Import Statements

```
In [4]:
        import os
        import keras
        from keras.models import Sequential
        from keras.layers import Conv2D, MaxPooling2D, Flatten, Dense, Dropout, BatchNormal
        from PIL import Image
        import numpy as np
        import pandas as pd
        import matplotlib.pyplot as plt
        plt.style.use('dark_background')
        from sklearn.model_selection import train_test_split
        from sklearn.preprocessing import OneHotEncoder
In [2]: !pip install keras
        Requirement already satisfied: keras in c:\users\arivu\appdata\local\programs\pyth
        on\python310\lib\site-packages (2.10.0)
        [notice] A new release of pip available: 22.3 -> 22.3.1
        [notice] To update, run: python.exe -m pip install --upgrade pip
```

One Hot Encoding the Target Classes

```
In [5]: encoder = OneHotEncoder()
  encoder.fit([[0], [1]])

# 0 - cancer
# 1 - normal

Out[5]: OneHotEncoder()
```

Creating 3 Important Lists --

- 1. data list for storing image data in numpy array form
- 2. paths list for storing paths of all images
- 3. result list for storing one hot encoded form of target class whether normal or cancer

```
In [6]:
    data = []
    paths = []
    result = []

for r, d, f in os.walk(r"C:\Users\arivu\OneDrive\Dataset_vani"):
        for file in f:
            if '.bmp' in file:
                 paths.append(os.path.join(r, file))

for path in paths:
    img = Image.open(path)
    img = img.resize((128,128))
    img = np.array(img)
    if(img.shape == (128,128,3)):
        data.append(np.array(img))
        result.append(encoder.transform([[0]]).toarray())
```

```
In [8]:
         paths = []
         for r, d, f in os.walk(r"C:\Users\arivu\OneDrive\Dataset vani"):
             for file in f:
                 if '.bmp' in file:
                     paths.append(os.path.join(r, file))
         for path in paths:
             img = Image.open(path)
             img = img.resize((128,128))
             img = np.array(img)
             if(img.shape == (128,128,3)):
                 data.append(np.array(img))
                  result.append(encoder.transform([[1]]).toarray())
In [9]: data = np.array(data)
         data.shape
         (1365, 128, 128, 3)
Out[9]:
In [11]:
         result = np.array(result)
         result = result.reshape(1365,2)
```

Splitting the Data into Training & Testing

```
In [12]: x_train,x_test,y_train,y_test = train_test_split(data, result, test_size=0.2, shuf-
In [13]: # Normalization
    x_train = x_train/255.0
    x_test = x_test/255.0
```

Model Building

Batch normalization is a technique for training very deep neural networks that standardizes the inputs to a layer for each mini-batch. This has the effect of stabilizing the learning process and dramatically reducing the number of training epochs required to train deep networks.

```
In [14]: model = Sequential()
    model.add(Conv2D(32, kernel_size=(2, 2), input_shape=(128, 128, 3), padding = 'Same' model.add(Conv2D(32, kernel_size=(2, 2), activation = 'relu', padding = 'Same'))
    model.add(BatchNormalization())
    model.add(MaxPooling2D(pool_size=(2, 2)))
    model.add(Dropout(0.25))

model.add(Conv2D(64, kernel_size = (2,2), activation = 'relu', padding = 'Same'))
    model.add(Conv2D(64, kernel_size = (2,2), activation = 'relu', padding = 'Same'))
    model.add(BatchNormalization())
    model.add(MaxPooling2D(pool_size=(2,2), strides=(2,2)))
    model.add(Dropout(0.25))

model.add(Flatten())
```

```
model.add(Dense(512, activation='relu'))
model.add(Dropout(0.5))
model.add(Dense(2, activation='softmax'))

model.compile(loss = "categorical_crossentropy", optimizer='Adamax')
print(model.summary())
```

Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 128, 128, 32)	416
conv2d_1 (Conv2D)	(None, 128, 128, 32)	4128
<pre>batch_normalization (BatchN ormalization)</pre>	(None, 128, 128, 32)	128
<pre>max_pooling2d (MaxPooling2D)</pre>	(None, 64, 64, 32)	0
dropout (Dropout)	(None, 64, 64, 32)	0
conv2d_2 (Conv2D)	(None, 64, 64, 64)	8256
conv2d_3 (Conv2D)	(None, 64, 64, 64)	16448
<pre>batch_normalization_1 (Batc hNormalization)</pre>	(None, 64, 64, 64)	256
<pre>max_pooling2d_1 (MaxPooling 2D)</pre>	(None, 32, 32, 64)	0
dropout_1 (Dropout)	(None, 32, 32, 64)	0
flatten (Flatten)	(None, 65536)	0
dense (Dense)	(None, 512)	33554944
dropout_2 (Dropout)	(None, 512)	0
dense_1 (Dense)	(None, 2)	1026

Total params: 33,585,602 Trainable params: 33,585,410 Non-trainable params: 192

None

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

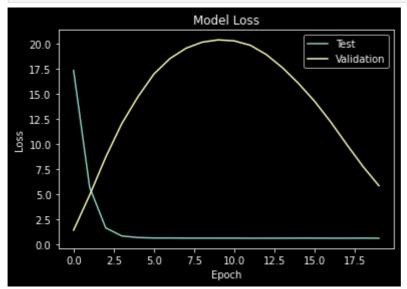
Out[23]: (4004, 2)

```
In [15]: history = model.fit(x_train, y_train, epochs = 20, batch_size = 40, verbose = 1,val
```

```
Epoch 1/20
1.4399
Epoch 2/20
28/28 [============ ] - 49s 2s/step - loss: 5.7070 - val loss: 4.
9389
Epoch 3/20
28/28 [============ ] - 52s 2s/step - loss: 1.6675 - val_loss: 8.
7149
Epoch 4/20
2.0532
Epoch 5/20
4.7119
Epoch 6/20
6.9932
Epoch 7/20
8.5327
Epoch 8/20
9.5581
Epoch 9/20
0.1596
Epoch 10/20
0.3900
Epoch 11/20
0.2893
Epoch 12/20
9.8524
Epoch 13/20
8.9206
Epoch 14/20
7.6169
Epoch 15/20
6.0451
Epoch 16/20
4.2701
Epoch 17/20
2.2198
Epoch 18/20
28/28 [============ ] - 40s 1s/step - loss: 0.6405 - val_loss: 9.
9718
Epoch 19/20
8009
Epoch 20/20
28/28 [============== ] - 40s 1s/step - loss: 0.6396 - val loss: 5.
8662
```

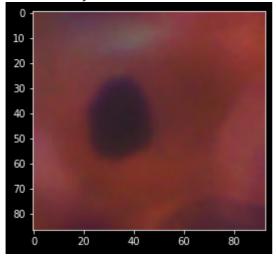
Plotting Losses

```
In [16]: plt.plot(history.history['loss'])
    plt.plot(history.history['val_loss'])
    plt.title('Model Loss')
    plt.ylabel('Loss')
    plt.xlabel('Epoch')
    plt.legend(['Test', 'Validation'], loc='upper right')
    plt.show()
```



Just Checking the Model

Affected by Cervical Cancer



In []: