

In [17]:

```
!pip install imutils
```

Requirement already satisfied: imutils in /opt/conda/lib/python3.7/site-packages (0.5.4)

WARNING: Running pip as the 'root' user can result in broken permissions and conflicting behaviour with the system package manager. It is recommended to use a virtual environment instead: <https://pip.pypa.io/warnings/venv>

In [18]:

```
# Importing Packages
```

```
from tensorflow.keras.preprocessing.image import ImageDataGenerator
from tensorflow.keras.layers import Dense, Flatten, MaxPooling2D, Input, Dropout, Activation, AveragePooling2D
from tensorflow.keras.applications import VGG16
from tensorflow.keras.utils import to_categorical
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.models import Model, Sequential
from keras.callbacks import ReduceLROnPlateau, LearningRateScheduler
from sklearn.metrics import classification_report
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelBinarizer
from sklearn.preprocessing import LabelEncoder
from tensorflow.keras.callbacks import EarlyStopping

import numpy as np # Linear algebra
import pandas as pd # data processing, CSV file I/O (e.g. pd.read_csv)
import matplotlib.pyplot as plt
import seaborn as sns
import os
import random
import shutil
import cv2
from imutils import paths
```

In [19]:

```
# root path
dataset_path = './dataset'
```

In [20]:

```
# Make directories for storing images
```

```
%%bash
rm -rf dataset
mkdir -p dataset/covid
mkdir -p dataset/normal
mkdir -p dataset/pneumonia
mkdir -p dataset/tuberculosis
```

In [21]:

```
# Set sample size
samples = 25
covid_dataset_path = '../input/covid-chest-xray'
```

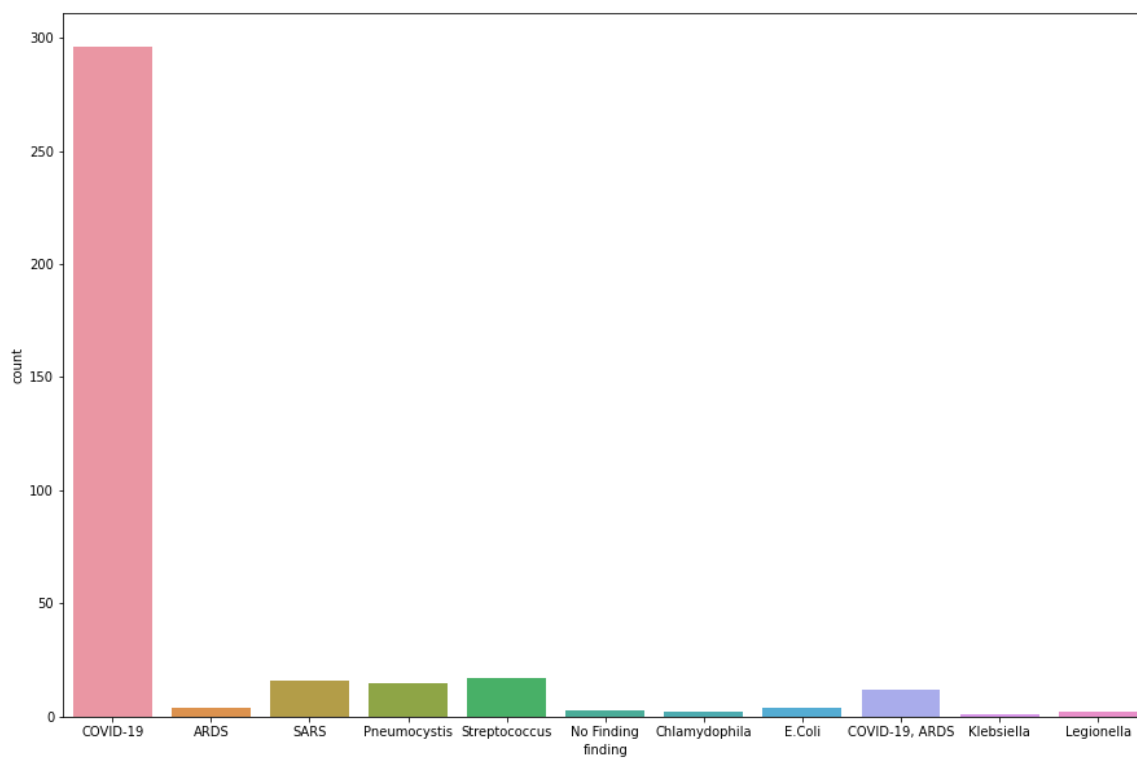
In [22]:

```
# Exploring both the datasets

# construct thvalue_counts the metadata CSV file and load it
csvPath = os.path.sep.join([covid_dataset_path, "metadata.csv"])
df = pd.read_csv(csvPath)

#EDA for COVID Xray Dataset
plt.figure(figsize=(15,10))
print(sns.countplot(x = df["finding"]))
```

