Gurjus Singh

MSDS 458 – Artificial Intelligence and Deep Learning

March 14th, 2021

Week 10: A.4 Fourth Research/Programming Assignment

Abstract

In this research, I came up with a way to classify X-Ray Chest Images with COVID-19. The goal of this research was to use supervised learning binary classification with Normal and COVID-19 X-Ray Images. I used a CNN architecture as this was important in image classification. I used accuracy for deciding which model was performing the best, but also looked at other metrics in my best model to see how it was performing. I also performed T-SNE on my best model as well as saw how the Convolution/Max Pooling by making a channel grid.

Introduction

In purpose of this research was to use supervised learning binary classification to classify X-Ray Chest Images of suspected COVID-19 patients. The dataset I used was from Kaggle and contained 2000 train images and 328 test images divided evenly between COVID Positive and COVID Negative images. The Neural Network architecture I used was a Convolution Neural Network architecture as this was suitable for image data. I tried out different CNN architectures before choosing my best model. I also decided to find out what features the layers of the CNN were learning by creating a grid of channels. I also decided to perform T-SNE on my best model. After experimenting with my best model, I then made a management recommendation.

Literature Review

As COVID-19 spreads, and vaccines are distributed, there continues to be growing research on how to diagnose patients. One such paper that I found, talked about how countries

have adopted reverse transcription polymerase chain reaction to diagnose COVID-19 [1]. The authors of the paper point out this test takes 4-6 hours or even a whole day to get results [1]. The test can also give false positives and false negatives [1]. The authors then point out that one solution to preventing false positives/negatives is to test COVID-19 infections by using XRAY/CT Scans of the Chest in COVID patients [1].

The paper goes on to share the results of XRAY/CT Scan classification [1]. They use multi-image classification to decrease overfitting [1]. This is a way to generate more images for the CNN architecture to train on [1]. In the results of the experiments with a 70 percent-30 percent training-testing split the authors obtained a 95.38 percent accuracy for CT Scans, while for X-ray they obtained a 98.97 percent accuracy [1].

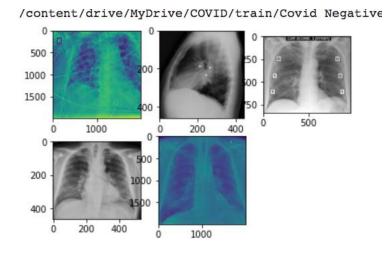
Methods

I first start out this research by importing the necessary packages to complete this project. I found the keras and tensoflow packages important for implementing the CNN architecture. I also found sklearn package important for computing metrics. I also found matplotlib package important for creating the EDA, and the os package was important for extracting the files from the directories. The imported packages are shown below in 1-1.

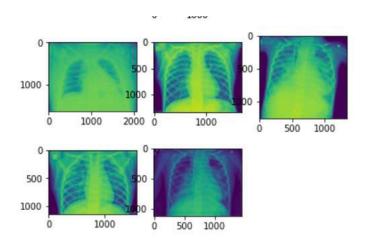
```
import matgletlib pyplot as plt
import matgletlib pyplot as plt
import pandas as pd
import pandas import dequential
from koras.models import deal import pangabutaGenerator
from koras.models import pandas import pangabutaGenerator
from koras.models import pandas import pangabutaGenerator
from kalean.metrica import pangabutaGenerator
from kalean.metrica import pangabutaGenerator
from kalean.metrica import pensions.geore
import tori
im
```

Import Packages 1-1

After importing the packages my next step was to do some EDA. I plotted the first 5 images of COVID-POSITIVE patients as seen in 1-2 and plotted first 5 photos of COVID NEGATIVE patients as seen in 1-3.

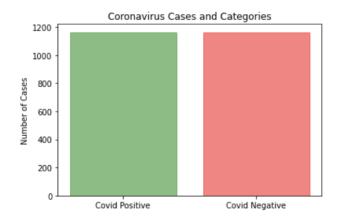


COVID-POSITIVE PATIENT XRAYS 1-2



COVID-NEGATIVE PATIENT XRAYS 1-2

After taking an initial look at the data, the next thing I did is loaded the data in the RGB format and also created labels which we will use for training. I also resized each image to 64 x64. After loading of the images and creation of the labels, I looked how many COVID-POSTIVE and NEGATIVE X-RAY Images were in the dataset as seen in 1-3. I noticed there were even number of both sets of images.



Number of COVID CASES 1-3

I also wanted to see what the Train and Test split percentage was as it was already split when taking the dataset from Kaggle. What I noticed was that it was a 85.9 percent to 14.1 percent Train to Test split as seen in 1-4.



Training-Test Split 1-4

After looking at the split I then extracted the labels I created and encoded and stored them in an array, I did the same with the images. I then normalized the images by dividing each image by 255 to get it between 0 and 1. After normalizing the images, I then added more images to the training data to prevent overfitting by using an ImageDataGenerator. This is known as Data Augmentation, and it creates more images by manipulating the existing images to create more images for training. After doing data augmentation, I then split the training data 80 percent to 20 percent to create a validation set.

I then went ahead to create my CNN Architectures which involved Max Pooling,
Convolution and Average Pooling layers. Max pooling helps in reducing the number of
parameters in the network by "downsizing feature maps" by only extracting the important
features of an image[2]. The convolutional layer functions in that tries to extract local patterns
from a window [2]. The window is called a filter which goes over each part of the image
extracting the features in the image [2]. Average pooling retains less important information of an
image by averaging the features [4].

After building each CNN Architecture, I then observed the accuracy and obtained the best model. On the best model, I performed TSNE, also built a grid for each layer in the CNN architecture, and computed other metrics such as F1 Score, Precision Score etc.

Results

In the research I performed a total of 17 experiments, 20 Epochs each, and also performed regularization techniques such as L1, L2 Regularizers, Dropout and Early Stopping. My Performance Summary is below in 1-5. What I noticed from these experiments was that Experiment 9 with 3 Max Pooling, 3 Convolution Layers with 128 Filters; and 1 Dense Layer with 256 Neurons was performing the best. I noticed it's testing accuracy was 98.47 while Training accuracy was 98.50 which means it was not overfitting. This was the highest testing accuracy that I had among the 17 models which was why I chose this model.

