

```
In [3]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from PIL import Image
from glob import glob

from sklearn.model_selection import train_test_split
from sklearn import metrics

import cv2
import gc
import os

import tensorflow as tf
from tensorflow import keras
from keras import layers

import warnings
warnings.filterwarnings('ignore')
```

```
In [4]: from zipfile import ZipFile

data_path = 'dataset.zip'

with ZipFile(data_path, 'r') as zip:
    zip.extractall()
    print('The data set has been extracted.')
```

The data set has been extracted.

```
In [5]: path = 'lung_colon_image_set/lung_image_sets'
classes = os.listdir(path)
classes
```

```
Out[5]: ['lung_aca', 'lung_n', 'lung_scc']
```

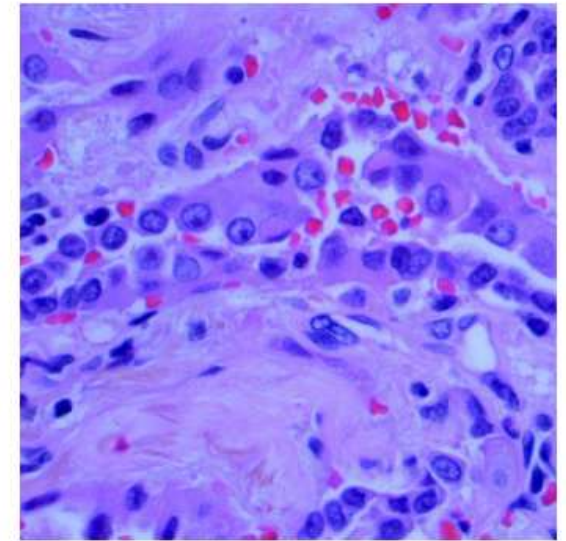
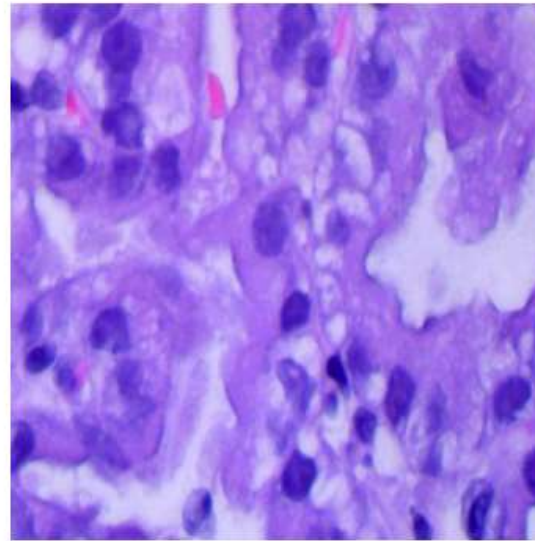
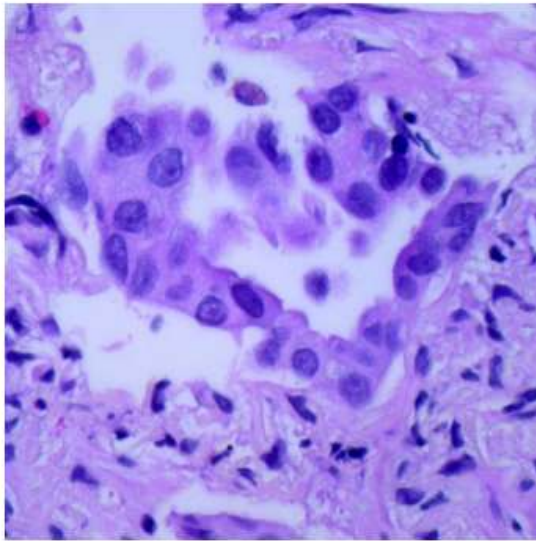
```
In [8]: path = 'lung_colon_image_set/lung_image_sets'

for cat in classes:
    image_dir = f'{path}/{cat}'
    images = os.listdir(image_dir)
```

