```
# IMPORTANT: RUN THIS CELL IN ORDER TO IMPORT YOUR KAGGLE DATA SOURCES,
# THEN FEEL FREE TO DELETE THIS CELL.
# NOTE: THIS NOTEBOOK ENVIRONMENT DIFFERS FROM KAGGLE'S PYTHON
# ENVIRONMENT SO THERE MAY BE MISSING LIBRARIES USED BY YOUR
# NOTEBOOK.
#import kagglehub
#paultimothymooney_chest_xray_pneumonia_path = kagglehub.dataset_download('paultimothymooney #awsaf49_pneumonia_chest_xray_npy_path = kagglehub.dataset_download('awsaf49/pneumonia-chest
#print('Data source import complete.')
```

## Import Libraries

```
import numpy as np # linear algebra
import pandas as pd
from tqdm import tqdm
from glob import glob
import os
import numpy as np
import matplotlib.pyplot as plt
#import tensorflow as tf
```

#### !pip install -U tensorflow

```
Requirement already satisfied: packaging in /usr/local/lib/python3.11/dist-packages (
Requirement already satisfied: protobuf!=4.21.0,!=4.21.1,!=4.21.2,!=4.21.3,!=4.21.4,!

Requirement already satisfied: requests<3,>=2.21.0 in /usr/local/lib/python3.11/dist-

Requirement already satisfied: setuptools in /usr/local/lib/python3.11/dist-packages

Requirement already satisfied: six>=1.12.0 in /usr/local/lib/python3.11/dist-packages

Requirement already satisfied: termcolor>=1.1.0 in /usr/local/lib/python3.11/dist-packages

Requirement already satisfied: typing-extensions>=3.6.6 in /usr/local/lib/python3.11/dist-package

Requirement already satisfied: wrapt>=1.11.0 in /usr/local/lib/python3.11/dist-packag

Requirement already satisfied: grpcio<2.0,>=1.24.3 in /usr/local/lib/python3.11/dist-

Collecting tensorboard~=2.19.0 (from tensorflow)
```

```
Requirement already satisfied: optree in /usr/local/lib/python3.11/dist-packages (fro A
Requirement already satisfied: charset-normalizer<4,>=2 in /usr/local/lib/python3.11/
Requirement already satisfied: idna<4,>=2.5 in /usr/local/lib/python3.11/dist-package
Requirement already satisfied: urllib3<3,>=1.21.1 in /usr/local/lib/python3.11/dist-p
Requirement already satisfied: certifi>=2017.4.17 in /usr/local/lib/python3.11/dist-p
Requirement already satisfied: markdown>=2.6.8 in /usr/local/lib/python3.11/dist-pack
Requirement already satisfied: tensorboard-data-server<0.8.0,>=0.7.0 in /usr/local/li
Requirement already satisfied: werkzeug>=1.0.1 in /usr/local/lib/python3.11/dist-pack
Requirement already satisfied: MarkupSafe>=2.1.1 in /usr/local/lib/python3.11/dist-pa
Requirement already satisfied: markdown-it-py>=2.2.0 in /usr/local/lib/python3.11/dis
Requirement already satisfied: pygments<3.0.0,>=2.13.0 in /usr/local/lib/python3.11/d
Requirement already satisfied: mdurl~=0.1 in /usr/local/lib/python3.11/dist-packages
Downloading tensorflow-2.19.0-cp311-cp311-manylinux 2 17 x86 64.manylinux2014 x86 64.
                                          - 644.9/644.9 MB 2.7 MB/s eta 0:00:00
Downloading ml dtypes-0.5.1-cp311-cp311-manylinux 2 17 x86 64.manylinux2014 x86 64.wh
                                          - 4.7/4.7 MB 64.3 MB/s eta 0:00:00
Downloading tensorboard-2.19.0-py3-none-any.whl (5.5 MB)
                                         -- 5.5/5.5 MB 75.4 MB/s eta 0:00:00
Installing collected packages: ml-dtypes, tensorboard, tensorflow
 Attempting uninstall: ml-dtypes
    Found existing installation: ml-dtypes 0.4.1
   Uninstalling ml-dtypes-0.4.1:
      Successfully uninstalled ml-dtypes-0.4.1
 Attempting uninstall: tensorboard
    Found existing installation: tensorboard 2.18.0
   Uninstalling tensorboard-2.18.0:
      Successfully uninstalled tensorboard-2.18.0
 Attempting uninstall: tensorflow
    Found existing installation: tensorflow 2.18.0
   Uninstalling tensorflow-2.18.0:
      Successfully uninstalled tensorflow-2.18.0
ERROR: pip's dependency resolver does not currently take into account all the package
tensorflow-text 2.18.1 requires tensorflow<2.19,>=2.18.0, but you have tensorflow 2.1
tensorflow-decision-forests 1.11.0 requires tensorflow==2.18.0, but you have tensorfl
tf-keras 2.18.0 requires tensorflow<2.19,>=2.18, but you have tensorflow 2.19.0 which
Successfully installed ml-dtypes-0.5.1 tensorboard-2.19.0 tensorflow-2.19.0
```