EXPERIMENTS - 20 EPOCHS EACH;	Regularization	TRAIN ACCURACY	VAL ACCURACY	TEST ACCURACY	TRAIN LOSS	VAL LOSS	TEST LOSS	TIME (HR, MIN, SECS
- 1 MAX POOLING AND CONV LAYER - 32								
ILTERS; 1 DENSE LAYER 32 NEURONS	None	97.62%	96.25%	96.03%	0.0725	0.0823	0.1094	4.23 SECS
- 1 MAX POOLING AND CONV LAYER - 64 ILTERS; 1 DENSE LAYER 64 NEURONS	None	98.50%	97.50%	96.95%	0.0532	0.0634	0.0866	5.04 SECS
- 1 MAX POOLING AND CONV LAYER - 128 ILTERS; 1 DENSE LAYER 128 NEURONS	None	98%	96.75%	96.03%	6.52E-02	0.0752	0.0958	7.1 SECS
 2 MAX POOLING AND CONV LAYER - 32 ILTERS; 1 DENSE LAYER 256 NEURONS 	None	98%	97.75%	96.95%	6.39E-02	0.0564	0.0762	4.54 SECS
- 2 MAX POOUNG AND CONV LAYER - 64 ILTERS; 1 DENSE LAYER 256 NEURONS	None	98%	97.75%	97.25%	0.0584	0.658	0.0834	6.1 SECS
- 2 MAX POOUNG AND CONV LAYER - 128 ILTERS; 1 DENSE LAYER 256 NEURONS	None	98%	96.25%	97.86%	0.0603	0.0714	0.0614	9.6 SECS
- 3 MAX POOLING AND CONV LAYER - 64 ILTERS; 1 DENSE LAYER 256 NEURONS	None	98.45%	99.00%	99.08%	0.0442	0.0342	0.0384	6 SECS
- 3 AVERAGE POOLING AND CONV LAYER - 64 ILTERS; 1 DENSE LAYER 256 NEURONS	None	97.74%	97.75%	94.82%	0.0674	0.0655	0.1216	6.05 SECS
- 3 MAX POOLING AND CONV LAYER - 128 ILTERS; 1 DENSE LAYER 256 NEURONS	None	98.50%	98.25%	98.47%	0.0514	0.053	0.0485	9.32 SECS
0 - 3 MAX POOLING AND CONV LAYER - 256 ILTERS; 1 DENSE LAYER 256 NEURONS	None	97.61%	98.75%	97.56%	0.0643	0.0338	0.0577	21.2 SECS
1 - 3 MAX POOUNG AND CONV LAYER - 128 ILTERS; 1 DENSE LAYER 256 NEURONS	1 DROPOUT LAYER 20%	88.50%	97.25%	97.56%	1.5564	0.0753	0.083	9.29 SECS
2 - 3 MAX POOLING AND CONV LAYER - 128 ILTERS; 1 DENSE LAYER 256 NEURONS	2 DROPOUT LAYER 20%	98.33%	99.00%	96.95%	0.0516	0.0365	0.0777	9.7 SECS
3 - 3 MAX POOLING AND CONV LAYER - 128 ILTERS; 1 DENSE LAYER 256 NEURONS	3 DROPOUT LAYER 20%	98.66%	97.75%	96.95%	0.0574	0.0518	0.0872	9.4 SECS
4- 3 MAX POOLING AND CONV LAYER - 256 ILTERS; 1 DENSE LAYER 256 NEURONS	EARLY STOPPIN PATIENCE 1	55.99%	49.75%	50.00%	0.6849	0.7115	0.7338	2.4 SECS
5 - 3 MAX POOLING AND CONV LAYER - 256 ILTERS; 1 DENSE LAYER 256 NEURONS	EARLY STOPPIN PATIENCE 2	94.39%	94.50%	94.21%	0.161	0.1336	0.1306	8.26 SECS
6 - 3 MAX POOLING AND CONV LAYER - 256 ILTERS; 1 DENSE LAYER 256 NEURONS	L1 REGULARIZER	98.09%	97.25%	97.86%	0.155	0.1615	0.1612	33.8 SECS
7 - 3 MAX POOLING AND CONV LAYER - 256 ILTERS; 1 DENSE LAYER 256 NEURON	L2 REGULARIZER	98.51%	98.50%	98.17%	0.059	0.0483	0.0659	18.8 SECS

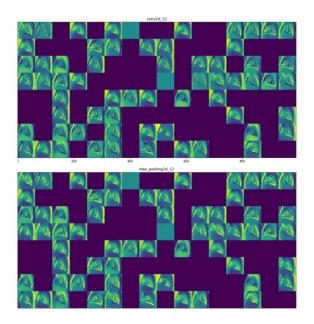
Performance Summary; See Excel Sheet attached for better look 1-5

Next, I looked into the metrics to see how good of a job the model 9 was doing. From the results in 1-6, 1-7. From the Confusion Matrices, I felt like the model was doing a fantastic job. I also noticed that Precision and Recall scores were around 0.98, and the F1 Scores were high as well being close to 1 which is the best score.

Training Set Metrics 1-6

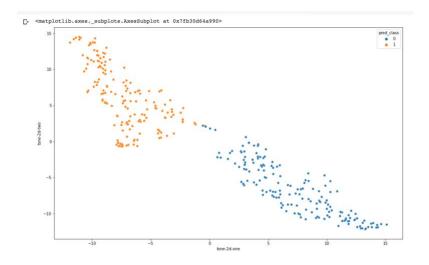
Testing Set Metrics 1-7

After looking at key metrics for my model, I then looked at the grid for how the layers in the architecture learned in 1-8. I noticed the layers was trying to learn the features of the bones in the X-ray.



Grid for Layers 1-8

Lastly, I performed T-SNE, which is dimensionality reduction unsupervised learning technique to visualize higher dimension data as seen in 1-9 [3].



T-SNE on COVID Dataset 1-9

Conclusion

In this experiment, I did binary classification on COVID-19 Xray images using CNN Architecture. From doing 17 experiments I chose Experiment 9 which had an architecture of 3 Max Pooling, 3 Convolution Layers with 128 Filters; and 1 Dense Layer with 256 Neurons as the best experiment. After choosing the best model, I also computed key metrics in 1-6, 1-7, saw how the layers learned by creating a grid in 1-8, and performed T-SNE in 1-9. For my recommendation to management, I choose an architecture of 3 Max Pooling, 3 Convolution Layers with 128 Filters; and 1 Dense Layer with 256 Neurons to diagnose COVID Patients. In the future, I hope to see if the CNN Architecture can distinguish between other fluid like illnesses like Pneumonia.