AxesSubplot(0.125,0.125;0.775x0.755)



In [23]:

```
# Transferring the COVID-19 images to the COVID folder

# Loop over the rows of the COVID-19 data frame
for (i, row) in df.iterrows():
    # if (1) the current case is not COVID-19 or (2) this is not
    # a 'PA' view, then ignore the row
    if row["finding"] != "COVID-19" or row["view"] != "PA":
        continue

    # build the path to the input image file
    imagePath = os.path.sep.join([covid_dataset_path, "images", row["filename"]])

    # if the input image file does not exist (there are some errors in
    # the COVID-19 metadata file), ignore the row
    if not os.path.exists(imagePath):
        continue

    # extract the filename from the image path and then construct the
    # path to the copied image file
    filename = row["filename"].split(os.path.sep)[-1]
    outputPath = os.path.sep.join([f"{dataset_path}/covid", filename])

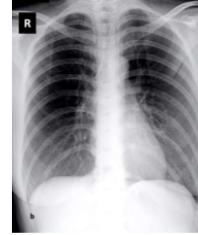
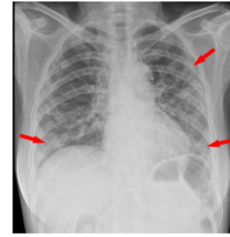
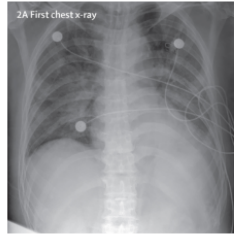
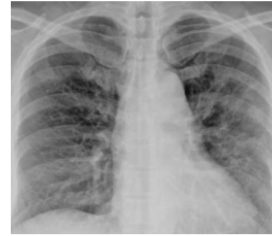
    # copy the image
    shutil.copy2(imagePath, outputPath)

# Display the images

covid = os.listdir("./dataset/covid")
covid_dir = "./dataset/covid"
plt.figure(figsize=(20, 10))

for i in range(9):
    plt.subplot(3, 3, i + 1)
    img = plt.imread(os.path.join(covid_dir, covid[i]))
    plt.imshow(img, cmap='gray')
    plt.axis('off')

plt.tight_layout()
```



In [24]:

```
# Transferring the normal images to the normal folder

normal_dataset_path = '../input/chest-xray-pneumonia/chest_xray'

basePath = os.path.sep.join([normal_dataset_path, "train", "NORMAL"])
ImagePaths = list(paths.list_images(basePath))

random.seed(42)
random.shuffle(ImagePaths)
ImagePaths = ImagePaths[:samples]

for (i, ImagePath) in enumerate(ImagePaths):
    filename = ImagePath.split(os.path.sep)[-1]
    outputPath = os.path.sep.join([f"{dataset_path}/normal", filename])

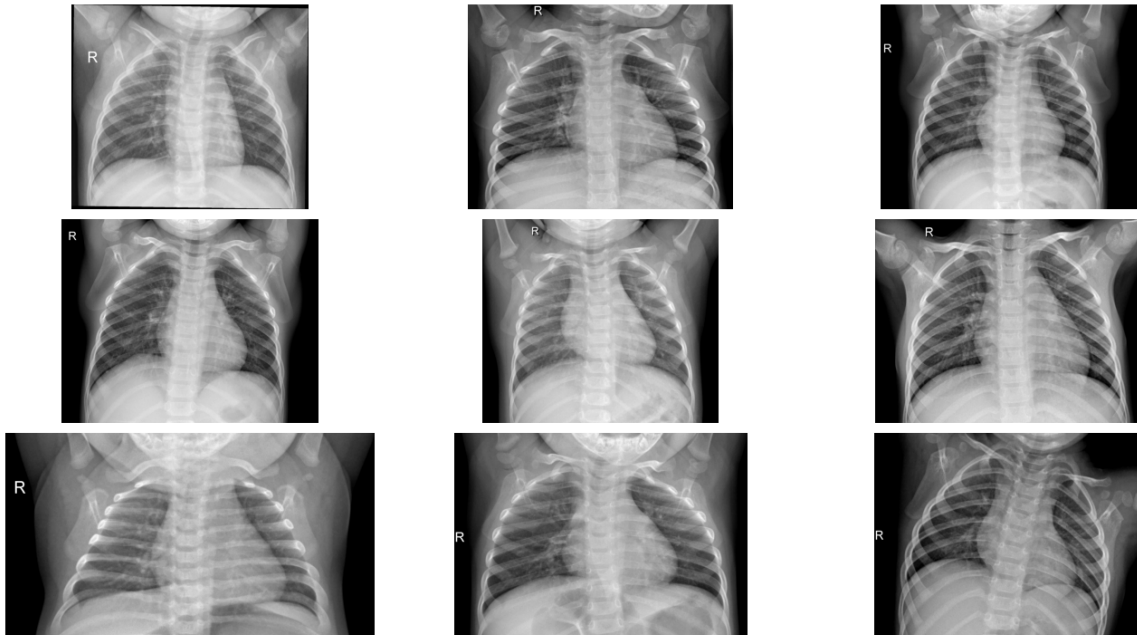
    shutil.copy2(ImagePath, outputPath)

# Display the images

normal = os.listdir("../dataset/normal")
normal_dir = "../dataset/normal"
plt.figure(figsize=(20, 10))

for i in range(9):
    plt.subplot(3, 3, i + 1)
    img = plt.imread(os.path.join(normal_dir, normal[i]))
    plt.imshow(img, cmap='gray')
    plt.axis('off')

plt.tight_layout()
```



