          124,  
          133,  
          103,  
          148,  
          130,  
          118,  
          127,  
          142]
```

```
In [29]: sns.scatterplot(x=x,y=y)
```

```
Out [29]: <Axes: >
```



```
In [30]: np.mean(x)
```

```
Out[30]: 130.92538461538462
```

```
In [31]: np.mean(y)
```

```
Out[31]: 130.75
```

```
In [32]: image_shape = (130,130,3)
```

Preparing the Data for the model

```
In [33]: uninfected_cell.max()
```

```
Out[33]: 0.85490197
```

```
In [34]: para_img.max()
```

```
Out[34]: 0.8980392
```

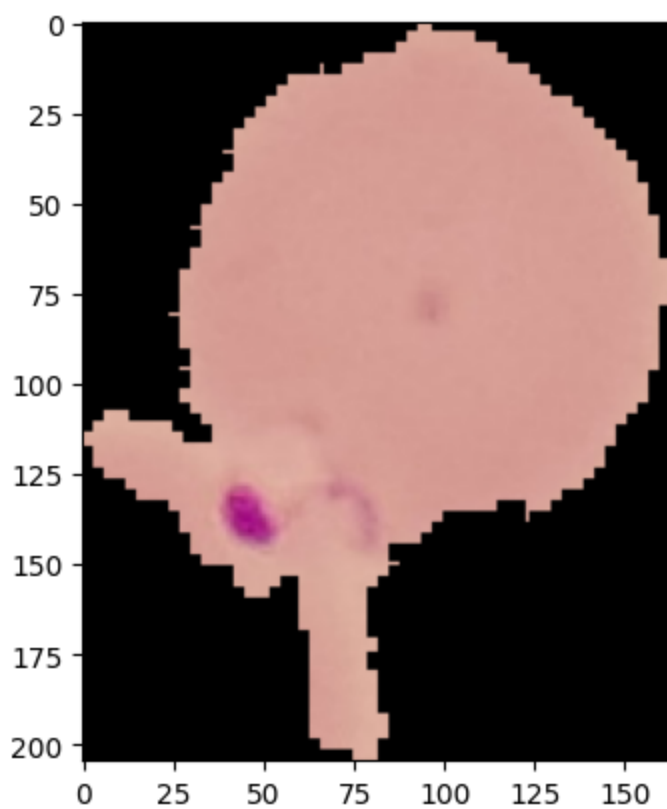
```
In [35]: from tensorflow.keras.preprocessing.image import ImageDataGenerator
```

```
In [93]: #help(ImageDataGenerator)
```

[illegible]

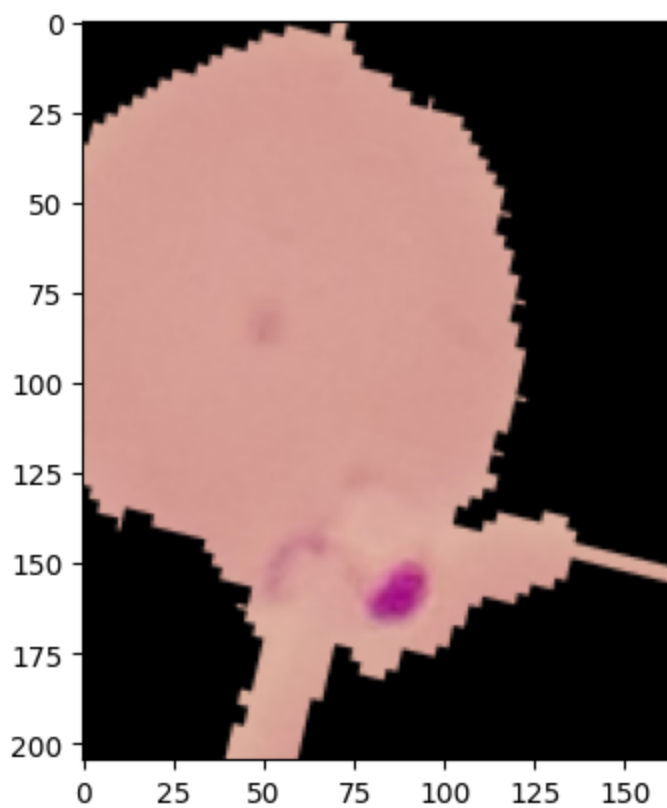

```
In [37]: plt.imshow(para_img)
```

```
Out[37]: <matplotlib.image.AxesImage at 0x2830f00a0>
```



```
In [38]: plt.imshow(image_gen.random_transform(para_img))
```

```
Out[38]: <matplotlib.image.AxesImage at 0x28444df00>
```



```
In [39]: image_gen.flow_from_directory(train_path)
```

Found 24958 images belonging to 2 classes.

Out [39]: <keras.preprocessing.image.DirectoryIterator at 0x2844d8d60>

In [40]: `image_gen.flow_from_directory(test_path)`

Found 2600 images belonging to 2 classes.

Out [40]: <keras.preprocessing.image.DirectoryIterator at 0x2844d9ab0>

Creating the Model

In [41]: `from tensorflow.keras.models import Sequential`
`from tensorflow.keras.layers import Activation, Dropout, Flatten, Dense, Conv2D, MaxPool`

In [42]: `model = Sequential()`