References

- [1] Purohit, K., Kesarwani, A., Kisku, D. R., & Dalui, M. (2020, January 1). *COVID-19*detection on chest x-ray and ct scan images using multi-image augmented deep learning

 model. BioRxiv. https://www.biorxiv.org/content/10.1101/2020.07.15.205567v2.full
- [2] Chollet, F. (2017). Deep Learning with Python. Manning Publications Company.
- [3] Introduction to t-SNE. (n.d.). Data Camp. Retrieved February 6, 2021, from https://www.datacamp.com/community/tutorials/introduction-t-sne

pooling-global-max-pooling-and-global-average-pooling/

[4] C. (2020, January 30). What are Max Pooling, Average Pooling, Global Max Pooling and Global Average Pooling? –. MachineCurve.

https://www.machinecurve.com/index.php/2020/01/30/what-are-max-pooling-average-

- APPENDIX

```
import matplotlib.pyplot as plt
import seaborn as sns
import pandas as pd
import keras
from keras.models import Sequential
from keras.layers import Dense, Conv2D , MaxPool2D , Flatten , Dro
from keras.preprocessing.image import ImageDataGenerator
from keras.optimizers import Adam
from sklearn.metrics import classification report, confusion matrix
from sklearn.metrics import f1 score
from sklearn.metrics import precision score
from sklearn.metrics import recall_score
import tensorflow as tf
from sklearn.decomposition import PCA
import cv2
import os
import numpy as np
from sklearn.model selection import train test split
# TensorFlow and tf.keras
import tensorflow as tf
from tensorflow import keras
from tensorflow.keras.utils import to categorical
from tensorflow.keras import models, layers
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Conv2D, MaxPooling2D
from tensorflow.keras.layers import Dropout, Flatten, Input, Dense
# Load the Drive helper and mount
from google.colab import drive
```

drive.mount('/content/drive')

Mounted at /content/drive

```
import matplotlib.image as mpimg
directory=os.listdir('/content/drive/MyDrive/COVID/train/')
for each in directory:
     plt.figure()
     currentFolder = '/content/drive/MyDrive/COVID/train/' + each
     for i, file in enumerate(os.listdir(currentFolder)[0:5]):
                                            + "/" + file
          fullpath = currentFolder
          print(fullpath)
          img=mpimg.imread(fullpath)
          plt.subplot(2, 3, i+1)
          plt.imshow(img)
    /content/drive/MyDrive/COVID/train/Covid Positive/00870a9c.jpg
    /content/drive/MyDrive/COVID/train/Covid Positive/000025-1.jpg
    /content/drive/MyDrive/COVID/train/Covid Positive/11547 2020 1200 Fig2 HTML-a.png
    /content/drive/MyDrive/COVID/train/Covid Positive/000024-1.jpg
    /content/drive/MyDrive/COVID/train/Covid Positive/1052b0fe.jpg
    /content/drive/MyDrive/COVID/train/Covid Negative/person108 virus 199.jpeg
    /content/drive/MyDrive/COVID/train/Covid Negative/person120 virus 226.jpeg
    /content/drive/MyDrive/COVID/train/Covid Negative/person124 virus 238.jpeg
    /content/drive/MyDrive/COVID/train/Covid Negative/person130 virus 263.jpeg
    /content/drive/MyDrive/COVID/train/Covid Negative/person124_virus_236.jpeg
     500
     1000
     1500
                                       500
                              400
            1000
                         200
                    0
     200
     400
           200
               400
                         1000
                   500
     1000
                  1000
            1000
                  2000 0
                          1000
                                     500
                                         1000
     500
     1000
              1000
                           1000
                     Ó
                        500
```

import matplotlib.image as mpimg
directory=os.listdir('/content/drive/MyDrive/COVID/train/')

```
for each in directory:
     plt.figure()
     currentFolder = '/content/drive/MyDrive/COVID/train/' + each
     for i, file in enumerate(os.listdir(currentFolder)[0:5]):
                                            + "/" + file
          fullpath = currentFolder
          print(fullpath)
          img=mpimg.imread(fullpath)
          plt.subplot(2, 3, i+1)
          plt.imshow(img)
    /content/drive/MyDrive/COVID/train/Covid Positive/00870a9c.jpg
    /content/drive/MyDrive/COVID/train/Covid Positive/000025-1.jpg
    /content/drive/MyDrive/COVID/train/Covid Positive/11547 2020 1200 Fig2 HTML-a.png
    /content/drive/MyDrive/COVID/train/Covid Positive/000024-1.jpg
    /content/drive/MyDrive/COVID/train/Covid Positive/1052b0fe.jpg
    /content/drive/MyDrive/COVID/train/Covid Negative/person108 virus 199.jpeg
    /content/drive/MyDrive/COVID/train/Covid Negative/person120 virus 226.jpeg
    /content/drive/MyDrive/COVID/train/Covid Negative/person124 virus 238.jpeg
    /content/drive/MyDrive/COVID/train/Covid Negative/person130 virus 263.jpeg
    /content/drive/MyDrive/COVID/train/Covid Negative/person124 virus 236.jpeg
     500
     1000
     1500
            1000
                         200
                    0
       0
     200
                  1000
     400
                400
            200
                         1000
                                 0
                   5bo
     1000
                  1000
            1000
                  2000
                           1000
       0
     500
     1000
              1000
                     Ó
                        500
                           1000
labels = ['Covid Negative', 'Covid Positive']
img size = 64
def get data(data dir):
```

data = []

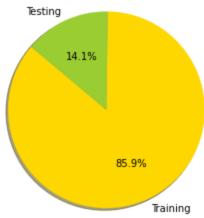
for label in labels:

```
path = os.path.join(data dir, label)
        class num = labels.index(label)
        for img in os.listdir(path):
            try:
                img arr = cv2.imread(os.path.join(path, img))[...,
                resized arr = cv2.resize(img arr, (img size, img s
                data.append([resized arr, class num])
            except Exception as e:
                print(e)
    return np.array(data)
train = get_data('/content/drive/MyDrive/COVID/train/')
test = get_data('/content/drive/MyDrive/COVID/test/')
   /usr/local/lib/python3.7/dist-packages/ipykernel launcher.py:15: VisibleDeprecationWarnir
     from ipykernel import kernelapp as app
path = '/content/drive/MyDrive/COVID/train/Covid Positive'
path1 = '/content/drive/MyDrive/COVID/test/Covid Positive'
path2 = '/content/drive/MyDrive/COVID/train/Covid Negative'
path3 = '/content/drive/MyDrive/COVID/test/Covid Negative'
covidpositives = len([f for f in os.listdir(path)if os.path.isfile
covidnegatives = len([f for f in os.listdir(path2)if os.path.isfil
Cats = ['Covid Positive', 'Covid Negative']
y pos = np.arange(len(Cats))
barlist = plt.bar(y pos,[covidpositives, covidnegatives], align='c
barlist[0].set color('g')
barlist[1].set color('r')
plt.xticks(y_pos,['Covid Positive', 'Covid Negative'])
plt.ylabel('Number of Cases')
plt.title('Coronavirus Cases and Categories')
plt.show()
```

```
Coronavirus Cases and Categories

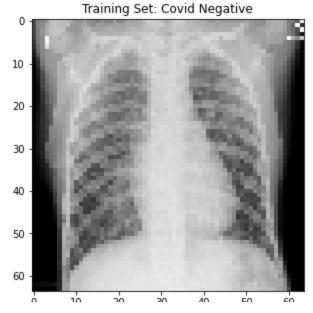
1200 -
1000 -
8800 -
600 -
```

```
path = '/content/drive/MyDrive/COVID/test/Covid Negative'
path1 = '/content/drive/MyDrive/COVID/test/Covid Positive'
path2 = '/content/drive/MyDrive/COVID/train/Covid Negative'
path3 = '/content/drive/MyDrive/COVID/train/Covid Positive'
Test = len([f for f in os.listdir(path)if os.path.isfile(os.path.j
Train = len([f for f in os.listdir(path2)if os.path.isfile(os.path
# Data to plot
labels = 'Training', 'Testing'
sizes = [Train, Test]
colors = ['gold', 'yellowgreen']
explode = (0, 0) # explode 1st slice
# Plot
plt.pie(sizes, explode=explode, labels=labels, colors=colors,
autopct='%1.1f%%', shadow=True, startangle=140)
plt.axis('equal')
plt.show()
```