In [25]:

```
# Transferring the pneumonia images to the pneumonia folder

pneumonia_dataset_path = '../input/chest-xray-pneumonia/chest_xray'

PneuPath = os.path.sep.join([pneumonia_dataset_path, "train", "PNEUMONIA"])
ImagePaths = list(paths.list_images(PneuPath))

random.seed(42)
random.shuffle(ImagePaths)
ImagePaths = ImagePaths[:samples]

for (i, ImagePath) in enumerate(ImagePaths):
    filename = ImagePath.split(os.path.sep)[-1]
    outputPath = os.path.sep.join([f"{dataset_path}/pneumonia", filename])

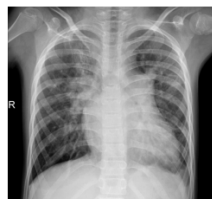
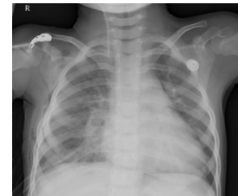
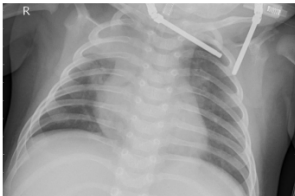
    shutil.copy2(ImagePath, outputPath)

# Display the images

pneumonia = os.listdir("../dataset/pneumonia")
pneumonia_dir = "../dataset/pneumonia"
plt.figure(figsize=(20, 10))

for i in range(9):
    plt.subplot(3, 3, i + 1)
    img = plt.imread(os.path.join(pneumonia_dir, pneumonia[i]))
    plt.imshow(img, cmap='gray')
    plt.axis('off')

plt.tight_layout()
```



