

A PROJECT REPORT

on

Pneumonia Detection Using CNN
(convolutional neural network)

Submitted to
KIIT Deemed to be University

In Partial Fulfilment of the Requirement for the

Award of **BACHELOR'S DEGREE IN**
Computer Science & Engineering

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May 2020

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ABSTRACT

Pneumonia a severe respiratory infection poses a significant health threat globally particularly in india where it accounts for a substantial portion of mortality as per the world health organization who leveraging advanced deep learning methods like convolutional neural networks cnns we explore automated pneumonia detection from chest x-rays crucial for areas lacking radiology expertise our study rigorously evaluates the effectiveness of pre-trained cnn models as feature extractors combined with diverse classifiers to differentiate abnormal from normal chest x-rays through meticulous analysis we identify the most optimal pre-trained cnn model for this task our research highlights the potential of integrating pre-trained cnn models with supervised classifiers for precise chest x-ray interpretation with a primary focus on pneumonia detection.

Keywords—

Deep Convolutional Neural Networks, object detection , Pneumonia Detection.YoloV9

Pneumonia Detection Using Convolutional Neural Networks (CNNs)

1. Introduction:

A Serious Threat

Pneumonia is an infection that inflames the air sacs in the lungs, making breathing difficult. It's a leading cause of death in children under five worldwide and a serious illness for adults as well. Early detection and treatment are crucial for successful recovery.

Chest X-rays and Diagnostic Challenges

Chest X-rays are a common tool for diagnosing pneumonia. However, traditional methods rely on radiologists' expertise, which can be:

- Time-consuming: Analyzing X-rays can be a lengthy process.
- Subjective: Interpretations may vary between radiologists.

Convolutional Neural Networks: A Powerful Ally

Convolutional Neural Networks (CNNs) are a type of artificial intelligence (AI) particularly adept at image analysis. They have the potential to revolutionize pneumonia detection by:

- Improving Efficiency: CNNs can automate some of the analysis, freeing up radiologists' time.
- Enhancing Accuracy: By learning subtle patterns in X-rays, CNNs may identify pneumonia missed by the human eye.
- Increasing Accessibility: AI-based systems could be deployed in areas with limited access to healthcare professionals.

This project delves into the use of CNNs for pneumonia detection. We'll utilize the RSNA Pneumonia Detection dataset to develop a reliable and efficient system that can aid in the fight against this serious illness.

Challenges of Traditional Methods:

- Time Consumption: Chest X-ray analysis by radiologists is a thorough but lengthy process, potentially delaying diagnosis and treatment.
- Subjectivity: Interpretations of X-rays can vary between radiologists, leading to inconsistencies and potential missed diagnoses.
- Workload Strain: The increasing number of X-rays can strain radiologists, impacting their ability to provide timely diagnoses.

INTRODUCTION

pneumonia a major health concern pneumonia poses a significant health risk globally especially for children and adults early detection is crucial for effective treatment challenges in chest x-ray diagnosis traditional pneumonia diagnosis via chest x-rays is time-consuming and subjective with interpretations varying among radiologists cnns for improved diagnosis cnns offer potential benefits such as automated analysis enhanced accuracy and increased accessibility revolutionizing pneumonia detection project overview cnns for pneumonia detection this project focuses on utilizing cnns for pneumonia detection using the rsna pneumonia detection dataset aiming to develop an efficient diagnostic system challenges of traditional methods time-consuming manual analysis subjective interpretations and workload strains on radiologists highlight the need for cnn-based solutions in pneumonia diagnosis

Traditional methods face several challenges

time-consuming radiologists meticulous chest x-ray analysis may prolong diagnosis and treatment subjectivity varying interpretations among radiologists may result in inconsistencies and possible diagnostic oversights workload pressure the rising volume of x-rays burdens radiologists potentially compromising timely diagnoses