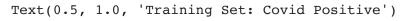


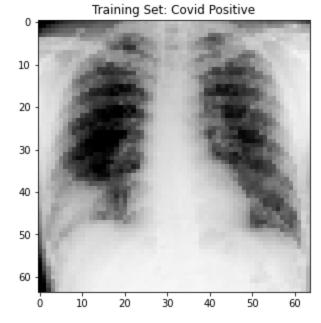
```
plt.figure(figsize = (5,5))
plt.imshow(train[1][0])
plt.title('Training Set: Covid Negative')
```

Text(0.5, 1.0, 'Training Set: Covid Negative')



```
plt.figure(figsize = (5,5))
plt.imshow(train[-1][0])
plt.title('Training Set: Covid Positive')
```





```
x_train = []
y_train = []
x_test = []
y_test = []

for feature, label in train:
    x_train.append(feature)
    y_train.append(label)
```

```
for feature, label in test:
    x test.append(feature)
   y test.append(label)
  # Normalize the data
 x_train = np.array(x_train) / 255
 x \text{ test} = np.array(x \text{ test}) / 255
 x train.reshape(-1, img size, img size, 1)
 y train = np.array(y train)
 x test.reshape(-1, img size, img size, 1)
 y test = np.array(y test)
  datagen = ImageDataGenerator(
          featurewise_center=False, # set input mean to 0 over the
          samplewise_center=False, # set each sample mean to 0
          featurewise_std_normalization=False, # divide inputs by s
          samplewise std normalization=False, # divide each input h
          zca_whitening=False, # apply ZCA whitening
          rotation_range = 30, # randomly rotate images in the range
          zoom range = 0.2, # Randomly zoom image
          width shift range=0.1, # randomly shift images horizontal
          height_shift_range=0.1, # randomly shift images verticall
          horizontal flip = True, # randomly flip images
          vertical flip=True) # randomly flip images
 datagen.fit(x train)
 X train, X val, y train, y val = train test split(x train, y trair
Experiment 1
```

model.add(layers.Conv2D(filters=32, kernel_size=(3, 3), strides=(1

model.add(layers.MaxPooling2D((2, 2),strides=2))

model.add(layers.Dense(units=32, activation=tf.nn.relu))

model = models.Sequential()

model.add(layers.Flatten())

```
model.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
 model.compile(optimizer='adam',
                loss=tf.keras.losses.BinaryCrossentropy(),
                metrics=['accuracy'])
  %%time
 history = model.fit(X_train,
                      y train,
                      validation data = (X val, y val),
                      epochs=20,
                      batch size=512
  import numpy as np
 loss, accuracy = model.evaluate(x test, y test)
 print('test set accuracy: ', accuracy * 100)
Experiment 2
 model2 = models.Sequential()
 model2.add(layers.Conv2D(filters=64, kernel size=(3, 3), strides=(
 model2.add(layers.MaxPooling2D((2, 2),strides=2))
 model2.add(layers.Flatten())
 model2.add(layers.Dense(units=64, activation=tf.nn.relu))
 model2.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
```

```
batch_size=512
)

loss, accuracy = model2.evaluate(x_test, y_test)
print('test set accuracy: ', accuracy * 100)
```

▼ Experiment 3

```
model3 = models.Sequential()
model3.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides=
model3.add(layers.MaxPooling2D((2, 2),strides=2))
model3.add(layers.Flatten())
model3.add(layers.Dense(units=128, activation=tf.nn.relu))
model3.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
model3.compile(optimizer='adam',
              loss=tf.keras.losses.BinaryCrossentropy(),
              metrics=['accuracy'])
%%time
history3 = model3.fit(X_train,
                    y train,
                    validation data = (X val, y val),
                    epochs=20,
                    batch size=512
loss, accuracy = model3.evaluate(x test, y test)
print('test set accuracy: ', accuracy * 100)
```

▼ Experiment 4

```
model4 = models.Sequential()
model4.add(layers.Conv2D(filters=32, kernel_size=(3, 3), strides=(
model4.add(layers.MaxPooling2D((2, 2), strides=2))
model4.add(layers.Conv2D(filters=32, kernel_size=(3, 3), strides=(3, 3), strides=(3, 3), strides=(3, 3)
```

```
moder4.add(rayers.maxPoorring2D((2, 2),Strides-2))
 model4.add(layers.Flatten())4
 model4.add(layers.Dense(units=256, activation=tf.nn.relu))
 model4.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
 model4.compile(optimizer='adam',
                loss=tf.keras.losses.BinaryCrossentropy(),
                metrics=['accuracy'])
  %%time
 history4 = model4.fit(X train,
                      y train,
                      validation data = (X val, y val),
                      epochs=20,
                      batch size=512
  loss, accuracy = model4.evaluate(x test, y test)
 print('test set accuracy: ', accuracy * 100)
▼ Experiment 5
 model5 = models.Sequential()
 model5.add(layers.Conv2D(filters=64, kernel size=(3, 3), strides=(
 model5.add(layers.MaxPooling2D((2, 2),strides=2))
 model5.add(layers.Conv2D(filters=64, kernel_size=(3, 3), strides=(
 model5.add(layers.MaxPooling2D((2, 2),strides=2))
 model5.add(layers.Flatten())
 model5.add(layers.Dense(units=256, activation=tf.nn.relu))
 model5.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
 model5.compile(optimizer='adam',
                loss=tf.keras.losses.BinaryCrossentropy(),
                metrics=['accuracy'])
  %%time
 history5 = model5.fit(X train,
                      y_train,
                      validation data = (X val, y val),
```

epochs=20.

```
batch_size=512
)

loss, accuracy = model5.evaluate(x_test, y_test)
print('test set accuracy: ', accuracy * 100)
```