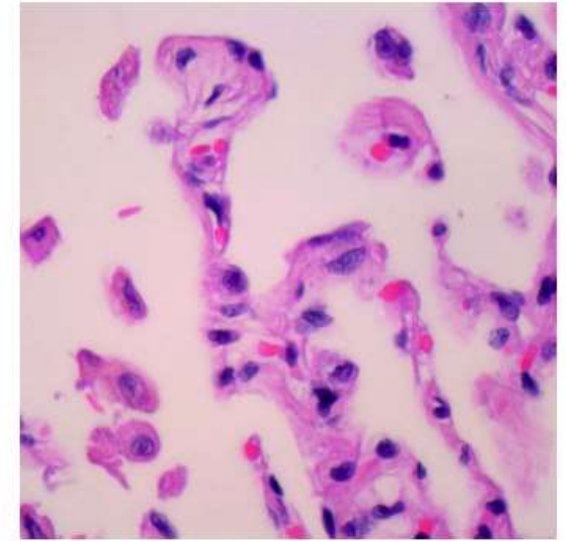
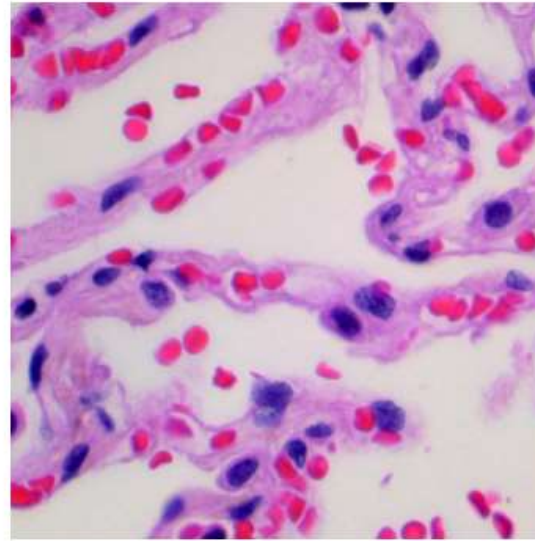
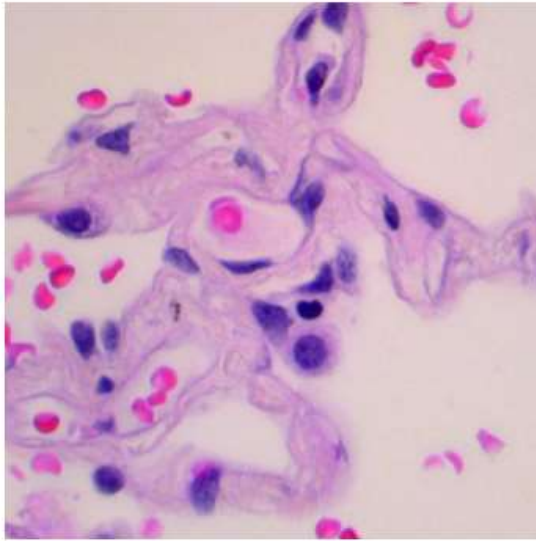
```
fig, ax = plt.subplots(1, 3, figsize=(15, 5))
fig.suptitle(f'Images for {cat} category . . . .', fontsize=20)

for i in range(3):
    k = np.random.randint(0, len(images))
    img = np.array(Image.open(f'{path}/{cat}/{images[k]}'))
    ax[i].imshow(img)
    ax[i].axis('off')
plt.show()
```