# Extrating files

from google.colab import drive
drive.mount('/content/drive')

→ Mounted at /content/drive

!ls /content/drive/MyDrive/pfc2T1

**→** archive

```
import numpy as np

base_path = '/content/drive/MyDrive/pfc2T1/archive/Array128/Array128'
!ls '/content/drive/MyDrive/pfc2T1/Array128/Array128'
normal = np.load(f'{base_path}/train_Normal_128.npy')
viral = np.load(f'{base_path}/train_Virus_128.npy')
bacterial = np.load(f'{base_path}/train_bacteria_128.npy')
```

🚁 ls: cannot access '/content/drive/MyDrive/pfc2T1/Array128/Array128': No such file or dir

# Loading Training Files

I have converted all images to numpy array to boost speed

```
#normal = np.load('../input/pneumonia-chest-xray-npy/Array128/Array128/train_Normal_128.npy'
#viral = np.load('../input/pneumonia-chest-xray-npy/Array128/Array128/train_Virus_128.npy')
#bacterial = np.load('../input/pneumonia-chest-xray-npy/Array128/Array128/train_bacteria_128

normal.shape, viral.shape, bacterial.shape

((1341, 128, 128, 1), (1345, 128, 128, 1), (2530, 128, 128, 1))
```

#### Create Labels

```
label_normal = np.zeros(len(normal))
label_bacterial = np.ones(len(bacterial))
label_viral = np.full(len(viral),2, dtype = int)

train_data = np.concatenate((normal,bacterial,viral),axis=0)
train_label = np.concatenate((label_normal,label_bacterial,label_viral),axis=0)

train_label.shape, train_data.shape

((5216,), (5216, 128, 128, 1))
```

### Visualization

#### Normal

```
n_row = 2
n_col = 4

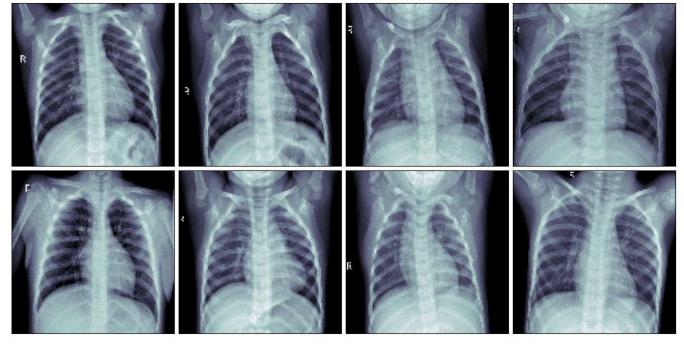
fig, ax = plt.subplots(n_row, n_col, figsize = (n_col*3, n_row*3), constrained_layout = True

for row in tqdm(range(n_row)):

    for col in range(n_col):

        ax[row][col].imshow(normal[row*n_col + col,:,:,0], cmap = 'bone')
        ax[row][col].set_xticks([])
        ax[row][col].set_yticks([])
```

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### Viral Pneumonia

```
n_row = 2
n_col = 4

fig, ax = plt.subplots(n_row, n_col, figsize = (n_col*3, n_row*3), constrained_layout = True

for row in tqdm(range(n_row)):
    for col in range(n_col):
        ax[row][col].imshow(viral[row*n_col + col,:,:,0], cmap = 'bone')
        ax[row][col].set_xticks([])
        ax[row][col].set_yticks([])
```