Experiment 6

```
model6 = models.Sequential()
model6.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides=
model6.add(layers.MaxPooling2D((2, 2),strides=2))
model6.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides=
model6.add(layers.MaxPooling2D((2, 2),strides=2))
model6.add(layers.Flatten())
model6.add(layers.Dense(units=256, activation=tf.nn.relu))
model6.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
model6.compile(optimizer='adam',
              loss=tf.keras.losses.BinaryCrossentropy(),
              metrics=['accuracy'])
%%time
history6 = model6.fit(X_train,
                    y train,
                    validation data = (X val, y val),
                    epochs=20,
                    batch size=512
loss, accuracy = model6.evaluate(x test, y test)
print('test set accuracy: ', accuracy * 100)
```

Experiment 7

```
model7 = models.Sequential()
model7.add(layers.Conv2D(filters=64, kernel_size=(3, 3), strides=()
model7.add(layers.MaxPooling2D((2, 2), strides=2))
model7.add(layers.Conv2D(filters=64, kernel_size=(3, 3), strides=()
```

```
model7.add(layers.MaxPooling2D((2, 2),strides=2))
 model7.add(layers.Conv2D(filters=64, kernel size=(3, 3), strides=(
 model7.add(layers.MaxPooling2D((2, 2),strides=2))
 model7.add(layers.Flatten())
 model7.add(layers.Dense(units=256, activation=tf.nn.relu))
 model7.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
 model7.compile(optimizer='adam',
                loss=tf.keras.losses.BinaryCrossentropy(),
                metrics=['accuracy'])
 %%time
 history7 = model7.fit(X train,
                      y train,
                      validation data = (X val, y val),
                      epochs=20,
                      batch size=512
                     )
 loss, accuracy = model7.evaluate(x test, y test)
 print('test set accuracy: ', accuracy * 100)
Experiment 8
 from keras.layers import AveragePooling2D
 model8 = models.Sequential()
 model8.add(layers.Conv2D(filters=64, kernel size=(3, 3), strides=(
 model8.add(layers.AveragePooling2D((2, 2),strides=2))
 model8.add(layers.Conv2D(filters=64, kernel size=(3, 3), strides=(
 model8.add(layers.AveragePooling2D((2, 2),strides=2))
 model8.add(layers.Conv2D(filters=64, kernel size=(3, 3), strides=(
 model8.add(layers.AveragePooling2D((2, 2),strides=2))
 model8.add(layers.Flatten())
 model8.add(layers.Dense(units=256, activation=tf.nn.relu))
```

model8.add(layers.Dense(units=1, activation=tf.nn.sigmoid))

```
%%time
 history8 = model8.fit(X_train,
                     y train,
                     validation data = (X_val, y_val),
                     epochs=20,
                     batch size=512
 loss, accuracy = model8.evaluate(x test, y test)
 print('test set accuracy: ', accuracy * 100)
Experiment 9
 model9 = models.Sequential()
 model9.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides=
 model9.add(layers.MaxPooling2D((2, 2),strides=2))
 model9.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides=
 model9.add(layers.MaxPooling2D((2, 2),strides=2))
 model9.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides=
 model9.add(layers.MaxPooling2D((2, 2),strides=2))
 model9.add(layers.Flatten())
 model9.add(layers.Dense(units=256, activation=tf.nn.relu))
 model9.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
 model9.compile(optimizer='adam',
               loss=tf.keras.losses.BinaryCrossentropy(),
               metrics=['accuracy'])
  %%time
 history9 = model9.fit(X train,
                     y train,
                     validation data = (X val, y val),
                     epochs=20,
                     batch size=512
                    )
     Epoch 1/20
```

4/4 [================] - 1s 146ms/step - loss: 0.6498 - accuracy: 0.7436 -

Epoch 2/20

```
Epoch 3/20
   Epoch 4/20
   4/4 [==============] - 1s 148ms/step - loss: 0.3948 - accuracy: 0.8584 -
   Epoch 5/20
   Epoch 6/20
   4/4 [============== ] - 1s 146ms/step - loss: 0.1593 - accuracy: 0.9396 -
   Epoch 7/20
   4/4 [============== ] - 1s 148ms/step - loss: 0.1313 - accuracy: 0.9500 -
   Epoch 8/20
   4/4 [=============== ] - 1s 144ms/step - loss: 0.1165 - accuracy: 0.9555 -
   Epoch 9/20
   4/4 [============== ] - 1s 145ms/step - loss: 0.0998 - accuracy: 0.9671 -
   Epoch 10/20
   4/4 [===========] - 1s 146ms/step - loss: 0.0963 - accuracy: 0.9609 -
   Epoch 11/20
   4/4 [============= ] - 1s 145ms/step - loss: 0.0561 - accuracy: 0.9823 -
   Epoch 12/20
   4/4 [============== ] - 1s 146ms/step - loss: 0.0782 - accuracy: 0.9717 -
   Epoch 13/20
   4/4 [============== ] - 1s 146ms/step - loss: 0.0618 - accuracy: 0.9727 -
   Epoch 14/20
   Epoch 15/20
   4/4 [============== ] - 1s 147ms/step - loss: 0.0717 - accuracy: 0.9764 -
   Epoch 16/20
   Epoch 17/20
   4/4 [============== ] - 1s 148ms/step - loss: 0.0480 - accuracy: 0.9832 -
   Epoch 18/20
   4/4 [============= ] - 1s 146ms/step - loss: 0.0365 - accuracy: 0.9877 -
   Epoch 19/20
   Epoch 20/20
   4/4 [============= ] - 1s 143ms/step - loss: 0.0350 - accuracy: 0.9848 -
   CPU times: user 7.58 s, sys: 3.79 s, total: 11.4 s
   Wall time: 12.9 s
loss, accuracy = model9.evaluate(x test, y test)
print('test set accuracy: ', accuracy * 100)
   test set accuracy: 98.47561120986938
y_pred = (model9.predict(X_train) > 0.5).astype("int32")
confusion matrix(y train, y pred)
   array([[787, 14],
       [ 5, 794]])
f1_score(y_train, y_pred, average='macro')
   0.9881247727007274
```

recall score(v train, v pred, average='macro')