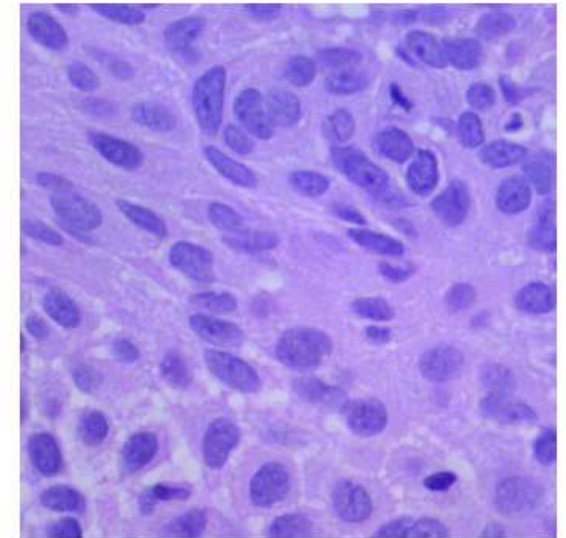
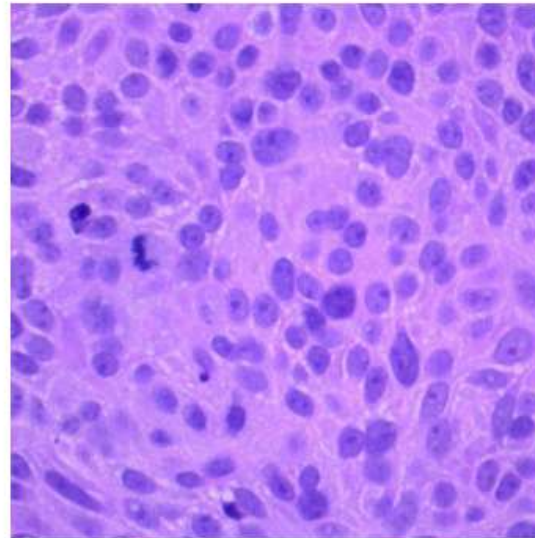
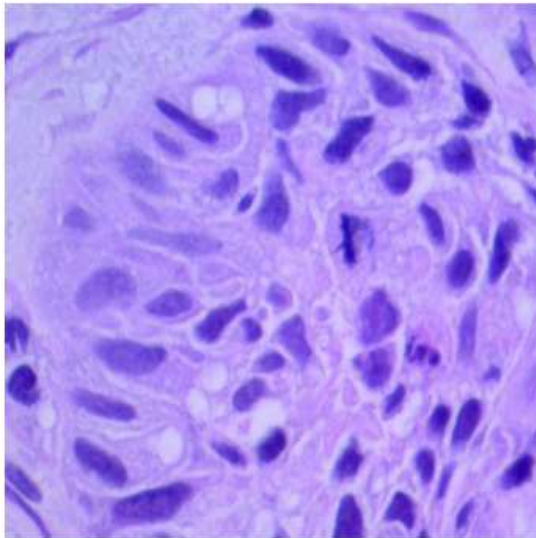
Images for lung\_aca category . . . .



## Images for lung\_n category . . . .



## Images for lung\_scc category . . . .



```
In [9]: IMG_SIZE = 256
        SPLIT = 0.2
        EPOCHS = 10
        BATCH_SIZE = 64
```

```
In [10]: X = []
        Y = []

        for i, cat in enumerate(classes):
            images = glob(f'{path}/{cat}/*.jpeg')

            for image in images:
                img = cv2.imread(image)

                X.append(cv2.resize(img, (IMG_SIZE, IMG_SIZE)))
                Y.append(i)

        X = np.asarray(X)
        one_hot_encoded_Y = pd.get_dummies(Y).values
```

```
In [11]: X_train, X_val, Y_train, Y_val = train_test_split(X, one_hot_encoded_Y,
                                                            test_size = SPLIT,
                                                            random_state = 2022)

        print(X_train.shape, X_val.shape)

        (12000, 256, 256, 3) (3000, 256, 256, 3)
```

```
In [12]: model = keras.models.Sequential([
            layers.Conv2D(filters=32,
                          kernel_size=(5, 5),
                          activation='relu',
                          input_shape=(IMG_SIZE,
                                        IMG_SIZE,
                                        3),
                          padding='same'),
            layers.MaxPooling2D(2, 2),

            layers.Conv2D(filters=64,
                          kernel_size=(3, 3),
                          activation='relu',
                          padding='same'),
            layers.MaxPooling2D(2, 2),
```

```
layers.Conv2D(filters=128,
              kernel_size=(3, 3),
              activation='relu',
              padding='same'),
layers.MaxPooling2D(2, 2),

layers.Flatten(),
layers.Dense(256, activation='relu'),
layers.BatchNormalization(),
layers.Dense(128, activation='relu'),
layers.Dropout(0.3),
layers.BatchNormalization(),
layers.Dense(3, activation='softmax')
])
```