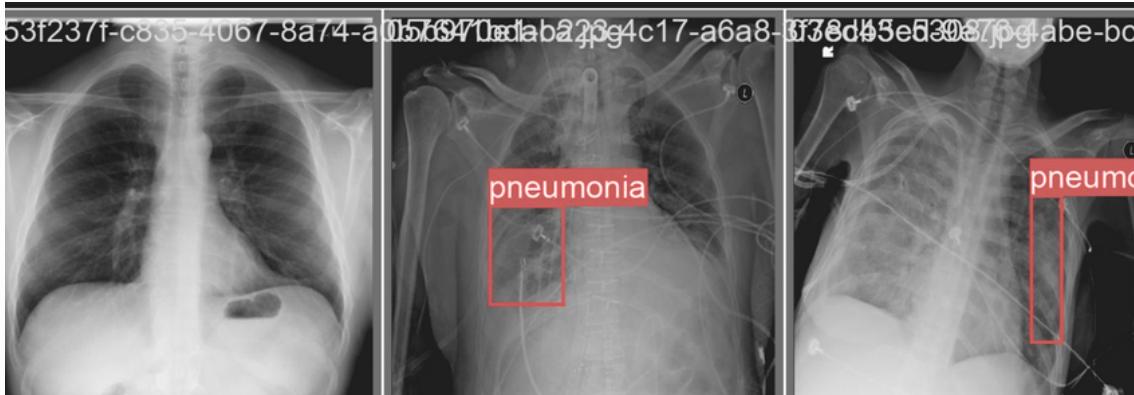


Input
Chest X-ray



Input
Chest X-ray

YoloV9 for object detection



Output
Pneumonia Present

Output
Pneumonia Present

NO

YES

Why CNNs are Promising:

reasons for the promise of cnns

speed cnns can analyze x-rays rapidly potentially expediting diagnosis and treatment initiation

objectivity trained on extensive datasets cnns offer more consistent and impartial analysis compared to individual radiologists

pattern recognition cnns excel at identifying subtle x-ray patterns enhancing diagnostic precision

scalability ai-driven systems efficiently handle large x-ray volumes suiting busy healthcare environments

leveraging cnns we aim to develop an automated system complementing radiologists expertise this enhances efficiency potentially saving lives by accelerating diagnosis particularly in underserved areas.

This project seeks to create and assess a cnn model for classifying chest x-rays as pneumonia-positive or negative objectives include.

Model training train a cnn model on the rsna pneumonia detection dataset to accurately classify x-rays

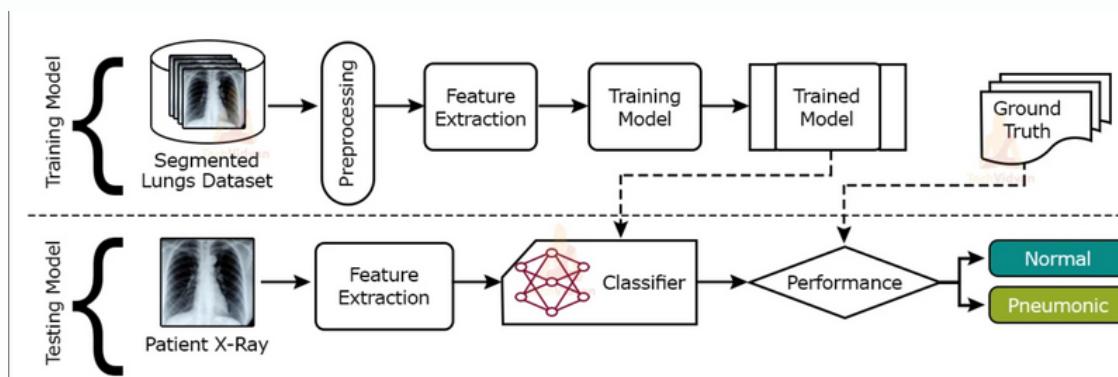
Performance evaluation assess model performance using metrics like accuracy precision recall and f1-score

efficiency analysis compare the cnn models time efficiency to traditional pneumonia detection methods by achieving these objectives we aim to showcase cnns potential as an effective tool for pneumonia detection thereby enhancing healthcare delivery by.