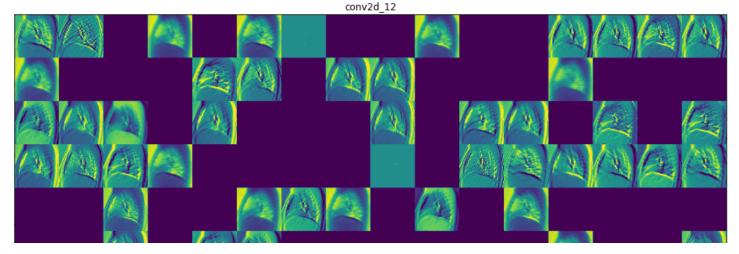
```
0.9881320127062698
precision score(y train, y pred, average='macro')
   0.9881800680068007
y pred = (model9.predict(x test) > 0.5).astype("int32")
confusion matrix(y test, y pred)
   array([[162, 2],
        [ 3, 161]])
f1 score(y test, y pred, average='macro')
   0.9847559558666332
recall score(y test, y pred, average='macro')
   0.9847560975609756
precision_score(y_test, y_pred, average='macro')
   0.9847741215839375
from keras.preprocessing import image
import numpy as np
img tensor = image.img to array(x test[2])
img tensor = np.expand dims(img tensor, axis=0)
# Remember that the model was trained on inputs
# that were preprocessed in the following way:
img tensor /= 255.
from keras import models
# Extracts the outputs of the top 8 layers:
layer outputs = [layer.output for layer in model9.layers[:2]]
# Creates a model that will return these outputs, given the model
activation_model = models.Model(inputs=model9.input, outputs=layer
```

```
# This will return a list of 5 Numpy arrays:
# one array per layer activation
activations = activation model.predict(img tensor)
first layer activation = activations[-1]
print(first layer activation.shape)
import keras
# These are the names of the layers, so can have them as part of o
layer names = []
for layer in model9.layers[:8]:
    layer names.append(layer.name)
images per row = 16
# Now let's display our feature maps
for layer name, layer activation in zip(layer names, activations):
    # This is the number of features in the feature map
    n features = layer activation.shape[-1]
    # The feature map has shape (1, size, size, n features)
    size = layer activation.shape[1]
    # We will tile the activation channels in this matrix
    n cols = n features // images per row
    display grid = np.zeros((size * n cols, images per row * size)
    # We'll tile each filter into this big horizontal grid
    for col in range(n cols):
        for row in range(images per row):
            channel_image = layer_activation[0,
                                             col * images per row
            # Post-process the feature to make it visually palatak
            channel image -= channel image.mean()
            channel image /= channel image.std()
            channel image *= 64
            channel image += 128
            channel image = np.clip(channel image, 0, 255).astype(
            display grid[col * size : (col + 1) * size,
                        ---- + --- · /--- · 1) + --- - - -----
```

row * size : (row + r) * size = channer_

NING:tensorflow:11 out of the last 11 calls to <function Model.make_predict_function.<locally, 31, 128)

r/local/lib/python3.7/dist-packages/ipykernel_launcher.py:57: RuntimeWarning: invalid value



pred_classes = (model9.predict(x_test) > 0.5).astype("int32").rave
pred_classes

```
0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1,
dtype=int32)
```

layer_outputs = [layer.output for layer in model9.layers]
activation_model = models.Model(inputs=model9.input, outputs=layer
layer_outputs

Get the outputs of all the hidden nodes for each of the 60000 tr

```
output layer activations = activations[8]
hidden layer activation.shape # each of the 128 hidden nodes ha
   (328, 256)
#Get the dataframe of all the node values
activation data = {'pred class':pred classes[0:328]}
for k in range(0,256):
    activation_data[f"act_val_{k}"] = hidden_layer_activation[:,k]
activation df = pd.DataFrame(activation data)
activation df.head()
      pred_class act_val_0 act_val_1 act_val_2 act_val_3 act_val_4 act_val_5 act_val_6
    0
                     0.0
                          0.261437
                                       0.0
                                               0.0
                                                    2.302884
                                                                 0.0
                                                                         0.0
                          0.000000
    1
              0
                     0.0
                                       0.0
                                               0.0
                                                    2.508843
                                                                 0.0
                                                                         0.0
    2
                     0.0
                          0.000000
                                                                         0.0
              0
                                       0.0
                                               0.0
                                                    2.587324
                                                                 0.0
                     0.0
                          0.000000
                                               0.0
                                                    2.679549
                                                                         0.0
    3
              0
                                       0.0
                                                                 0.0
    4
              0
                     0.0
                          0.000000
                                       0.0
                                               0.0
                                                    3.288160
                                                                 0.0
                                                                         0.0
   5 rows × 257 columns
# Separating out the features
features = [*activation_data][1:] # ['act_val_0', 'act_val_1',...]
x = activation df.loc[:, features].values
pca = PCA(n_components=3)
principalComponents = pca.fit transform(x)
principalDf = pd.DataFrame(data = principalComponents
               , columns = ['pca-one', 'pca-two', 'pca-three'])
principalDf.head()
```

accivación moaci-picaicc(x

hidden layer activation = activations[7]

	pca-one	pca-two	pca-three
0	-5.853893	2.599480	1.266800
1	-9.155988	1.404219	-0.198507
2	-10.286340	2.204174	-0.435203
3	-8.915661	1.304460	0.064735
4	-12.442112	5.414866	-0.132032

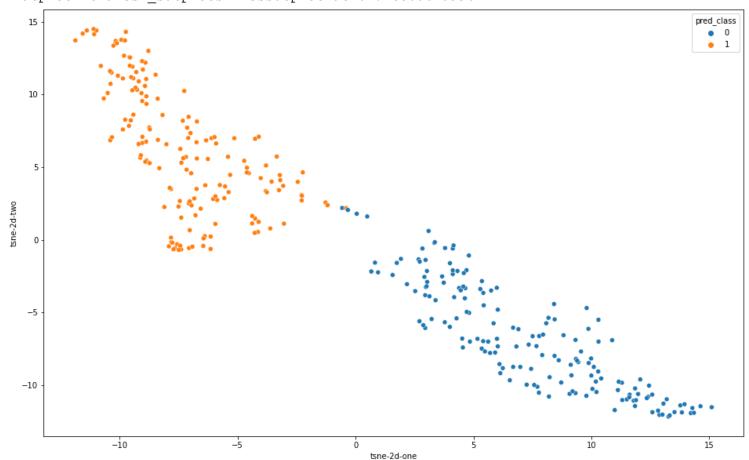
```
activation_pca_df = pd.concat([principalDf, activation_df[['pred_c
activation pca df.head()
```

```
pca-one pca-two pca-three pred_class
    0 -5.853893 2.599480
                       1.266800
    1 -9.155988 1.404219
                       -0.198507
    2 -10.286340 2.204174
                       -0.435203
    3 -8.915661 1.304460
                       0.064735
    4 -12.442112 5.414866 -0.132032
N = 10000
activation df subset = activation df.iloc[:N].copy()
activation df subset.shape
    (328, 257)
data subset = activation df subset[features].values
data subset.shape
    (328, 256)
from sklearn.manifold import TSNE
tsne = TSNE(n components=2, verbose=1, perplexity=40, n iter=300)
tsne results = tsne.fit transform(data subset)
    [t-SNE] Computing 121 nearest neighbors...
    [t-SNE] Indexed 328 samples in 0.004s...
    [t-SNE] Computed neighbors for 328 samples in 0.040s...
    [t-SNE] Computed conditional probabilities for sample 328 / 328
    [t-SNE] Mean sigma: 1.609862
    [t-SNE] KL divergence after 250 iterations with early exaggeration: 52.930374
    [t-SNE] KL divergence after 300 iterations: 0.244041
activation df subset['tsne-2d-one'] = tsne results[:,0]
activation df subset['tsne-2d-two'] = tsne results[:,1]
plt.figure(figsize=(16,10))
sns.scatterplot(
    x="tsne-2d-one", y="tsne-2d-two",
    hue="pred class",
    palette=sns.color palette(n colors = 2),
    data=activation df subset,
```