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### Bacterial Pneumonia

```
n_row = 2
n_col = 4
```

```
fig, ax = plt.subplots(n_row, n_col, figsize = (n_col*3, n_row*3), constrained_layout = True

for row in tqdm(range(n_row)):
    for col in range(n_col):
        ax[row][col].imshow(bacterial[row*n_col + col,:,:,0], cmap = 'bone')
        ax[row][col].set_xticks([])
        ax[row][col].set_yticks([])
```

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### Loading Test Data

#test\_normal = np.load('../input/pneumonia-chest-xray-npy/Array128/Array128/test\_Normal\_128.
#test\_viral = np.load('../input/pneumonia-chest-xray-npy/Array128/Array128/test\_Virus\_128.np
#test\_bacterial = np.load('../input/pneumonia-chest-xray-npy/Array128/Array128/test\_bacteria

```
test_normal = np.load(f'{base_path}/test_Normal_128.npy')
test_viral = np.load(f'{base_path}/test_Virus_128.npy')
test_bacterial = np.load(f'{base_path}/test_bacteria_128.npy')

test_normal.shape, test_viral.shape , test_bacterial.shape

((234, 128, 128, 1), (148, 128, 128, 1), (242, 128, 128, 1))
```

### Create Labels

```
label_test_normal = np.zeros(len(test_normal))
label_test_bacterial = np.ones(len(test_bacterial))
label_test_viral = np.full(len(test_viral),2, dtype = int)

test_data = np.concatenate((test_normal, test_bacterial, test_viral),axis=0)
test_label = np.concatenate((label_test_normal,label_test_bacterial,label_test_viral),axis=0

test_label.shape, test_data.shape

((624,), (624, 128, 128, 1))
```

### Visualization

#### Normal

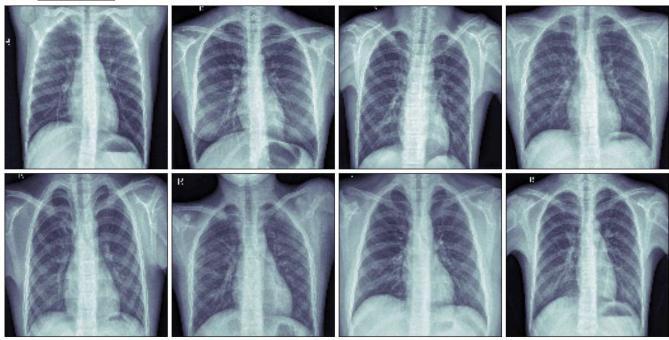
```
n_row = 2
n_col = 4

fig, ax = plt.subplots(n_row, n_col, figsize = (n_col*3, n_row*3), constrained_layout = True

for row in tqdm(range(n_row)):
    for col in range(n_col):
        ax[row][col].imshow(test_normal[row*n_col + col,:,:,0], cmap = 'bone')
        ax[row][col].set_xticks([])
        ax[row][col].set_yticks([])
```



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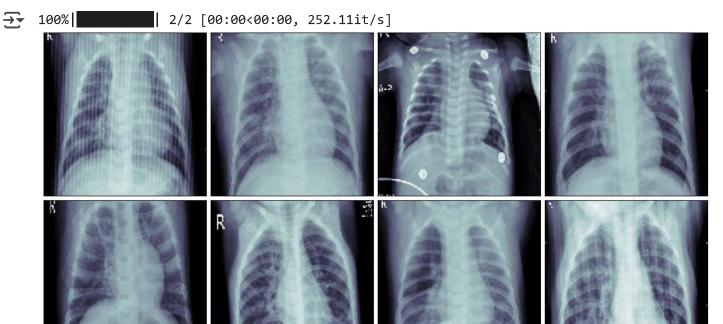


### Viral Pneumonia

```
n_row = 2
n_col = 4

fig, ax = plt.subplots(n_row, n_col, figsize = (n_col*3, n_row*3), constrained_layout = True

for row in tqdm(range(n_row)):
    for col in range(n_col):
        ax[row][col].imshow(test_viral[row*n_col + col,:,:,0], cmap = 'bone')
        ax[row][col].set_xticks([])
        ax[row][col].set_yticks([])
```