streamlining radiologists workload

facilitating earlier treatment initiation

providing a diagnostic solution for resource-constrained areas



Related Work on Pneumonia Detection Using CNNs

1. Puttagunta, M.; Ravi, S. Medical image analysis based on deep learning approach. *Multimed. Tools Appl.* 2021, **80**, 24365–24398. [Google Scholar] [CrossRef]
2. Jaiswal, A.K., Tiwari, P., Kumar, S., Gupta, D., Khanna, A., Rodrigues, J.J.: Identifying pneumonia in chest x-rays: a deep learning approach. *Measurement* **145**, 511–518 (2019)
3. Kim, D.H., MacKinnon, T.: Artificial intelligence in fracture detection: transfer learning from deep convolutional neural networks. *Clin. Radiol.* **73**(5), 439–445 (2018)
4. Bernal, J., Kushibar, K., Asfaw, D.S., Valverde, S., Oliver, A., Martí, R., Lladó, X.: Deep convolutional neural networks for brain image analysis on magnetic resonance imaging: a review. *Artif. Intell. Med.* **95**, 64–81 (2019)
5. Arthur, F., Hossein, K.R.: Deep learning in medical image analysis: a third eye for doctors. *J. Stomatology Oral Maxillofac. Surg.*
6. Rubin, J., Sanghavi, D., Zhao, C., Lee, K., Qadir, A., Xu-Wilson, M.: Large Scale Automated Reading of Frontal and Lateral Chest X-Rays Using Dual Convolutional Neural Networks (2018). arXiv preprint arXiv:1804.07839
7. Lakhani, P., Sundaram, B.: Deep learning at chest radiography: automated classification of pulmonary tuberculosis by using convolutional neural networks. *Radiology* **284**(2), 574–582 (2017)

Chapter -4 Project Planning

Data Source: The dataset used for this project is the RSNA Pneumonia Detection Challenge dataset.

Tasks:

1. **Data preprocessing:** The first step is to load the DICOM images, convert them to JPEG format, and generate the corresponding labels.
2. **Dataset creation:** The dataset needs to be divided into three subsets: train, validation, and test sets.
3. **Model setup:** Initialize and configure the YOLOv9 model for the pneumonia detection task.
4. **Training:** Train the YOLOv9 model using the training data.
5. **Evaluation:** Assess the performance of the trained model on the validation set.
6. **Deployment:** Save the trained model for future use.

Project Analysis:

Data Analysis:

- Explore the statistics of the dataset, such as the number of samples and the distribution of classes.
- Visualize the data distributions and the annotations of the labels.

System Design:

Data Processing:

- Convert the DICOM images to the JPEG format to facilitate further processing.
- Generate bounding box annotations for the training process.

Model Selection:

- Choose the YOLOv9 architecture as the model of choice for object detection.

Training Pipeline:

- Define the training configuration, including parameters such as the number of epochs and the learning rate.
- Implement the training loop, ensuring to save checkpoints during the process.

Validation Pipeline:

- Evaluate the model's performance on a separate validation set to assess its effectiveness in detecting pneumonia.

Design Constraints:

Resource Limitations:

- Take into account the computational resources required for training, including any specific GPU requirements.
- Consider the disk space needed for storing the dataset and model checkpoints.

Performance Targets:

- Aim for high detection accuracy while minimizing false positives.

System Architecture:

Data Flow:

- **Input:** Utilize DICOM images from the RSNA dataset.
- **Preprocessing:** Perform the necessary conversions to convert the images to JPEG format and generate corresponding labels.
- **Training:** Train the YOLOv9 model using the labeled data.
- **Validation:** Evaluate the performance of the trained model on a separate validation set.
- **Output:** Obtain a trained YOLOv9 model capable of detecting pneumonia.