legend="full"

```
alpha = 1
```

<matplotlib.axes. subplots.AxesSubplot at 0x7fb30d64a990>



Experiment 10

```
model10 = models.Sequential()
model10.add(layers.Conv2D(filters=256, kernel_size=(3, 3), strides
model10.add(layers.MaxPooling2D((2, 2), strides=2))
model10.add(layers.Conv2D(filters=256, kernel_size=(3, 3), strides
model10.add(layers.MaxPooling2D((2, 2), strides=2))
model10.add(layers.Conv2D(filters=256, kernel_size=(3, 3), strides
model10.add(layers.MaxPooling2D((2, 2), strides=2))
model10.add(layers.Flatten())
model10.add(layers.Dense(units=256, activation=tf.nn.relu))
model10.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
```

```
model10.compile(optimizer='adam',
                loss=tf.keras.losses.BinaryCrossentropy(),
                metrics=['accuracy'])
   %%time
 history10 = model10.fit(X train,
                      y train,
                      validation data = (X val, y val),
                      epochs=20,
                      batch size=512
  loss, accuracy = model10.evaluate(x test, y test)
 print('test set accuracy: ', accuracy * 100)
Experiment 11
 model11 = models.Sequential()
 model11.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides
 model11.add(layers.MaxPooling2D((2, 2),strides=2))
 model11.add(layers.Dropout(.2))
 model11.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides
 model11.add(layers.MaxPooling2D((2, 2),strides=2))
 model11.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides
 model11.add(layers.MaxPooling2D((2, 2),strides=2))
 model11.add(layers.Flatten())
 model11.add(layers.Dense(units=256, activation=tf.nn.relu))
 model11.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
 model11.compile(optimizer='adam',
                loss=tf.keras.losses.BinaryCrossentropy(),
                metrics=['accuracy'])
   %%time
 history11 = model11.fit(X train,
                      y train,
```

validation data = (X_val, y_val),

epochs=20,

batch size=512

```
loss, accuracy = model11.evaluate(x_test, y_test)
print('test set accuracy: ', accuracy * 100)
```

Experiment 12

```
model12 = models.Sequential()
model12.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides
model12.add(layers.MaxPooling2D((2, 2),strides=2))
model12.add(layers.Dropout(.2))
model12.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides
model12.add(layers.MaxPooling2D((2, 2),strides=2))
model11.add(layers.Dropout(.2))
model12.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides
model12.add(layers.MaxPooling2D((2, 2),strides=2))
model12.add(layers.Flatten())
model12.add(layers.Dense(units=256, activation=tf.nn.relu))
model12.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
model12.compile(optimizer='adam',
              loss=tf.keras.losses.BinaryCrossentropy(),
              metrics=['accuracy'])
 %%time
history12 = model12.fit(X train,
                    y train,
                    validation data = (X val, y val),
                    epochs=20,
                    batch size=512
loss, accuracy = model12.evaluate(x_test, y_test)
print('test set accuracy: ', accuracy * 100)
```

Experiment 13

model13 = models.Sequential()

```
model13.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides
model13.add(layers.MaxPooling2D((2, 2),strides=2))
model13.add(layers.Dropout(.2))
model13.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides
model13.add(layers.MaxPooling2D((2, 2),strides=2))
model13.add(layers.Dropout(.2))
model13.add(layers.Conv2D(filters=128, kernel size=(3, 3), strides
model13.add(layers.MaxPooling2D((2, 2),strides=2))
model13.add(layers.Dropout(.2))
model13.add(layers.Flatten())
model13.add(layers.Dense(units=256, activation=tf.nn.relu))
model13.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
model13.compile(optimizer='adam',
              loss=tf.keras.losses.BinaryCrossentropy(),
              metrics=['accuracy'])
 %%time
history13 = model13.fit(X train,
                    y train,
                    validation data = (X val, y val),
                    epochs=20,
                    batch size=512
loss, accuracy = model13.evaluate(x_test, y_test)
print('test set accuracy: ', accuracy * 100)
```

Experiment 14

model14 = models.Sequential()

```
model14.add(layers.Conv2D(filters=256, kernel size=(3, 3), strides
model14.add(layers.MaxPooling2D((2, 2),strides=2))
model14.add(layers.Conv2D(filters=256, kernel size=(3, 3), strides
model14.add(layers.MaxPooling2D((2, 2),strides=2))
model14.add(layers.Conv2D(filters=256, kernel size=(3, 3), strides
model14.add(layers.MaxPooling2D((2, 2),strides=2))
model14.add(layers.Flatten())
model14.add(layers.Dense(units=256, activation=tf.nn.relu))
```

```
model14.add(layers.Dense(units=1, activation=ti.nn.sigmoid))
 model14.compile(optimizer='adam',
                loss=tf.keras.losses.BinaryCrossentropy(),
                metrics=['accuracy'])
   %%time
 history14 = model14.fit(X train,
                      y train,
                      validation_data = (X_val, y_val),
                      epochs=20,
                      batch size=512,
                     callbacks = [tf.keras.callbacks.EarlyStopping(m
  loss, accuracy = model14.evaluate(x test, y test)
 print('test set accuracy: ', accuracy * 100)
▼ Experiment 15
 model15 = models.Sequential()
 model15.add(layers.Conv2D(filters=256, kernel size=(3, 3), strides
 model15.add(layers.MaxPooling2D((2, 2),strides=2))
 model15.add(layers.Conv2D(filters=256, kernel size=(3, 3), strides
 model15.add(layers.MaxPooling2D((2, 2),strides=2))
 model15.add(layers.Conv2D(filters=256, kernel size=(3, 3), strides
 model15.add(layers.MaxPooling2D((2, 2),strides=2))
 model15.add(layers.Flatten())
 model15.add(layers.Dense(units=256, activation=tf.nn.relu))
 model15.add(layers.Dense(units=1, activation=tf.nn.sigmoid))
 model15.compile(optimizer='adam',
                loss=tf.keras.losses.BinaryCrossentropy(),
                metrics=['accuracy'])
   %%time
 history15 = model15.fit(X train,
                      y train,
                      validation data = (X_val, y_val),
                      epochs=20,
```

