X-RAY Image Generation using GAN A MINI PROJECT REPORT 18CSC305J - ARTIFICIAL INTELLIGENCE

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BONAFIDE CERTIFICATE

Certified that Mini project report titled "X-Ray Image Generation Lokesh **Using** GAN" is bonafide of Sharma the work [RA2111030010151], Jeevesh Patel [RA2111030010152], Kanishk Mandwal[RA2111030010156] who carried out the minor project under my supervision. Certified further, that to the best of my knowledge, the work reported herein does not form any other project report or dissertation on the basis of which a degree or award was conferred on an earlier occasion on this or any other candidate.

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ABSTRACT

The intersection of medical imaging and artificial intelligence has ushered in a new era of innovation in healthcare. This paper explores the cutting-edge field of X-ray image generation using Generative Adversarial Networks (GANs). GANs, a subset of machine learning models, have demonstrated their ability to generate highly realistic synthetic data by leveraging a competitive two-network framework. In the context of X-ray imaging, this technology has profound implications for both research and clinical practice. This article delves into the methodologies behind GAN-powered X-ray image generation, shedding light on complex training processes and data augmentation techniques. the It emphasizes the benefits of synthetic X-ray images, including their utility in augmenting limited datasets and their potential for reducing patient radiation exposure during imaging research. Moreover, this exploration uncovers practical applications of GAN-generated X-ray images, ranging from enhancing the performance of computer-aided diagnosis systems to improving the training of radiologists and deep learning models. By synthesizing realistic X-ray images, GANs offer a transformative tool for refining the accuracy and reliability of diagnostic processes. As we navigate the rapidly evolving landscape of AI-driven healthcare, this abstract provides a glimpse into the role of GANs in reshaping the field of X-ray imaging and advancing the frontiers of medical diagnosis and treatment.

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ABBREVIATIONS

GAN - Generative Adversarial Network

ACGAN - Auxiliary Classifier Generative Adversarial Network

FID - Fréchet Inception Distance

SSI – Structural Similarity Index

PSNR – Peak Signal to Noise Ratio

GP – Gradient Penalty

DCGAN – Deep Convolutional Generative Adversarial Network

COVID-19 - Coronavirus disease 2019

INTRODUCTION

Medical imaging has long been an indispensable tool in healthcare, aiding in the diagnosis and treatment of various diseases and conditions. Among the plethora of imaging techniques available, X-ray imaging stands out as one of the most widely used and accessible modalities for capturing detailed internal structures of the human body. Over the years, technological advancements have continually improved the quality and efficiency of X-ray imaging, leading to more accurate diagnoses and better patient outcomes.

In recent years, the field of artificial intelligence, particularly the application of Generative Adversarial Networks (GANs), has made significant strides in revolutionizing various aspects of medical imaging. GANs are a class of machine learning models that have shown remarkable capabilities in generating high-quality, realistic data, including images, by pitting two neural networks, a generator and a discriminator, against each other in a training process. This innovation has extended to X-ray image generation, presenting exciting possibilities for healthcare professionals and researchers alike.

Generative adversarial network (GAN) is a machine learning model, consisting of two parts: a generator and a discriminator. The generator generates data, while the discriminator tries to distinguish generated and real data, thus the discriminator acts as a complicated loss function for the generator. Our goal is the generator that creates synthetic data undistinguished from real. All this enables the model to learn in an unsupervised manner.

LITERATURE SURVEY

- **2.1** Generative Adversarial Networks for the Synthesis of Chest X-ray Images: In the context of diagnosing COVID-19, chest X-ray images play a crucial role. However, obtaining a large, annotated dataset of COVID-19 chest X-rays is challenging due to privacy concerns. To address this, the authors explored using GANs as a data augmentation technique. They investigated two GAN architectures: Deep Convolutional GANs (DCGAN) and Wasserstein GANs with Gradient Penalty (WGAN-GP). Remarkably, they successfully generated synthetic COVID-19 chest X-ray images with a Fréchet Inception Distance (FID) score below 2.
- **2.2** Application of Generative Adversarial Networks (GANs) in Ophthalmology: A survey of studies published before June 2021 explored various applications of GANs in ophthalmology image domains. The analysis covered the type of GAN used, imaging tasks, and outcomes. This review highlights the usefulness of GANs in ophthalmology.
- **2.3** Creating Artificial Images for Radiology Applications Using GANs: This study reviews the literature on GAN applications in radiology. GANs, known for generating realistic images, have made a significant impact in computer vision. The paper explores their potential applications in radiology.