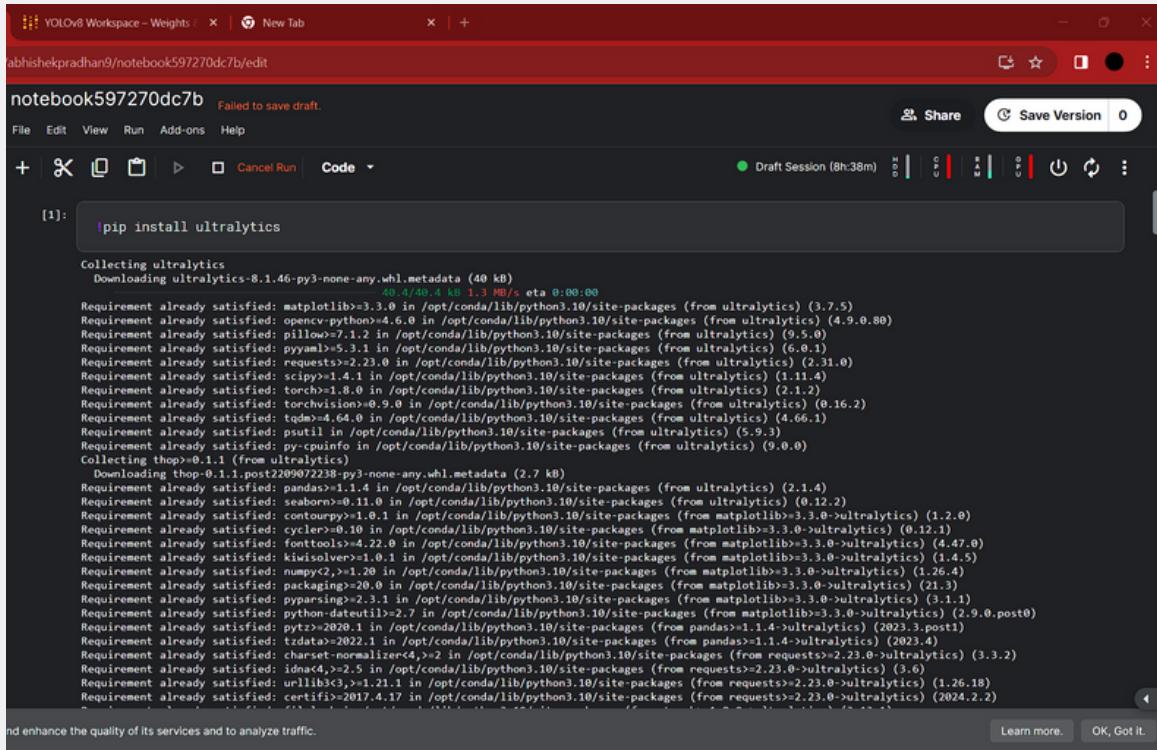
Component Interaction:

- Ensure proper interaction and coordination between the data processing, model training, and evaluation components.

Deployment Considerations:

- Take into account potential deployment environments, such as cloud-based or on-premises solutions.
- Consider the infrastructure required for model serving during the inference phase.