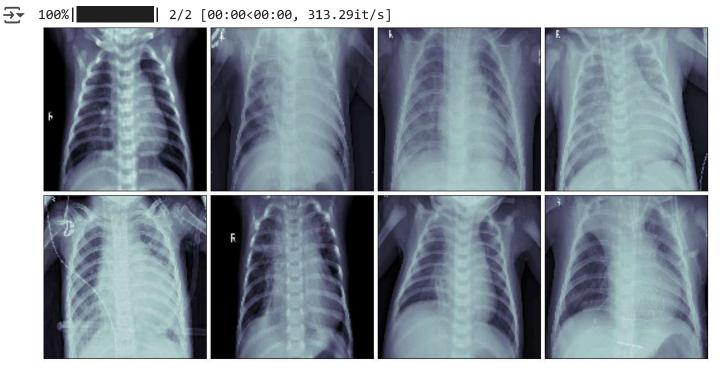


### Bacterial Pneumonia

```
n_row = 2
n_col = 4

fig, ax = plt.subplots(n_row, n_col, figsize = (n_col*3, n_row*3), constrained_layout = True

for row in tqdm(range(n_row)):
    for col in range(n_col):
        ax[row][col].imshow(test_bacterial[row*n_col + col,:,:,0], cmap = 'bone')
        ax[row][col].set_xticks([])
        ax[row][col].set_yticks([])
```



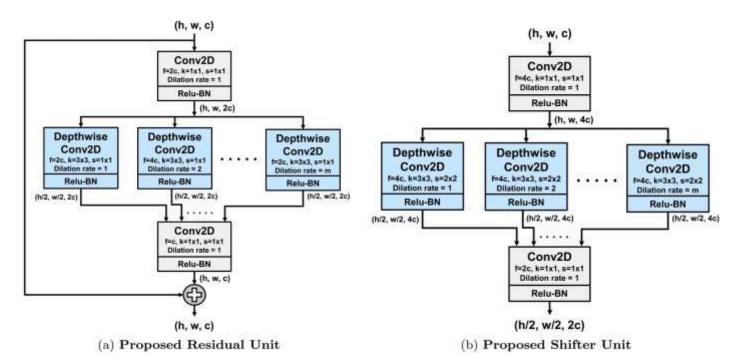
### Class Imbalance

## **CovXNet**

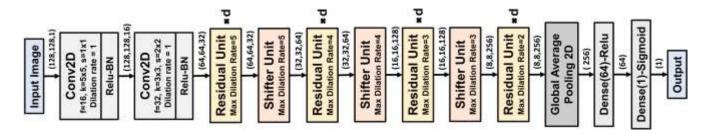
Title: CovXNet: A multi-dilation convolutional neural network for automatic COVID-19 and other pneumonia detection from chest X-ray images with transferable multi-receptive feature optimization

Code: here

### Residual & Shifter Unit:



#### Model:



### CovXNet128

```
# model.summary()
# plot_model(model, show_shapes=True)
```

## One Hot Encoding the labels

```
from tensorflow.keras.utils import to_categorical
train_label = to_categorical(train_label, num_classes= 3)
test_label = to_categorical(test_label, num_classes = 3)
```

### ImageDataGenerator

```
from tensorflow.keras.preprocessing.image import ImageDataGenerator
train_datagen = ImageDataGenerator(rescale = 1/255,
                                   width shift range = 0.1,
                                   height_shift_range = 0.1,
                                   fill_mode = 'constant',
                                   zoom range = 0.2,
                                   rotation_range = 30)
val_datagen = ImageDataGenerator(rescale = 1/255)
train_ds = train_datagen.flow(train_data,
                                     train_label,
                                     batch size = 16,
                                      shuffle = True)
val_ds = val_datagen.flow(test_data,
                                 test_label,
                                 batch size = 16,
                                 shuffle = False)
```