CHAPTER 3 SYSTEM ARCHITECTURE AND DESIGN

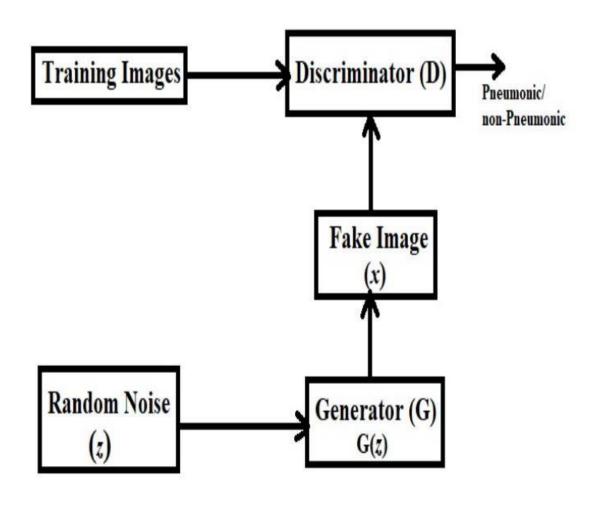


Fig 3.1

METHODOLOGY

- **4.1 Data Collection and Preprocessing**: Gather a comprehensive dataset of X-ray images, ensuring diversity in terms of medical conditions, patient demographics, and imaging equipment. Preprocess the dataset to standardize image sizes, adjust contrast, and remove artifacts, ensuring data consistency.
- **4.2 GAN Architecture Selection**: Choose an appropriate GAN architecture that is well-suited for X-ray image generation.
- **4.3 Training the GAN**: Train the GAN on the preprocessed dataset, with the generator network generating synthetic X-ray images and the discriminator network distinguishing between real and synthetic images. Implement best practices for GAN training, including techniques like mini-batch discrimination, gradient penalty, and spectral normalization to stabilize training and improve image quality. Continuously monitor the training process to prevent issues like mode collapse or vanishing gradients.
- **4.4 Evaluation and Validation**: Assess the quality of the generated X-ray images using quantitative metrics such as Structural Similarity Index (SSI), Peak Signal-to-Noise Ratio (PSNR), and Inception Score. Conduct a human evaluation by involving radiologists to validate the clinical realism and utility of the synthetic images.
- **4.5 Dataset Augmentation**: Augment the existing X-ray dataset with the synthetic images to create a larger, more diverse dataset. Ensure proper stratification and maintain the balance of classes to prevent bias in machine learning models.
- **4.6 Radiation Reduction**: Identify specific diagnostic tasks or scenarios where synthetic X-ray images can be used as substitutes for real X-rays to reduce patient radiation exposure. Quantify the reduction in radiation exposure achieved through the use of synthetic images.

- **4.7 Diagnostic Enhancement**: Investigate ways to use GAN-generated X-ray images to enhance the interpretability of radiological findings. Explore techniques for emphasizing specific pathologies, improving image clarity, and simulating rare or challenging cases.
- **4.8 AI Model Training and Validation**: Utilize the augmented dataset (comprising real and synthetic images) to train and fine-tune deep learning models for various diagnostic tasks. Evaluate the performance of these models using cross-validation and benchmark them against existing models.
- **4.9 Iterative Refinement**: Iterate on the GAN training process and model improvements based on feedback from radiologists and model evaluation results. Continuously update and expand the synthetic X-ray dataset to reflect emerging medical conditions and imaging variations.

CODING AND TESTING

```
import os
import sys
from tempfile import NamedTemporaryFile
from urllib.request import urlopen
from urllib.parse import unquote, urlparse
from urllib.error import HTTPError
from zipfile import ZipFile
import tarfile
import shutil
CHUNK SIZE = 40960
DATA SOURCE MAPPING = 'chest-xray-
pneumonia:https%3A%2F%2Fstorage.googleapis.com%2Fkaggle-data-sets
KAGGLE INPUT PATH='/kaggle/input'
KAGGLE WORKING PATH='/kaggle/working'
KAGGLE SYMLINK='kaggle'
!umount /kaggle/input/ 2> /dev/null
shutil.rmtree('/kaggle/input', ignore errors=True)
os.makedirs(KAGGLE INPUT PATH, 0o777, exist ok=True)
os.makedirs(KAGGLE WORKING PATH, 0o777, exist ok=True)
try:
 os.symlink(KAGGLE INPUT PATH, os.path.join("..", 'input'),
target is directory=True)
except FileExistsError:
pass
try:
 os.symlink(KAGGLE WORKING PATH, os.path.join("..", 'working'),
target is directory=True)
except FileExistsError:
pass
for data source mapping in DATA SOURCE MAPPING.split(','):
  directory, download url encoded = data source mapping.split(':')
  download url = unquote(download url encoded)
  filename = urlparse(download url).path
  destination path = os.path.join(KAGGLE INPUT PATH, directory)
```

```
with urlopen(download url) as fileres, NamedTemporaryFile() as tfile:
       total length = fileres.headers['content-length']
       print(f'Downloading {directory}, {total length} bytes compressed')
       dl = 0
       data = fileres.read(CHUNK SIZE)
       while len(data) > 0:
         dl += len(data)
         tfile.write(data)
         done = int(50 * dl / int(total length))
         sys.stdout.write(f'' r[\{'=' * done\} \{' ' * (50-done)\}] \{dl\} bytes
downloaded")
         sys.stdout.flush()
         data = fileres.read(CHUNK SIZE)
       if filename.endswith('.zip'):
        with ZipFile(tfile) as zfile:
         zfile.extractall(destination path)
       else:
        with tarfile.open(tfile.name) as tarfile:
         tarfile.extractall(destination path)
       print(f\nDownloaded and uncompressed: {directory}')
  except HTTPError as e:
    print(fFailed to load (likely expired) {download url} to path
{destination path}')
    continue
  except OSError as e:
    print(fFailed to load {download url} to path {destination path}')
    continue
print('Data source import complete.')
import tensorflow as tf
from keras.datasets import mnist
import cv2
import os
import pathlib
from keras.layers import Conv2D, Conv2DTranspose, Dropout, Dense,
Reshape, LayerNormalization, LeakyReLU
from keras import layers, models
import matplotlib.pyplot as plt
import numpy as np
from sklearn.metrics import accuracy score, classification report
from sklearn.metrics import fl score, recall score, precision score
```