hatch size=512

```
callbacks = [tf.keras.callbacks.EarlyStopping(m
 loss, accuracy = model15.evaluate(x test, y test)
 print('test set accuracy: ', accuracy * 100)
Experiment 16
 model16 = models.Sequential()
 model16.add(layers.Conv2D(filters=256, kernel size=(3, 3), strides
 model16.add(layers.MaxPooling2D((2, 2),strides=2))
 model16.add(layers.Conv2D(filters=256, kernel size=(3, 3), strides
 model16.add(layers.MaxPooling2D((2, 2),strides=2))
 model16.add(layers.Conv2D(filters=256, kernel size=(3, 3), strides
 model16.add(layers.MaxPooling2D((2, 2),strides=2))
 model16.add(layers.Flatten())
 model16.add(layers.Dense(units=256, activation=tf.nn.relu))
 model16.add(layers.Dense(units=1, activation=tf.nn.sigmoid, kernel
 model16.compile(optimizer='adam',
              loss=tf.keras.losses.BinaryCrossentropy(),
             metrics=['accuracy'])
  %%time
 history16 = model16.fit(X train,
                   y train,
                   validation data = (X val, y val),
                   epochs=20,
                   batch size=512)
    Epoch 1/20
    Epoch 2/20
    4/4 [============] - 1s 262ms/step - loss: 0.8844 - accuracy: 0.5331 -
    Epoch 3/20
    Epoch 4/20
    Epoch 5/20
    4/4 [============== ] - 1s 263ms/step - loss: 0.6602 - accuracy: 0.8174 -
    Epoch 6/20
    Epoch 7/20
    4/4 [============== ] - 1s 260ms/step - loss: 0.3564 - accuracy: 0.9365 -
```

```
Epoch 8/20
   4/4 [===============] - 1s 260ms/step - loss: 0.3122 - accuracy: 0.9358 -
   Epoch 9/20
   4/4 [============== ] - 1s 259ms/step - loss: 0.3390 - accuracy: 0.9288 -
   Epoch 10/20
   4/4 [=============== ] - 1s 261ms/step - loss: 0.2863 - accuracy: 0.9482 -
   Epoch 11/20
   4/4 [============== ] - 1s 260ms/step - loss: 0.2545 - accuracy: 0.9641 -
   Epoch 12/20
   4/4 [============= ] - 1s 261ms/step - loss: 0.2311 - accuracy: 0.9642 -
   Epoch 13/20
   4/4 [============== ] - 1s 266ms/step - loss: 0.1969 - accuracy: 0.9788 -
   Epoch 14/20
   4/4 [============= ] - 1s 263ms/step - loss: 0.2070 - accuracy: 0.9730 -
   Epoch 15/20
   4/4 [===========] - 1s 261ms/step - loss: 0.1974 - accuracy: 0.9742 -
   Epoch 16/20
   4/4 [===============] - 1s 259ms/step - loss: 0.1684 - accuracy: 0.9816 -
   Epoch 17/20
   4/4 [============= ] - 1s 263ms/step - loss: 0.1659 - accuracy: 0.9813 -
   Epoch 18/20
   4/4 [============] - 1s 267ms/step - loss: 0.1605 - accuracy: 0.9809 -
   Epoch 19/20
   4/4 [============= ] - 1s 267ms/step - loss: 0.1694 - accuracy: 0.9806 -
   Epoch 20/20
   4/4 [============= ] - 1s 263ms/step - loss: 0.1550 - accuracy: 0.9809 -
   CPU times: user 19.5 s, sys: 14.3 s, total: 33.8 s
   Wall time: 1min 7s
loss, accuracy = model16.evaluate(x test, y test)
print('test set accuracy: ', accuracy * 100)
   test set accuracy: 97.86585569381714
```

Experiment 17

model17.compile(optimizer='adam',

```
model17 = models.Sequential()
model17.add(layers.Conv2D(filters=256, kernel_size=(3, 3), strides
model17.add(layers.MaxPooling2D((2, 2), strides=2))
model17.add(layers.Conv2D(filters=256, kernel_size=(3, 3), strides
model17.add(layers.MaxPooling2D((2, 2), strides=2))
model17.add(layers.Conv2D(filters=256, kernel_size=(3, 3), strides
model17.add(layers.MaxPooling2D((2, 2), strides=2))
model17.add(layers.Flatten())
model17.add(layers.Dense(units=256, activation=tf.nn.relu))
model17.add(layers.Dense(units=1, activation=tf.nn.sigmoid, kernel
```

loss=tf.keras.losses.BinaryCrossentropy(),

```
metrics=['accuracy'])
```

```
%%time
history17 = model17.fit(X train,
                  y train,
                  validation data = (X val, y val),
                  epochs=20,
                  batch size=512)
   Epoch 1/20
   4/4 [===============] - 2s 323ms/step - loss: 0.8549 - accuracy: 0.5537 -
   Epoch 2/20
   4/4 [===============] - 1s 270ms/step - loss: 0.7012 - accuracy: 0.4921 -
   Epoch 3/20
   4/4 [============== ] - 1s 269ms/step - loss: 0.6303 - accuracy: 0.6534 -
   Epoch 4/20
   4/4 [===============] - 1s 272ms/step - loss: 0.6281 - accuracy: 0.6660 -
   Epoch 5/20
   Epoch 6/20
   4/4 [============== ] - 1s 272ms/step - loss: 0.3985 - accuracy: 0.8938 -
   Epoch 7/20
   4/4 [============] - 1s 272ms/step - loss: 0.2443 - accuracy: 0.9416 -
   Epoch 8/20
   4/4 [=============== ] - 1s 276ms/step - loss: 0.2414 - accuracy: 0.9226 -
   Epoch 9/20
   4/4 [============== ] - 1s 276ms/step - loss: 0.2119 - accuracy: 0.9321 -
   Epoch 10/20
   Epoch 11/20
   4/4 [================ ] - 1s 269ms/step - loss: 0.1328 - accuracy: 0.9552 -
   Epoch 12/20
   4/4 [============== ] - 1s 276ms/step - loss: 0.1518 - accuracy: 0.9455 -
   Epoch 13/20
   4/4 [============] - 1s 275ms/step - loss: 0.1502 - accuracy: 0.9485 -
   Epoch 14/20
   Epoch 15/20
   4/4 [============= ] - 1s 280ms/step - loss: 0.0969 - accuracy: 0.9733 -
   Epoch 16/20
   4/4 [================ ] - 1s 273ms/step - loss: 0.0864 - accuracy: 0.9765 -
   Epoch 17/20
   Epoch 18/20
   4/4 [============] - 1s 272ms/step - loss: 0.0676 - accuracy: 0.9834 -
   Epoch 19/20
   Epoch 20/20
   CPU times: user 11.5 s, sys: 7.28 s, total: 18.8 s
   Wall time: 23.7 s
```

loss, accuracy = model17.evaluate(x test, y test)

print('test set accuracy: ', accuracy * 100)

test set accuracy: 98.17073345184326