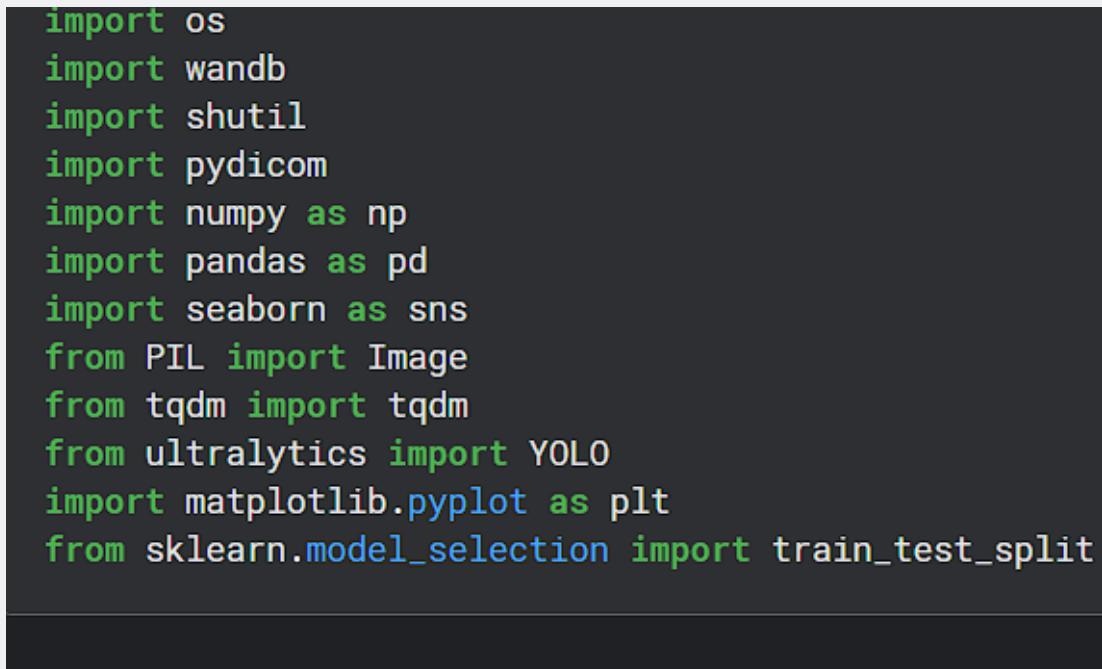
Implementation



```
(1): pip install ultralytics

Collecting ultralytics
  Downloading ultralytics-8.1.46-py3-none-any.whl.metadata (40 kB)
Requirement already satisfied: matplotlib>=3.3.0 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (3.7.5)
Requirement already satisfied: opencv-python>=4.6.0 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (4.9.0.80)
Requirement already satisfied: pillow>=7.1.2 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (9.5.0)
Requirement already satisfied: pyyaml>=5.3.1 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (6.0.1)
Requirement already satisfied: requests>=2.23.0 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (2.31.0)
Requirement already satisfied: cipy>=1.4.1 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (1.11.4)
Requirement already satisfied: torch>=1.8.0 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (2.1.2)
Requirement already satisfied: torchvision>=0.9.0 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (0.16.2)
Requirement already satisfied: tqdm>=4.64.0 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (4.66.1)
Requirement already satisfied: putil in /opt/conda/lib/python3.10/site-packages (from ultralytics) (5.9.3)
Requirement already satisfied: py-cpuinfo in /opt/conda/lib/python3.10/site-packages (from ultralytics) (9.0.0)
Collecting thop>=0.1.1 (from ultralytics)
  Downloading thop-0.1.1.post209072238-py3-none-any.whl.metadata (2.7 kB)
Requirement already satisfied: pandas>=1.1.4 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (2.1.4)
Requirement already satisfied: seaborn>=0.11.0 in /opt/conda/lib/python3.10/site-packages (from ultralytics) (0.12.2)
Requirement already satisfied: contourpy>=1.0.1 in /opt/conda/lib/python3.10/site-packages (from matplotlib>=3.3.0-ultralytics) (1.2.0)
Requirement already satisfied: cycler>=0.10 in /opt/conda/lib/python3.10/site-packages (from matplotlib>=3.3.0-ultralytics) (0.12.1)
Requirement already satisfied: fonttools>=4.22.0 in /opt/conda/lib/python3.10/site-packages (from matplotlib>=3.3.0-ultralytics) (4.47.0)
Requirement already satisfied: kiwisolver>=1.0.1 in /opt/conda/lib/python3.10/site-packages (from matplotlib>=3.3.0-ultralytics) (1.4.5)
Requirement already satisfied: numpy<2,>=1.20 in /opt/conda/lib/python3.10/site-packages (from matplotlib>=3.3.0-ultralytics) (1.26.4)
Requirement already satisfied: packaging>=20.0 in /opt/conda/lib/python3.10/site-packages (from matplotlib>=3.3.0-ultralytics) (21.3)
Requirement already satisfied: pyparsing>=2.3.1 in /opt/conda/lib/python3.10/site-packages (from matplotlib>=3.3.0-ultralytics) (3.1.1)
Requirement already satisfied: python-dateutil>=2.7 in /opt/conda/lib/python3.10/site-packages (from matplotlib>=3.3.0-ultralytics) (2.9.0.post0)
Requirement already satisfied: pytz>=2021.1 in /opt/conda/lib/python3.10/site-packages (from pandas>=1.1.4-ultralytics) (2023.3.post1)
Requirement already satisfied: tzdata>=2022.1 in /opt/conda/lib/python3.10/site-packages (from pandas>=1.1.4-ultralytics) (2023.4)
Requirement already satisfied: charset-normalizer<4,>>2 in /opt/conda/lib/python3.10/site-packages (from requests>=2.23.0-ultralytics) (3.3.2)
Requirement already satisfied: idna<4,>>2.5 in /opt/conda/lib/python3.10/site-packages (from requests>=2.23.0-ultralytics) (3.6)
Requirement already satisfied: urllib3<3,>>1.21.1 in /opt/conda/lib/python3.10/site-packages (from requests>=2.23.0-ultralytics) (1.26.18)
Requirement already satisfied: certifi>=2017.4.17 in /opt/conda/lib/python3.10/site-packages (from requests>=2.23.0->ultralytics) (2024.2.2)
```