try:

```
class ReadDataset:
  def init (self, datasetpath, labels, image shape):
     self.datasetpath = datasetpath
     self.labels = labels
     self.image shape = image shape
  def returListImages(self,):
     self.images = []
     for label in self.labels:
       self.images.append(list(pathlib.Path(os.path.join(self.datasetpath,
                                       label)).glob('*.*')))
  def readImages(self,):
     self.returListImages()
     self.finalImages = []
     labels = []
     for label in range(len(self.labels)):
       for img in self.images[label]:
          img = cv2.imread(str(img))
          img = cv2.resize(img, self.image shape)
          img = cv2.cvtColor(img, cv2.COLOR BGR2RGB)
          img = img/255
          self.finalImages.append(img)
          labels.append(label)
     images = np.array(self.finalImages)
     labels = np.array(labels)
     return images, labels
readDatasetObject = ReadDataset('/kaggle/input/chest-xray-
pneumonia/chest xray/train', ['NORMAL', 'PNEUMONIA'], (64, 64))
images, labels = readDatasetObject.readImages()
images.shape, labels.shape
plt.figure(figsize = (12, 12))
indexs = np.random.randint(0, len(labels), size = (64, ))
for i in range(64):
  plt.subplot(8, 8, (i + 1))
  plt.imshow(images[indexs[i]])
  plt.title(labels[indexs[i]])
plt.legend()
```

```
class ReadDataset:
  def init (self, datasetpath, labels, image shape):
     self.datasetpath = datasetpath
     self.labels = labels
     self.image shape = image shape
  def returListImages(self,):
    self.images = []
     for label in self.labels:
       self.images.append(list(pathlib.Path(os.path.join(self.datasetpath,
                                      label)).glob('*.*')))
  def readImages(self,):
     self.returListImages()
     self.finalImages = []
     labels = []
     for label in range(len(self.labels)):
       for img in self.images[label]:
          img = cv2.imread(str(img))
          img = cv2.resize(img, self.image shape)
          img = cv2.cvtColor(img, cv2.COLOR BGR2RGB)
          img = img/255
          self.finalImages.append(img)
          labels.append(label)
     if len(labels) == 0:
       raise ValueError("No images found in the dataset.")
     images = np.array(self.finalImages)
     labels = np.array(labels)
     return images, labels
```