```
model.add(Conv2D(filters=64, kernel_size=(3,3), input_shape=image_shape, padding='same',
model.add(MaxPooling2D(pool_size=(2, 2)))

model.add(Conv2D(filters=128, kernel_size=(3,3), padding='same', activation='relu',))
model.add(Conv2D(filters=128, kernel_size=(3,3), padding='same', activation='relu',))
model.add(MaxPooling2D(pool_size=(2, 2)))

model.add(Conv2D(filters=256, kernel_size=(3,3), padding='same', activation='relu',))
model.add(Conv2D(filters=256, kernel_size=(3,3), padding='same', activation='relu',))
model.add(Conv2D(filters=256, kernel_size=(3,3), padding='same', activation='relu',))
model.add(MaxPooling2D(pool_size=(2, 2)))

model.add(Conv2D(filters=128, kernel_size=(3,3), padding='same', activation='relu',))
model.add(MaxPooling2D(pool_size=(2, 2)))

model.add(Conv2D(filters=64, kernel_size=(3,3), padding='same', activation='relu',))
model.add(MaxPooling2D(pool_size=(2, 2)))

model.add(Flatten())

model.add(Dense(128))
model.add(Activation('relu'))

model.add(Dropout(0.5))

model.add(Dense(1))
model.add(Activation('sigmoid'))

model.compile(loss='binary_crossentropy',
              optimizer='adam',
              metrics=['accuracy'])
```

In [43]: `model.summary()`

Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 130, 130, 64)	1792
max_pooling2d (MaxPooling2D)	(None, 65, 65, 64)	0
conv2d_1 (Conv2D)	(None, 65, 65, 128)	73856
conv2d_2 (Conv2D)	(None, 65, 65, 128)	147584
max_pooling2d_1 (MaxPooling2D)	(None, 32, 32, 128)	0
conv2d_3 (Conv2D)	(None, 32, 32, 256)	295168
conv2d_4 (Conv2D)	(None, 32, 32, 256)	590080
conv2d_5 (Conv2D)	(None, 32, 32, 256)	590080
max_pooling2d_2 (MaxPooling2D)	(None, 16, 16, 256)	0
conv2d_6 (Conv2D)	(None, 16, 16, 128)	295040
max_pooling2d_3 (MaxPooling2D)	(None, 8, 8, 128)	0
conv2d_7 (Conv2D)	(None, 8, 8, 64)	73792
max_pooling2d_4 (MaxPooling2D)	(None, 4, 4, 64)	0
flatten (Flatten)	(None, 1024)	0
dense (Dense)	(None, 128)	131200
activation (Activation)	(None, 128)	0
dropout (Dropout)	(None, 128)	0
dense_1 (Dense)	(None, 1)	129
activation_1 (Activation)	(None, 1)	0
Total params: 2,198,721		
Trainable params: 2,198,721		
Non-trainable params: 0		

Early Stopping

```
In [44]: from tensorflow.keras.callbacks import EarlyStopping
```

```
In [45]: early_stop = EarlyStopping(monitor='val_loss',patience=5)
```

Training the Model

```
In [46]: batch_size = 32
train_image_gen = image_gen.flow_from_directory(train_path,
                                                target_size=(130,130),
                                                color_mode='rgb',
                                                batch_size=batch_size,
                                                class_mode='binary', shuffle=True, seed=4
```

Found 24958 images belonging to 2 classes.

```
In [47]: test_image_gen = image_gen.flow_from_directory(test_path,
                                                        target_size=(130,130),
                                                        color_mode='rgb',
                                                        batch_size=batch_size,
                                                        class_mode='binary', shuffle=True, seed=4
```

Found 2600 images belonging to 2 classes.

```
In [48]: train_image_gen.class_indices
```

```
Out[48]: {'parasitized': 0, 'uninfected': 1}
```

```
In [49]: test_image_gen.class_indices
```

```
Out[49]: {'parasitized': 0, 'uninfected': 1}
```

```
In [ ]: results = model.fit(train_image_gen, epochs=20,
                             validation_data=test_image_gen,
                             callbacks=[early_stop])
```

Epoch 1/20

780/780 [=====] - 1478s 2s/step - loss: 0.6961 - accuracy: 0.5082 - val_loss: 0.6931 - val_accuracy: 0.5000

Epoch 2/20

780/780 [=====] - 1458s 2s/step - loss: 0.6924 - accuracy: 0.5193 - val_loss: 0.6942 - val_accuracy: 0.4719

Epoch 3/20

780/780 [=====] - 1480s 2s/step - loss: 0.6925 - accuracy: 0.5335 - val_loss: 0.6888 - val_accuracy: 0.5238

Epoch 4/20

780/780 [=====] - 1710s 2s/step - loss: 0.6443 - accuracy: 0.6068 - val_loss: 0.3118 - val_accuracy: 0.9088

Epoch 5/20

780/780 [=====] - 1502s 2s/step - loss: 0.2005 - accuracy: 0.9423 - val_loss: 0.1772 - val_accuracy: 0.9458

Epoch 6/20

780/780 [=====] - 1507s 2s/step - loss: 0.1729 - accuracy: 0.9486 - val_loss: 0.1724 - val_accuracy: 0.9496

Epoch 7/20

780/780 [=====] - 1489s 2s/step - loss: 0.1604 - accuracy: 0.9512 - val_loss: 0.1640 - val_accuracy: 0.9504

Epoch 8/20

780/780 [=====] - 4497s 6s/step - loss: 0.1532 - accuracy: 0.9526 - val_loss: 0.1560 - val_accuracy: 0.9504

Epoch 9/20

95/780 [==>.....] - ETA: 3:15:47 - loss: 0.1424 - accuracy: 0.959