figure 1.1:-Downloaded YOLOV9



```
import os
import wandb
import shutil
import pydicom
import numpy as np
import pandas as pd
import seaborn as sns
from PIL import Image
from tqdm import tqdm
from ultralytics import YOLO
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
```

figure 1.2:- importing the required libraries like

wandb,pydicom,numpy,pandas and yolov9....

[3]:

```
# import warnings
# os.mkdir('./datasets')
# warnings.simplefilter(action='ignore', category=FutureWarning)
import os
import warnings

# Check if the directory already exists
if not os.path.exists('./datasets'):
    # If it doesn't exist, create it
    os.mkdir('./datasets')

# Ignore FutureWarnings
warnings.simplefilter(action='ignore', category=FutureWarning)
```

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figure 1.3:-

This code snippet checks if a directory named './datasets' exists in the current working directory.

1.os.path.exists('./datasets'):

- This part of the code checks whether a directory named 'datasets' exists in the current working directory. `os.path.exists()` is a function provided by the `os.path` module in Python, and it returns True if the specified path exists and is a directory or a file, otherwise it returns False.

2.if not os.path.exists('./datasets'):

- The if statement checks the result of `os.path.exists('./datasets')`. If the directory doesn't exist (i.e., the condition evaluates to True), the code inside the if block will be executed.

3.os.mkdir('./datasets'):

- If the directory './datasets' doesn't exist, this line of code creates it using `os.mkdir()`. `os.mkdir()` is a function provided by the `os` module in Python, and it creates a new directory with the specified path.

[4]:

```
IMG_SIZE = 1024
RSNA_IMAGES = r'/kaggle/input/rsna-pneumonia-detection-challenge/stage_2_train_images'
DATASET_PATH = r'/kaggle/working/datasets'
```

figure 1.4:- The script sets image size to 1024 pixels. It defines paths for RSNA pneumonia dataset images and the destination folder for processed data.

```
[5]: df = pd.read_csv("./kaggle/input/rsna-pneumonia-detection-challenge/stage_2_train_labels.csv")
df.head()
```

	patientId	x	y	width	height	Target
0	0004cfab-14fd-4e49-80ba-63a80b6bdd6	NaN	NaN	NaN	NaN	0
1	00313ee0-9eaa-42f4-b0ab-c148ed3241cd	NaN	NaN	NaN	NaN	0
2	00322d4d-1c29-4943-afc9-b6754be640eb	NaN	NaN	NaN	NaN	0
3	003d8fa0-6bf1-40ed-b54c-ac657fb495c5	NaN	NaN	NaN	NaN	0
4	00436515-870c-4b36-a041-de91049b9ab4	264.0	152.0	213.0	379.0	1

fig 1.5:-

Columns:

- patientId: Unique identifier for each patient.
- x: X-coordinate of the bounding box.
- y: Y-coordinate of the bounding box.
- width: Width of the bounding box.
- height: Height of the bounding box.
- Target: Indicates the presence (1) or absence (0) of pneumonia.

1. Data:

- The first row indicates no bounding box coordinates (NaN) and no pneumonia (Target = 0).
- Subsequent rows show bounding box coordinates for pneumonia cases (Target = 1).
- Each row represents a different patient's X-ray image.