In [13]: `model.summary()`

Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 256, 256, 32)	2432
max_pooling2d (MaxPooling2D)	(None, 128, 128, 32)	0
conv2d_1 (Conv2D)	(None, 128, 128, 64)	18496
max_pooling2d_1 (MaxPooling2D)	(None, 64, 64, 64)	0
conv2d_2 (Conv2D)	(None, 64, 64, 128)	73856
max_pooling2d_2 (MaxPooling2D)	(None, 32, 32, 128)	0
flatten (Flatten)	(None, 131072)	0
dense (Dense)	(None, 256)	33554688
batch_normalization (Batch Normalization)	(None, 256)	1024
dense_1 (Dense)	(None, 128)	32896
dropout (Dropout)	(None, 128)	0
batch_normalization_1 (Batch Normalization)	(None, 128)	512
dense_2 (Dense)	(None, 3)	387

```

=====
Total params: 33684291 (128.50 MB)
Trainable params: 33683523 (128.49 MB)
Non-trainable params: 768 (3.00 KB)
=====

```

```

In [14]: keras.utils.plot_model(
          model,
          show_shapes = True,

```



```
show_dtype = True,  
show_layer_activations = True  
)
```

You must install pydot (`pip install pydot`) and install graphviz (see instructions at <https://graphviz.gitlab.io/download/>) for plot\_model to work.

```
In [15]: model.compile(  
    optimizer = 'adam',  
    loss = 'categorical_crossentropy',  
    metrics = ['accuracy']  
)
```

```
In [16]: from keras.callbacks import EarlyStopping, ReduceLROnPlateau
```

```
class myCallback(tf.keras.callbacks.Callback):  
    def on_epoch_end(self, epoch, logs={}):  
        if logs.get('val_accuracy') > 0.90:  
            print('\n Validation accuracy has reached upto \  
                90% so, stopping further training.')  
            self.model.stop_training = True  
  
es = EarlyStopping(patience=3,  
                  monitor='val_accuracy',  
                  restore_best_weights=True)  
  
lr = ReduceLROnPlateau(monitor='val_loss',  