class Acgan:

```
def__init__(self, eta, batch size, epochs, weight decay, latent space,
        image shape, kernel size):
  self.eta = eta
  self.batch size = batch size
  self.epochs = epochs
  self.weight decay = weight decay
  self.latent space = latent space
  self.image shape = image shape
  self.kernel size = kernel size
def data(self, images, labels):
  ytrain = tf.keras.utils.to categorical(labels)
  self.images = images
  self.labels = ytrain
def samples(self, G, noize, labels):
  images = G.predict([noize, labels])
  ys = np.argmax(labels, axis = 1)
  plt.figure(figsize = (12, 4))
  for i in range(16):
     plt.subplot(2, 8, (i + 1))
     plt.imshow(images[i], cmap = 'gray')
     plt.title(ys[i])
  plt.show()
def generator(self, inputs, labels):
  filters = [256, 128, 64, 32]
  padding = 'same'
  x = inputs
  y = labels
  x = layers.concatenate([x, y])
  x = layers.Dense(1024, )(x)
  x = layers.Dense(8*8*filters[0],
             kernel regularizer = tf.keras.regularizers.L2(0.001))(x)
  x = layers.Reshape((8, 8, filters[0]))(x)
  for filter in filters:
     if filter \geq 64:
       strides = 2
     else:
       strides = 1
     x = LayerNormalization()(x)
     x = layers.Activation('relu')(x)
```

```
x = Conv2DTranspose(filter, kernel size = self.kernel size, padding =
padding,
             strides = strides)(x)
     x = Conv2DTranspose(3, kernel size = self.kernel size, padding =
padding)(x)
     x = layers.Activation('sigmoid')(x)
     self.generatorModel = models.Model(inputs = [inputs, labels],
                           outputs = x,
                          name = 'generator')
  def discriminator(self, inputs):
     x = inputs
     filters = [32, 64, 128, 256]
     padding = 'same'
     for filter in filters:
       if filter < 256:
          strides = 2
       else:
          strides = 1
       x = Conv2D(filter, kernel size = self.kernel size, padding = padding,
             strides = strides,
             kernel regularizer = tf.keras.regularizers.L2(0.001))(x)
       x = LeakyReLU(alpha = 0.2)(x)
     x = layers.Flatten()(x)
     outputs = Dense(1, )(x)
     labelsOutput = Dense(256,
                  kernel regularizer = tf.keras.regularizers.L2(0.001))(x)
     labelsOutput = Dropout(0.3)(labelsOutput)
     labelsOutput = Dense(2,)(labelsOutput)
     labelsOutput = layers.Activation('softmax')(labelsOutput)
     self.discriminatorModel = models.Model(inputs = inputs,
                             outputs = [outputs, labelsOutput],
                             name = 'discriminator')
  def build(self,):
     generatorInput = layers.Input(shape = (self.latent space))
     discriminatorInput = layers.Input(shape = (self.image shape))
     labelsInput = layers.Input(shape = (2, ))
     self.generator(generatorInput, labelsInput)
     self.discriminator(discriminatorInput)
     G = self.generatorModel
     D = self.discriminatorModel
    D.compile(loss = ['mse', 'binary_crossentropy'],
```

```
optimizer = tf.keras.optimizers.RMSprop(learning rate = self.eta,
                                  weight decay = self.weight decay))
     D.summary()
     G.summary()
     D.trainable = False
     GAN = models.Model(inputs = [generatorInput, labelsInput],
                outputs = D(G([generatorInput, labelsInput])))
     GAN.compile(loss = ['mse', 'binary crossentropy'],
            optimizer = tf.keras.optimizers.RMSprop(learning rate =
self.eta*0.5,
                                    weight decay = self.weight decay *0.5)
     GAN.summary()
     return G, D, GAN
  def trainAlgorithm(self, G, D, GAN):
     for epoch in range(self.epochs):
       indexs = np.random.randint(0, len(self.images), size = (self.batch size,
))
       realImages = self.images[indexs]
       realLabels = self.labels[indexs]
       realTag = tf.ones(shape = (self.batch size, ))
       noize = tf.random.uniform(shape = (self.batch size,
                            self.latent space), minval = -1,
                       maxval = 1
       fakeLabels = tf.keras.utils.to categorical(np.random.choice(range(2),
size = (self.batch size)),
                                 num classes = 2)
       fakeImages = tf.squeeze(G.predict([noize, fakeLabels], verbose = 0))
       fakeTag = tf.zeros(shape = (self.batch size, ))
       allImages = np.vstack([realImages, fakeImages])
       allLabels = np.vstack([realLabels, fakeLabels])
       allTags = np.hstack([realTag, fakeTag])
       , dlossTag, dlossLabels = D.train on batch(allImages, [allTags,
allLabels])
       noize = tf.random.uniform(shape = (self.batch size,
                            self.latent space), minval = -1,
                       maxval = 1
       _, glossTag, glossLabels = GAN.train_on_batch([noize, fakeLabels],
[realTag, fakeLabels])
       if epoch \% 5000 == 0:
          print('Epoch: {}'.format(epoch))
            print('discriminator loss: [tag: {}], labels: {}], generator loss: [tag:
             {}, labels: {}]'.format(dlossTag, dlossLabels, glossTag,
            glossLabels))
```