In [26]:

```
# Transferring the tuberculosis images to the tuberculosis folder

tuberculosis_dataset_path = '../input/chest-xray-pneumoniacovid19tuberculosis'

TubePath = os.path.sep.join([tuberculosis_dataset_path, "train", "TURBERCULOSIS"])
ImagePaths = list(paths.list_images(TubePath))

random.seed(42)
random.shuffle(ImagePaths)
ImagePaths = ImagePaths[:samples]

for (i, ImagePath) in enumerate(ImagePaths):
    filename = ImagePath.split(os.path.sep)[-1]
    outputPath = os.path.sep.join([f"{dataset_path}/tuberculosis", filename])

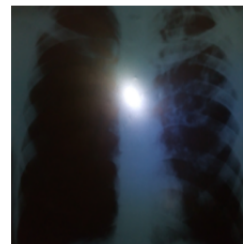
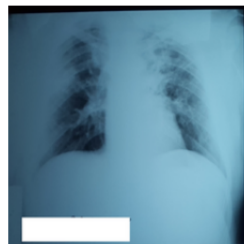
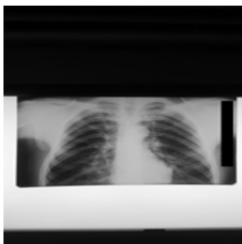
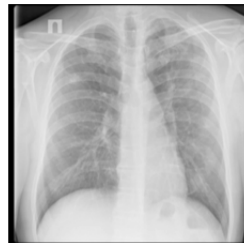
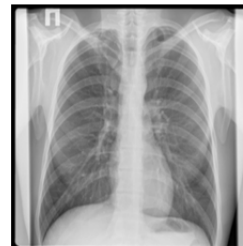
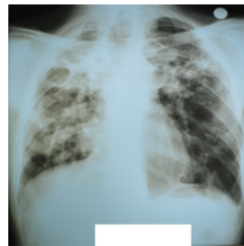
    shutil.copy2(ImagePath, outputPath)

# Display the images

tuberculosis = os.listdir("../dataset/tuberculosis")
tuberculosis_dir = "../dataset/tuberculosis"
plt.figure(figsize=(20, 10))

for i in range(9):
    plt.subplot(3, 3, i + 1)
    img = plt.imread(os.path.join(tuberculosis_dir, tuberculosis[i]))
    plt.imshow(img, cmap='gray')
    plt.axis('off')

plt.tight_layout()
```



In [27]:

```
# Data Preprocessing

# initialize the initial learning rate, number of epochs to train for,
# and batch size
LR = 1e-3
EPOCHS = 30
BS = 8
```

In [28]:

```
# Data Preprocessing - Converting the images to grey

imagePaths = list(paths.list_images(dataset_path))

data = []
labels = []

for image in imagePaths:
    label = image.split(os.path.sep)[-2]
    img = cv2.imread(image)
    img = cv2.cvtColor(img, cv2.COLOR_BGR2RGB)
    img = cv2.resize(img, (224, 224))

    data.append(img)
    labels.append(label)

data = np.array(data) / 255.0
labels = np.array(labels)
```

In [29]:

```
# One hot encoding for labels

lb = LabelEncoder()

labels = lb.fit_transform(labels)
labels = to_categorical(labels)

X_train, X_test, y_train, y_test = train_test_split(data, labels, test_size=0.20, random_state=42, stratify=labels)

trainAug = ImageDataGenerator(rotation_range=15, fill_mode="nearest")
```


In [30]:

```
# Creating the CNN

baseModel = VGG16(weights="imagenet", include_top=False, input_tensor=Input(shape=(224,
224, 3)))

mainModel = baseModel.output
mainModel = AveragePooling2D(pool_size = (4,4))(mainModel)
mainModel = Flatten(name = "flatten")(mainModel)
mainModel = Dense(64, activation = "relu")(mainModel)
mainModel = Dropout(0.5)(mainModel)
mainModel = Dense(4, activation = "softmax")(mainModel)

model = Model(inputs = baseModel.input, outputs = mainModel)

for layer in baseModel.layers:
    layer.trainable = False

opt = Adam(learning_rate=LR)

model.compile(loss= "categorical_crossentropy", optimizer = "adam", metrics = ["accuracy"])
```

In [31]:

```
# Setup the callbacks

early_stop = EarlyStopping(monitor='val_loss', patience=2, restore_best_weights=True)
```

In [32]:

```
# Fit the data to the model
```

```
H = model.fit_generator(  
    trainAug.flow(X_train, y_train, batch_size=BS),  
    steps_per_epoch= len(X_train) // BS,  
    validation_data= (X_test, y_test),  
    validation_steps = len(X_test) // BS,  
    epochs=EPOCHS, callbacks = early_stop)
```

```
/opt/conda/lib/python3.7/site-packages/keras/engine/training.py:1972: User
Warning: `Model.fit_generator` is deprecated and will be removed in a futu
re version. Please use `Model.fit`, which supports generators.
  warnings.warn("`Model.fit_generator` is deprecated and "
```