                      patience=2,  
                      factor=0.5,  
                      verbose=1)
```

```
In [17]: history = model.fit(X_train, Y_train,  
                             validation_data = (X_val, Y_val),  
                             batch_size = BATCH_SIZE,  
                             epochs = EPOCHS,  
                             verbose = 1,  
                             callbacks = [es, lr, myCallback()])
```

```

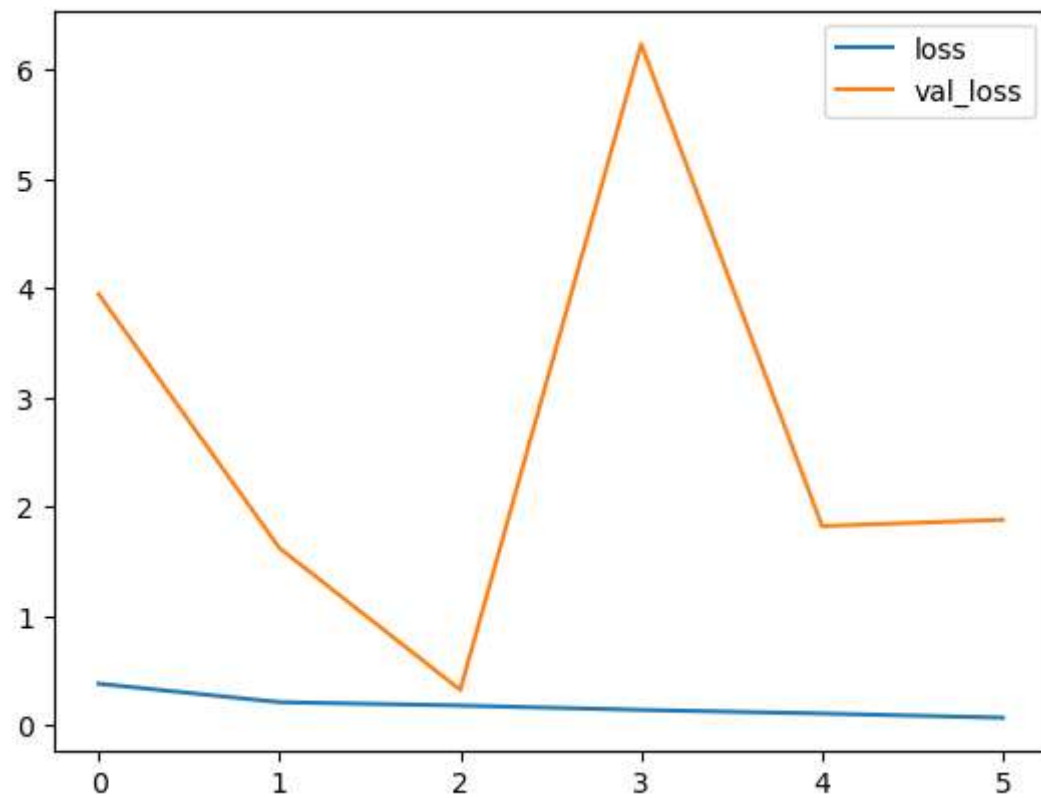
Epoch 1/10
188/188 [=====] - 636s 3s/step - loss: 0.3816 - accuracy: 0.8455 - val_loss: 3.9487 - val_accuracy: 0.6
620 - lr: 0.0010
Epoch 2/10
188/188 [=====] - 630s 3s/step - loss: 0.2126 - accuracy: 0.9128 - val_loss: 1.6230 - val_accuracy: 0.6
183 - lr: 0.0010
Epoch 3/10
188/188 [=====] - 625s 3s/step - loss: 0.1822 - accuracy: 0.9282 - val_loss: 0.3254 - val_accuracy: 0.8
803 - lr: 0.0010
Epoch 4/10
188/188 [=====] - 593s 3s/step - loss: 0.1416 - accuracy: 0.9436 - val_loss: 6.2358 - val_accuracy: 0.3
780 - lr: 0.0010
Epoch 5/10
188/188 [=====] - ETA: 0s - loss: 0.1091 - accuracy: 0.9603
Epoch 5: ReduceLROnPlateau reducing learning rate to 0.0005000000237487257.
188/188 [=====] - 595s 3s/step - loss: 0.1091 - accuracy: 0.9603 - val_loss: 1.8243 - val_accuracy: 0.5
987 - lr: 0.0010
Epoch 6/10
188/188 [=====] - 591s 3s/step - loss: 0.0706 - accuracy: 0.9748 - val_loss: 1.8819 - val_accuracy: 0.6
243 - lr: 5.0000e-04

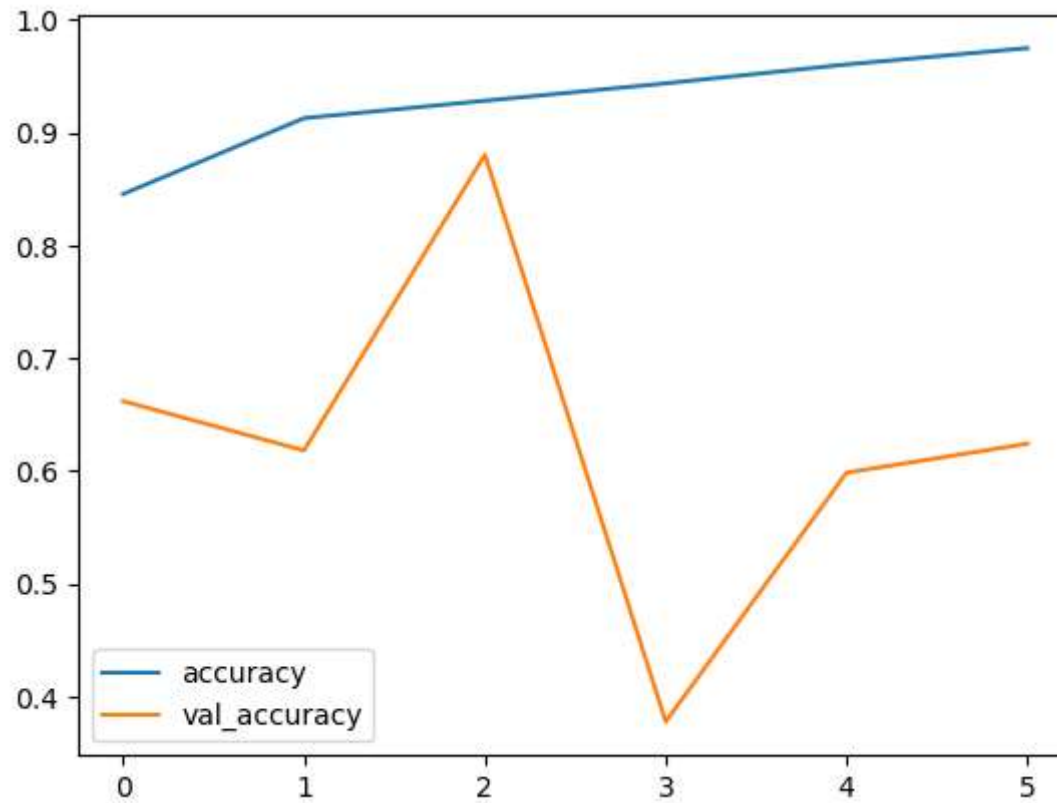
```

```
In [19]: model.save("model.h5")
```

```
In [18]: history_df = pd.DataFrame(history.history)
history_df.loc[:,['loss','val_loss']].plot()
history_df.loc[:,['accuracy','val_accuracy']].plot()
plt.show()
```







```
In [20]: Y_pred = model.predict(X_val)
Y_val = np.argmax(Y_val, axis=1)
Y_pred = np.argmax(Y_pred, axis=1)
```