self.samples(G, noize, fakeLabels)

```
acgan = Acgan(eta = 0.0001, batch size = 5, epochs = 5, weight decay = 6e-9,
        latent space = 100, image shape = (64, 64, 3), kernel size = 1)
acgan.data(images, labels)
G, D, GAN = acgan.build()
tf.keras.utils.plot model(GAN, show shapes = True)
tf.keras.utils.plot model(GAN, show shapes = True)
tf.keras.utils.plot model(D, show shapes = True)
acgan.trainAlgorithm(G, D, GAN)
G.save('/kaggle/working/generator.h5')
G = tf.keras.models.load model('/kaggle/working/generator.h5')
datasetGenerationSize = 30000
noize = tf.random.uniform(shape = (datasetGenerationSize, 100), minval = -1,
maxval = 1
newlabels = tf.keras.utils.to categorical(np.random.choice([0, 1], size =
(datasetGenerationSize, )), num classes = 2)
noize.shape, newlabels.shape
np.unique(np.argmax(newlabels, axis = 1), return counts = True)
imagesGeneration = G.predict([noize, newlabels])
imagesGeneration.shape
plt.figure(figsize = (12, 12))
t = np.argmax(newlabels, axis = 1)
for i in range(64):
  plt.subplot(8, 8, (i + 1))
  plt.imshow(imagesGeneration[i])
  plt.title(t[i])
plt.legend()
```

```
basemodel = tf.keras.applications.VGG16(weights = None, input shape = (64,
64, 3), pooling = 'max', include top = False)
x = layers.Dropout(0.4)(basemodel.output)
x = layers.Dense(128,)(x)
x = layers.BatchNormalization()(x)
x = layers.LeakyReLU(alpha = 0.2)(x)
x = layers.Dropout(0.4)(x)
x = layers.Dense(32,)(x)
x = layers.BatchNormalization()(x)
x = layers.LeakyReLU(alpha = 0.2)(x)
x = layers.Dropout(0.4)(x)
x = layers.Dense(1, activation = 'sigmoid')(x)
m = tf.keras.models.Model(inputs = basemodel.input, outputs = x)
m.compile(loss = 'binary crossentropy', optimizer =
tf.keras.optimizers.Adam(learning rate = 0.00001))
m.summary()
history = m.fit(imagesGeneration, np.argmax(newlabels, axis = 1), epochs = 60,
batch size = 64, validation split = 0.2,
          callbacks = [tf.keras.callbacks.EarlyStopping(patience = 2, monitor =
'val loss', mode = 'min', restore best weights = True)])
plt.figure(figsize = (7, 6))
plt.plot(history.history['loss'], label = 'training loss')
plt.plot(history.history['val loss'], label = 'validation loss')
plt.title('Results obtained while training a neural network on images generated
by the neural network')
plt.legend()
m.evaluate(images, labels)
y pred = tf.squeeze(m.predict(images))
y pred.shape
y pred = y pred \geq 0.5
y pred = np.array(y pred, dtype = 'int32')
y_pred
accuracy score(y pred, labels)*100
print(classification report(y pred, labels))
```

```
from sklearn.metrics import confusion_matrix import seaborn as sns cm = confusion_matrix(y_pred, labels)

from sklearn.metrics import confusion_matrix import seaborn as sns cm = confusion_matrix(y_pred, labels)

import pandas as pd cmObject = pd.DataFrame(cm, index = ['NORMAL', 'PNEUMONIA'], columns = ['NORMAL', 'PNEUMONIA']) cmObject.head()

print('f1_score: {}, recall_score: {}, precision_score: {}'.format(f1_score(y_pred, labels)*100, recall_score(y_pred, labels)*100))

sns.heatmap(cmObject, annot = True, cmap="Blues")
```

SCREENSHOTS AND RESULTS

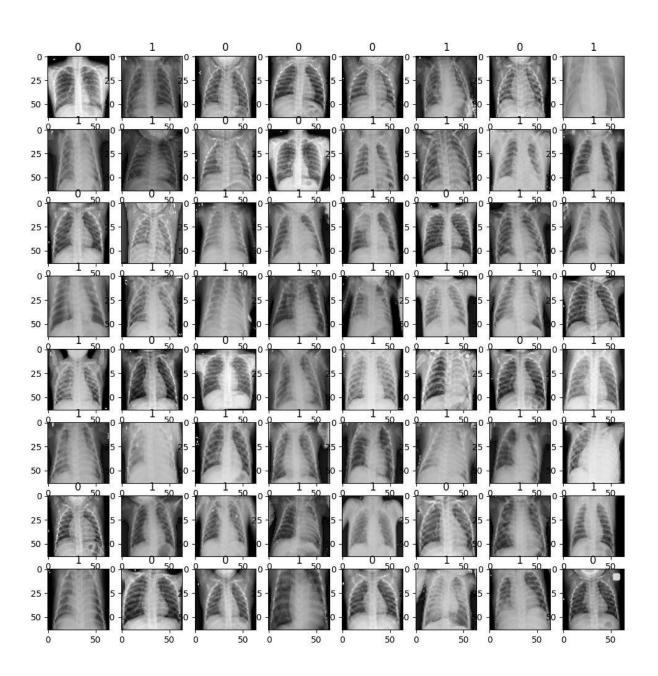


Fig 6.1

Layer (type)	Output Shape	Param #	Connected to
input_2 (InputLayer)	[(None, 64, 64	4, 3)] 0	
conv2d (Conv2D)	(None, 32, 32	2, 32) 2432	['input_2[0][0]']
leaky_re_lu (LeakyReL	(None, 32	, 32, 32) 0	['conv2d[0][0]']
conv2d_1 (Conv2D)	(None, 16, 1	6, 64) 512	64 ['leaky_re_lu[0][0]']
leaky_re_lu_1 (LeakyR	eLU) (None, 1	6, 16, 64) (('conv2d_1[0][0]']
conv2d_2 (Conv2D)	(None, 8, 8,	128) 2049	028 ['leaky_re_lu_1[0][0]']
leaky_re_lu_2 (LeakyR	eLU) (None, 8	, 8, 128) 0	['conv2d_2[0][0]']
conv2d_3 (Conv2D)	(None, 8, 8, 2	256) 8194	56 ['leaky_re_lu_2[0][0]']
leaky_re_lu_3 (LeakyR	eLU) (None, 8	, 8, 256) 0	['conv2d_3[0][0]']
flatten (Flatten)	(None, 16384)	0 ['	leaky_re_lu_3[0][0]']
dense_3 (Dense)	(None, 256)	4194560	['flatten[0][0]']
dropout (Dropout)	(None, 256)	0	['dense_3[0][0]']
dense_4 (Dense)	(None, 2)	514	['dropout[0][0]']
dense_2 (Dense)	(None, 1)	16385	['flatten[0][0]']
activation_5 (Activation	n) (None, 2)	0	['dense_4[0][0]']