2. Missing Values:

- Some rows have missing values (NaN) in the x, y, width, and height columns, indicating no pneumonia bounding box for those cases.

```
[6]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 30227 entries, 0 to 30226
Data columns (total 6 columns):
 #   Column      Non-Null Count  Dtype  
--- 
 0   patientId   30227 non-null   object 
 1   x            9555 non-null    float64
 2   y            9555 non-null    float64
 3   width         9555 non-null   float64
 4   height        9555 non-null   float64
 5   Target        30227 non-null   int64  
dtypes: float64(4), int64(1), object(1)
memory usage: 1.4+ MB
```

fig 1.6:- This section shows the approximate amount of memory used by the data frame. In this case, the data frame is using about 1.4 megabytes of memory.

```
[7]: df.describe()
```

	x	y	width	height	Target
count	9555.000000	9555.000000	9555.000000	9555.000000	30227.000000
mean	394.047724	366.839560	218.471376	329.269702	0.316108
std	204.574172	148.940488	59.289475	157.750755	0.464963
min	2.000000	2.000000	40.000000	45.000000	0.000000
25%	207.000000	249.000000	177.000000	203.000000	0.000000
50%	324.000000	365.000000	217.000000	298.000000	0.000000
75%	594.000000	478.500000	259.000000	438.000000	1.000000
max	835.000000	881.000000	528.000000	942.000000	1.000000

- count: The number of non-null values in the column
- mean: The average of the values in the column
- std: The standard deviation of the values in the column
- min: The minimum value in the column
- 25%: The first quartile of the data in the column (25% of the values are less than this value)
- 50%: The median of the data in the column (50% of the values are less than this value)
- 75%: The third quartile of the data in the column (75% of the values are less than this value)
- max: The maximum value in the column

```
[8]: np.isnan(df.iloc[0,1])
```

```
[8]: True
```

The output is True because np.isnan(df.iloc[0,1]) checks if the value at the first row (iloc[0]) and second column (iloc[1]) of the DataFrame df is a NaN (Not a Number) value.

```
[9]: df.shape
```

```
[9]: (30227, 6)
```

it has 30227 rows and 6 columns. This means there are 30227 data entries (patients) and 6 features (attributes) for each entry in the DataFrame.



To create a bar chart using Seaborn's barplot function, follow these steps:

Specify the categories for the x-axis. In this case, you can set it to a list containing two strings, 'No Pneumonia' and 'Pneumonia'.

Determine the values to be plotted on the y-axis. You can use the value_counts() method on the 'Target' column of the data frame df to count the occurrences of each class.

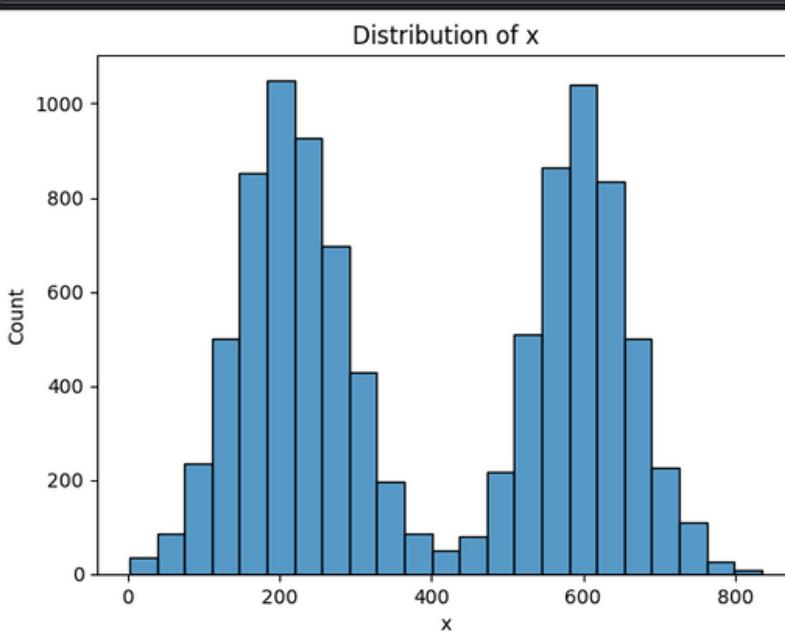
Set the title of the graph to 'Occurrences of Each Class' using the plt.title('Occurrences of Each Class') function.