94/94 [=====] - 24s 246ms/step

```
In [21]: metrics.confusion_matrix(Y_val, Y_pred)
```

```
Out[21]: array([[690,  53, 244],
               [  1, 976,   0],
               [ 61,   0, 975]], dtype=int64)
```

```
In [22]: print(metrics.classification_report(Y_val, Y_pred,
                                             target_names=classes))
```

	precision	recall	f1-score	support
lung_aca	0.92	0.70	0.79	987
lung_n	0.95	1.00	0.97	977
lung_scc	0.80	0.94	0.86	1036
accuracy			0.88	3000
macro avg	0.89	0.88	0.88	3000
weighted avg	0.89	0.88	0.88	3000

In [ ]:

In [ ]:

```
In [90]: # Now, you can load the model for testing
loaded_model = tf.keras.models.load_model('model.h5')

# Use the loaded model for predictions on new images
# new_image_path = 'Lungaca161.jpeg'
# new_image_path = 'Lungsc30.jpeg'
# new_image_path = 'Lungn29.jpeg'
new_image_path = 'lungsc44.jpeg'

new_image = tf.keras.preprocessing.image.load_img(new_image_path, target_size=(256, 256, 3))
new_image = tf.keras.preprocessing.image.img_to_array(new_image)
new_image = tf.expand_dims(new_image, axis=0)
result = loaded_model.predict(new_image)

print(result[0][1])

if result[0][1] >= 1:
    prediction = 'Cancerous'
else:
    prediction = 'Non-Cancerous'

print(f'The prediction for the new image is: {prediction}')
```

```
1/1 [=====] - 0s 171ms/step
1.0
The prediction for the new image is: Cancerous
```