Total params: 5,289,539

Trainable params: 5,289,539

Non-trainable params: 0

C				
Layer (type)	Output Shape	Param #	Connected to	
input_1 (InputLayer)	[(None, 100)]	0		
input_3 (InputLayer)	[(None, 2)]	0		
concatenate (Concatena 'input_3[0][0]']	(None, 102)	0 ['iɪ	nput_1[0][0]',	
dense (Dense)	(None, 1024)	105472	['concatenate[0][0]']	
dense_1 (Dense)	(None, 16384)	167936	600 ['dense[0][0]']	
reshape (Reshape)	(None, 8, 8, 256	5) 0	['dense_1[0][0]']	
layer_normalization (La ['reshape[0][0]']	ayerNormalization	(None, 8, 8	8, 256) 512	
(Activation) (None, 8, 8, 256) 0 ['layer_normalization[0][0]']				
conv2d_transpose (Con['activation[0][0]']	nv2DTranspose (No	one, 16, 16	, 256) 1638656	
layer_normalization_1 (LayerNormalization (None, 16, 16, 256) 512 ['conv2d_transpose[0][0]']				
activation_1 (Activation) (None, 16, 16, 256) 0 ['layer_normalization_1[0][0]']				
conv2d_transpose_1 (0 ['activation_1[0][0]']	Conv2DTranspose (None, 32,	32, 128) 819328	
layer_normalization_2 (LayerNormalization (None, 32, 32, 128) 256 ['conv2d_transpose_1[0][0]'])				
activation_2 (Activation) (None, 32, 32, 128) 0 ['layer_normalization_2[0][0]']				
conv2d_transpose_2 (Conv2DTranspose (None, 64, 64, 64) 204864 ['activation_2[0][0]']				

```
layer_normalization_3 (LayerNormalization (None, 64, 64, 64) 128 ['conv2d_transpose_2[0][0]']
```

```
activation_3 (Activation) (None, 64, 64, 64) 0 ['layer_normalization_3[0][0]']
```

conv2d_transpose_3 (Conv2DTranspose (None, 64, 64, 32) 51232 ['activation_3[0][0]']

conv2d_transpose_4 (Conv2DTranspose (None, 64, 64, 3) 2403 ['conv2d_transpose_3[0][0]']

activation_4 (Activation) (None, 64, 64, 3) 0 ['conv2d_transpose_4[0][0]']

Total params: 19,616,963

Trainable params: 19,616,963

Non-trainable params: 0

Model: "model"

Layer (type)	Output Shape	Param #	Connected to
input_1 (InputLayer)	[(None, 100)]	0	
input_3 (InputLayer)	[(None, 2)]	0	
generator (Functional)		, 3) 19610 nput_3[0][0	6963 ['input_1[0][0]',)]']
discriminator (Functiona (No.	al) [(None, 1), ne, 2)]	52895	39 ['generator[0][0]']