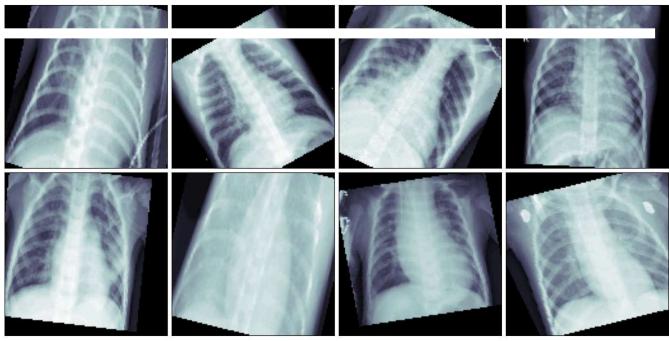
## Vizualization After Augmentation

```
#images, labels = train_ds.next()
images, labels = next(train_ds)

n_row = 2
n_col = 4
```

```
fig, ax = plt.subplots(n_row, n_col, figsize = (n_col*3, n_row*3), constrained_layout = True
for row in tqdm(range(n_row)):
    for col in range(n_col):
        ax[row][col].imshow(images[row*n_col + col,:,:,0], cmap = 'bone')
        ax[row][col].set_xticks([])
        ax[row][col].set_yticks([])
```

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### Callback

```
from tensorflow.keras.callbacks import ModelCheckpoint, CSVLogger
from tensorflow.keras.models import load_model
def get_callbacks():
    filepath = 'covxnet128.h5'
```

# Training

 $\overline{\Sigma}$ 

8/6/25, 11:31 p.m.

```
בטענוו סטווס
                            - 78s 238ms/step - accuracy: 0.8443 - loss: 0.3544 - val a ื
326/326 -
Epoch 67/75
                            - 78s 238ms/step - accuracy: 0.8589 - loss: 0.3434 - val a
326/326 -
Epoch 68/75
326/326 -
                            - 78s 239ms/step - accuracy: 0.8396 - loss: 0.3673 - val_a
Epoch 69/75
326/326 -
                            78s 240ms/step - accuracy: 0.8558 - loss: 0.3432 - val_a
Epoch 70/75
326/326 -
                             82s 240ms/step - accuracy: 0.8454 - loss: 0.3440 - val a
Epoch 71/75
326/326 -
                             78s 239ms/step - accuracy: 0.8613 - loss: 0.3329 - val a
Epoch 72/75
326/326 -
                            - 78s 239ms/step - accuracy: 0.8619 - loss: 0.3231 - val_a
Epoch 73/75
326/326 -
                            - 78s 239ms/step - accuracy: 0.8531 - loss: 0.3445 - val a
Epoch 74/75
326/326 •
                             82s 239ms/step - accuracy: 0.8537 - loss: 0.3387 - val a
Epoch 75/75
326/326 -
                            - 78s 240ms/step - accuracy: 0.8598 - loss: 0.3290 - val a
```

### Plotting History

```
import matplotlib.pyplot as plt
fig, axs = plt.subplots(2, 1, figsize=(10, 10))
# Plot training & validation accuracy values
axs[0].plot(history.history['accuracy'], label='Train Accuracy', color='blue', linestyle='da
axs[0].plot(history.history['val_accuracy'], label='Test Accuracy', color='blue')
axs[0].set_title('Model Accuracy')
axs[0].set_xlabel('Epoch')
axs[0].set_ylabel('Accuracy')
axs[0].legend(loc='lower right')
axs[0].grid(True)
max_val_acc = np.max(history.history['val_accuracy'])
max_val_acc_epoch = np.argmax(history.history['val_accuracy'])
axs[0].annotate(f'Best Acc: {max_val_acc:.2f}',
                xy=(max_val_acc_epoch, max_val_acc),
                xytext=(max_val_acc_epoch*0.8, max_val_acc*0.8),
                arrowprops=dict(facecolor='red', shrink=0.05))
# Plot training & validation loss values
axs[1].plot(history.history['loss'], label='Train Loss', color='orange', linestyle='dashed')
axs[1].plot(history.history['val_loss'], label='Test Loss', color='orange')
axs[1].set_title('Model Loss')
axs[1].set_xlabel('Epoch')
axs[1].set_ylabel('Loss')
```