Epoch 1/30
21/21 [=====] - 53s 2s/step - loss: 1.2666 - accuracy: 0.5366 - val_loss: 0.9943 - val_accuracy: 0.6591
Epoch 2/30
21/21 [=====] - 51s 2s/step - loss: 1.0540 - accuracy: 0.6220 - val_loss: 0.9679 - val_accuracy: 0.6591
Epoch 3/30
21/21 [=====] - 52s 2s/step - loss: 1.0041 - accuracy: 0.6585 - val_loss: 0.9399 - val_accuracy: 0.6591
Epoch 4/30
21/21 [=====] - 51s 2s/step - loss: 0.9685 - accuracy: 0.6646 - val_loss: 0.9066 - val_accuracy: 0.6591
Epoch 5/30
21/21 [=====] - 51s 2s/step - loss: 0.9636 - accuracy: 0.6548 - val_loss: 0.8910 - val_accuracy: 0.6591
Epoch 6/30
21/21 [=====] - 50s 2s/step - loss: 0.8901 - accuracy: 0.6463 - val_loss: 0.8412 - val_accuracy: 0.6591
Epoch 7/30
21/21 [=====] - 51s 2s/step - loss: 0.8589 - accuracy: 0.6829 - val_loss: 0.8103 - val_accuracy: 0.6818
Epoch 8/30
21/21 [=====] - 53s 3s/step - loss: 0.8389 - accuracy: 0.6890 - val_loss: 0.7669 - val_accuracy: 0.6818
Epoch 9/30
21/21 [=====] - 55s 3s/step - loss: 0.7966 - accuracy: 0.6890 - val_loss: 0.7577 - val_accuracy: 0.7045
Epoch 10/30
21/21 [=====] - 56s 3s/step - loss: 0.7913 - accuracy: 0.7073 - val_loss: 0.7133 - val_accuracy: 0.7045
Epoch 11/30
21/21 [=====] - 52s 3s/step - loss: 0.7273 - accuracy: 0.7012 - val_loss: 0.6938 - val_accuracy: 0.7045
Epoch 12/30
21/21 [=====] - 52s 2s/step - loss: 0.7302 - accuracy: 0.7500 - val_loss: 0.6706 - val_accuracy: 0.7273
Epoch 13/30
21/21 [=====] - 52s 3s/step - loss: 0.6758 - accuracy: 0.7439 - val_loss: 0.6457 - val_accuracy: 0.7273
Epoch 14/30
21/21 [=====] - 51s 2s/step - loss: 0.6515 - accuracy: 0.7500 - val_loss: 0.6497 - val_accuracy: 0.7955
Epoch 15/30
21/21 [=====] - 51s 2s/step - loss: 0.6353 - accuracy: 0.7988 - val_loss: 0.6065 - val_accuracy: 0.7500
Epoch 16/30
21/21 [=====] - 51s 2s/step - loss: 0.6124 - accuracy: 0.7683 - val_loss: 0.5847 - val_accuracy: 0.7727
Epoch 17/30
21/21 [=====] - 53s 3s/step - loss: 0.6116 - accuracy: 0.7561 - val_loss: 0.5684 - val_accuracy: 0.7955
Epoch 18/30
21/21 [=====] - 52s 2s/step - loss: 0.6196 - accuracy: 0.7927 - val_loss: 0.5690 - val_accuracy: 0.7955
Epoch 19/30
21/21 [=====] - 51s 2s/step - loss: 0.5830 - accuracy: 0.7866 - val_loss: 0.5456 - val_accuracy: 0.7955
Epoch 20/30
21/21 [=====] - 51s 2s/step - loss: 0.5233 - accuracy: 0.8171 - val_loss: 0.5309 - val_accuracy: 0.7955
Epoch 21/30

```

21/21 [=====] - 51s 2s/step - loss: 0.5687 - accu
racy: 0.8171 - val_loss: 0.5110 - val_accuracy: 0.8182
Epoch 22/30
21/21 [=====] - 51s 2s/step - loss: 0.5379 - accu
racy: 0.8049 - val_loss: 0.4983 - val_accuracy: 0.7955
Epoch 23/30
21/21 [=====] - 52s 2s/step - loss: 0.4742 - accu
racy: 0.8415 - val_loss: 0.4851 - val_accuracy: 0.7955
Epoch 24/30
21/21 [=====] - 51s 2s/step - loss: 0.5590 - accu
racy: 0.7988 - val_loss: 0.4642 - val_accuracy: 0.8182
Epoch 25/30
21/21 [=====] - 51s 2s/step - loss: 0.4932 - accu
racy: 0.8171 - val_loss: 0.4479 - val_accuracy: 0.8182
Epoch 26/30
21/21 [=====] - 55s 3s/step - loss: 0.5161 - accu
racy: 0.8415 - val_loss: 0.4452 - val_accuracy: 0.8182
Epoch 27/30
21/21 [=====] - 52s 3s/step - loss: 0.5306 - accu
racy: 0.7622 - val_loss: 0.4418 - val_accuracy: 0.8864
Epoch 28/30
21/21 [=====] - 50s 2s/step - loss: 0.4581 - accu
racy: 0.8537 - val_loss: 0.4120 - val_accuracy: 0.8409
Epoch 29/30
21/21 [=====] - 51s 2s/step - loss: 0.4230 - accu
racy: 0.8720 - val_loss: 0.3958 - val_accuracy: 0.8182
Epoch 30/30
21/21 [=====] - 50s 2s/step - loss: 0.4827 - accu
racy: 0.8049 - val_loss: 0.3899 - val_accuracy: 0.8409