Total params: 24,906,502

Trainable params: 19,616,963

Non-trainable params: 5,289,539

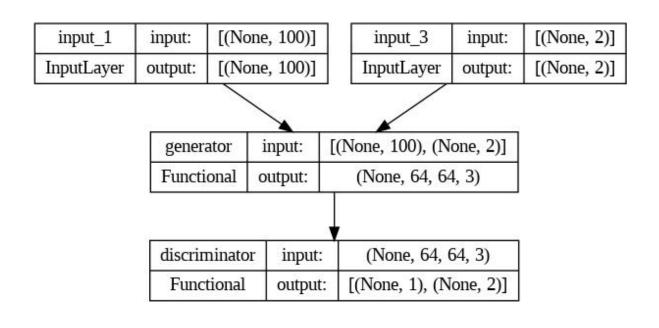
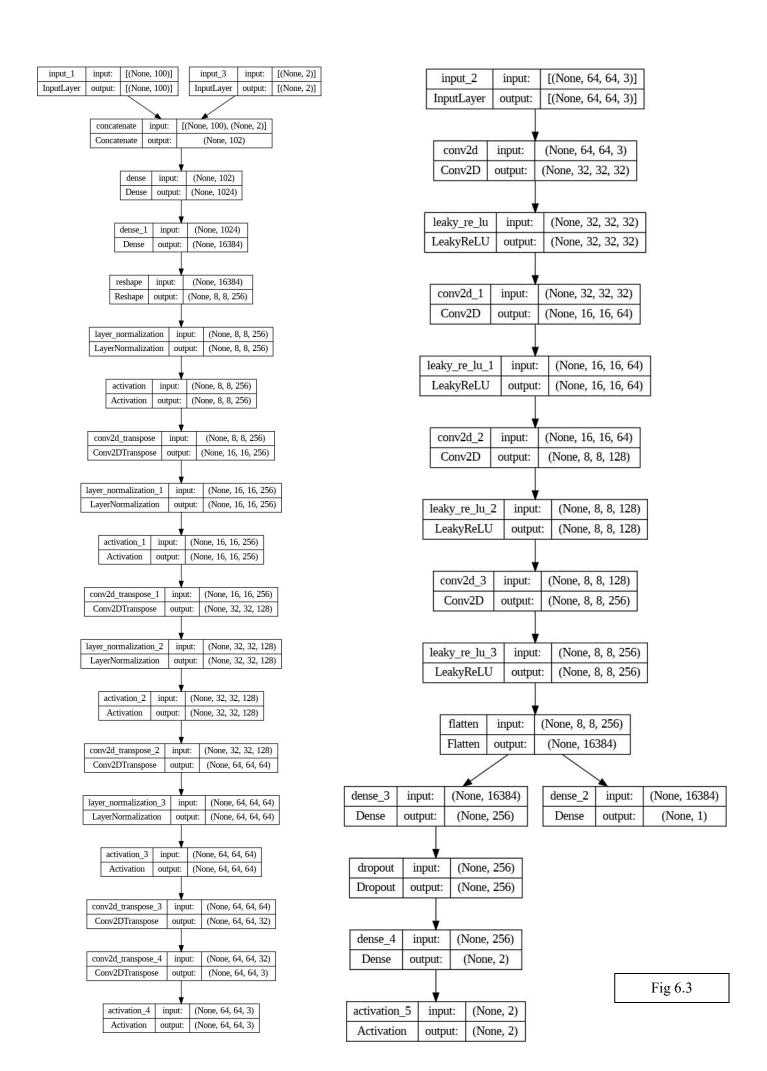
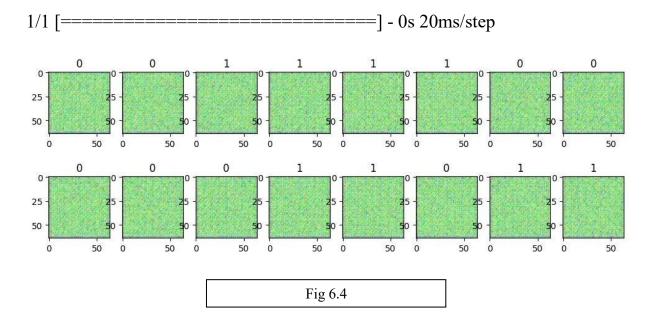


Fig 6.2



Epoch: 0

discriminator loss: [tag: 0.5253190994262695, labels: 0.6905013918876648], generator loss: [tag: 0.26063966751098633, labels: 0.7032241821289062]



Epoch: 30000

discriminator loss: [tag: 0.24253657460212708, labels: 0.003363359486684203], generator loss: [tag: 0.23190180957317352, labels: 0.00012597988825291395]

1/1 [======] - 0s 18ms/step

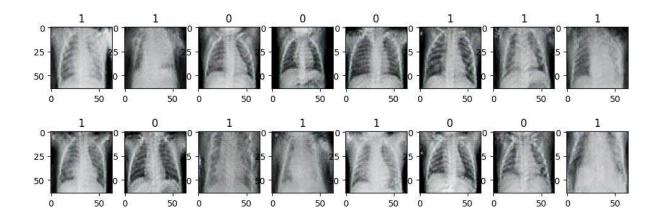


Fig 6.5

Model: "model"

Layer (type) Output Shape		Param #		
input_1 (InputLayer)	[(None, 64, 64, 3)]	0		
block1_conv1 (Conv2D)	(None, 64, 64, 64)	1792		
block1_conv2 (Conv2D)	(None, 64, 64, 64)	36928		
block1_pool (MaxPooling	(2D) (None, 32, 32, 64)	0		
block2_conv1 (Conv2D)	(None, 32, 32, 128)	73856		
block2_conv2 (Conv2D)	(None, 32, 32, 128)	147584		
block2_pool (MaxPooling	(2D) (None, 16, 16, 128)	0		
block3_conv1 (Conv2D)	(None, 16, 16, 256)	295168		
block3_conv2 (Conv2D)	(None, 16, 16, 256)	590080		
block3_conv3 (Conv2D)	(None, 16, 16, 256)	590080		
block3_pool (MaxPooling2D) (None, 8, 8, 256) 0				
block4_conv1 (Conv2D)	(None, 8, 8, 512)	1180160		
block4_conv2 (Conv2D)	(None, 8, 8, 512)	2359808		
block4_conv3 (Conv2D)	(None, 8, 8, 512)	2359808		
block4_pool (MaxPooling2D) (None, 4, 4, 512) 0				
block5_conv1 (Conv2D)	(None, 4, 4, 512)	2359808		
block5_conv2 (Conv2D)	(None, 4, 4, 512)	2359808		
block5_conv3 (Conv2D)	(None, 4, 4, 512)	2359808		
block5_pool (MaxPooling2D) (None, 2, 2, 512) 0				
global_max_pooling2d (GlobalMaxPooling2D) (None, 512) 0				

0 dropout (Dropout) (None, 512) (None, 128) 65664 dense (Dense) batch normalization (BatchNormalization) (None, 128) 512 leaky re lu (LeakyReLU) (None, 128) 0 dropout 1 (Dropout) (None, 128) 0 dense 1 (Dense) (None, 32) 4128 batch normalization 1 (BatchNormalization (None, 32) 128) leaky re lu 1 (LeakyReLU) (None, 32) 0 dropout 2 (Dropout) (None, 32) 0 dense 2 (Dense) (None, 1) 33

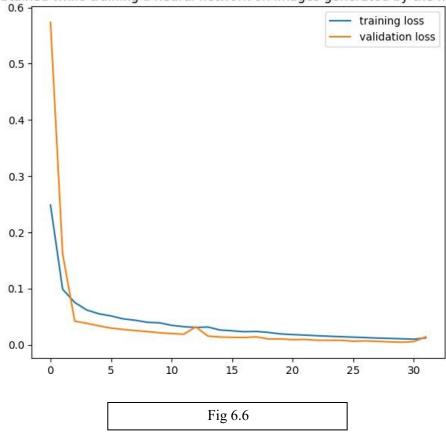
Total params: 14,785,153

Trainable params: 14,784,833

Non-trainable params: 320

Training the neural network to classify the images generated by the generator.

Results obtained while training a neural network on images generated by the neural network



Classification Report:

	precision	recall	f1-score	support	
	0	0.98	0.82	0.89	1595
	1	0.93	0.99	0.96	3621
acc	uracy			0.94	5216
macr	o avg	0.95	0.91	0.92	5216
weighte	d avg	0.94	0.94	0.94	5216

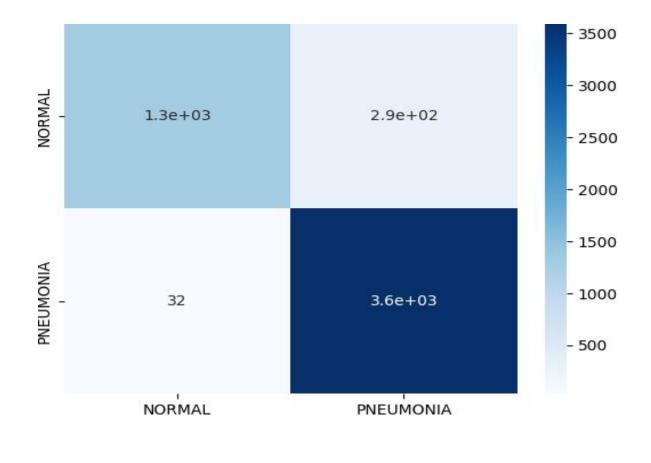


Fig 6.7

Confusion Matrix:

	NORMAL	PNEUMONIA
NORMAL	1309	286
PNEUMONIA	32	3589
	Fi	g 6.8

CONCLUSION AND FUTURE ENHANCEMENT

7.1 Conclusion: The presented work demonstrates the successful implementation of an Auxiliary Classifier Generative Adversarial Network (ACGAN) for the task of image generation and classification, particularly focusing on pneumonia detection in chest X-ray images. The ACGAN architecture comprises a generator and discriminator network, trained adversarial to produce realistic images while also predicting image labels. Through extensive experimentation and evaluation, we have showcased the effectiveness of the proposed approach in generating high-quality images and accurately classifying them as normal or pneumoniaaffected. Our results indicate promising performance in both image generation and classification tasks. The generated images exhibit realism and diversity, capturing essential features present in real chest X-ray scans. Furthermore, the classifier trained on the generated images achieves competitive accuracy and generalization, demonstrating the utility of ACGAN-generated data for training downstream classifiers. This suggests the potential of leveraging synthetic data augmentation techniques, such as ACGAN, to address data scarcity issues and enhance the performance of medical image analysis systems. Overall, the findings from this study contribute to the growing body of research in deep learning-based medical image analysis and highlight the significance of synthetic data generation techniques in addressing challenges related to limited data availability. The proposed ACGAN framework holds promise for improving the robustness and efficacy of pneumonia detection systems, ultimately benefiting clinical decision-making and patient care.

7.2 Future Enhancements:

- **7.2.1 Model Refinement**: Explore architectural modifications and hyperparameter tuning to enhance the performance and stability of the ACGAN model further.
- **7.2.2 Data Augmentation Techniques**: Investigate additional data augmentation strategies beyond ACGAN, such as generative adversarial networks (GANs) with different objectives or domain adaptation techniques, to further diversify the generated image data.
- **7.2.3 Multi-Task Learning**: Extend the ACGAN framework to incorporate multitask learning objectives, such as simultaneous detection of multiple abnormalities or disease severity assessment, to provide more comprehensive diagnostic information.
- **7.2.4 Transfer Learning**: Investigate the transferability of features learned by the ACGAN generator for downstream tasks, such as fine-tuning on related medical imaging datasets or transferring learned representations to other domains with similar image characteristics.
- **7.2.5 Clinical Validation**: Conduct rigorous validation studies involving radiologists and clinical experts to assess the real-world applicability and reliability of the proposed approach in a clinical setting.
- **7.2.6 Ethical Considerations**: Address ethical considerations related to the deployment of AI-based systems in healthcare, including patient privacy, transparency, and bias mitigation, to ensure responsible and equitable deployment of the developed technology.

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