```

In [33]:

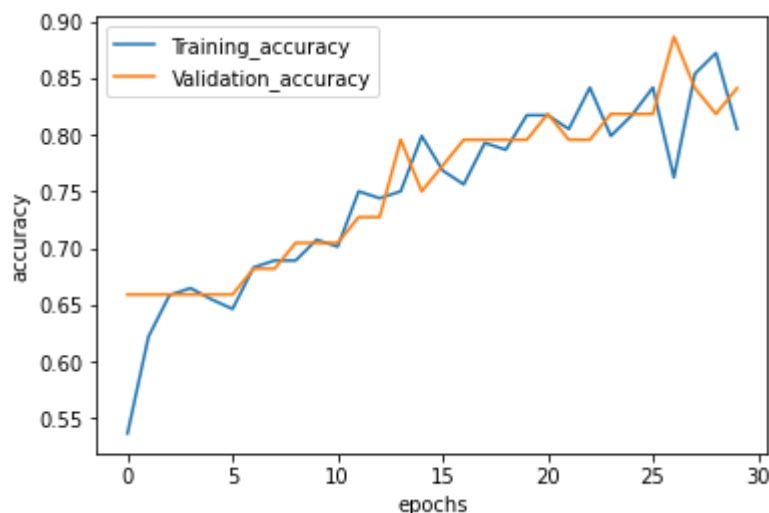
```
# Plot showing the accuracy across epochs
```

```

accs = H.history['accuracy']
val_accs = H.history['val_accuracy']

plt.plot(range(len(accs)),accs, label = 'Training_accuracy')
plt.plot(range(len(accs)),val_accs, label = 'Validation_accuracy')
plt.ylabel('accuracy')
plt.xlabel('epochs')
plt.legend()
plt.show()

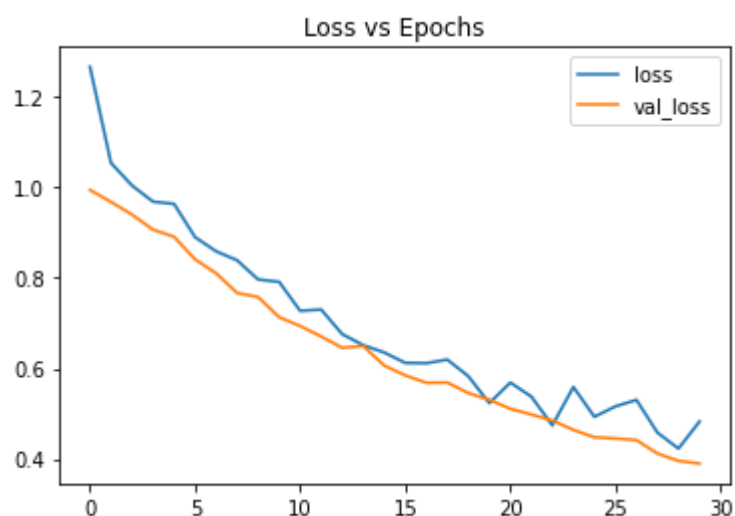
```



In [34]:

```
# Plot showing the loss across epochs
```

```
pd.DataFrame(model.history.history)[['loss', 'val_loss']].plot()  
plt.title("Loss vs Epochs")  
plt.show()
```



In [35]:

```
# Predictions
```

```
preds = model.predict(X_test, batch_size=BS)  
preds = np.argmax(preds, axis=1)  
print(classification_report(y_test.argmax(axis=1), preds, target_names=lb.classes_))
```

	precision	recall	f1-score	support
covid	0.83	1.00	0.91	29
normal	0.80	0.80	0.80	5
pneumonia	1.00	0.40	0.57	5
tuberculosis	1.00	0.40	0.57	5
accuracy			0.84	44
macro avg	0.91	0.65	0.71	44
weighted avg	0.86	0.84	0.82	44