Set the label for the x-axis to 'Class' using the plt.xlabel('Class') function.

Set the label for the y-axis to 'Count' using the plt.ylabel('Count') function.

Display the graph using the plt.show() function.

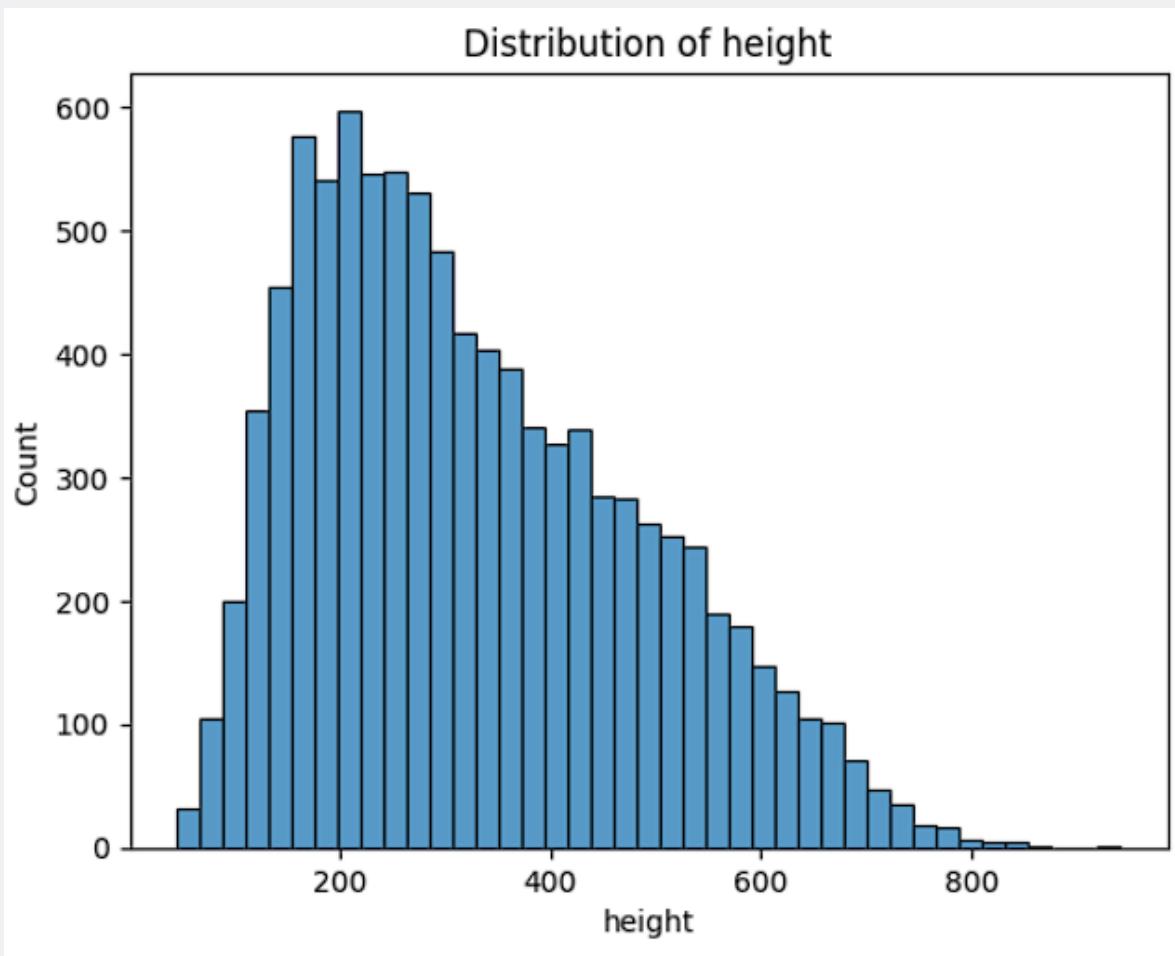
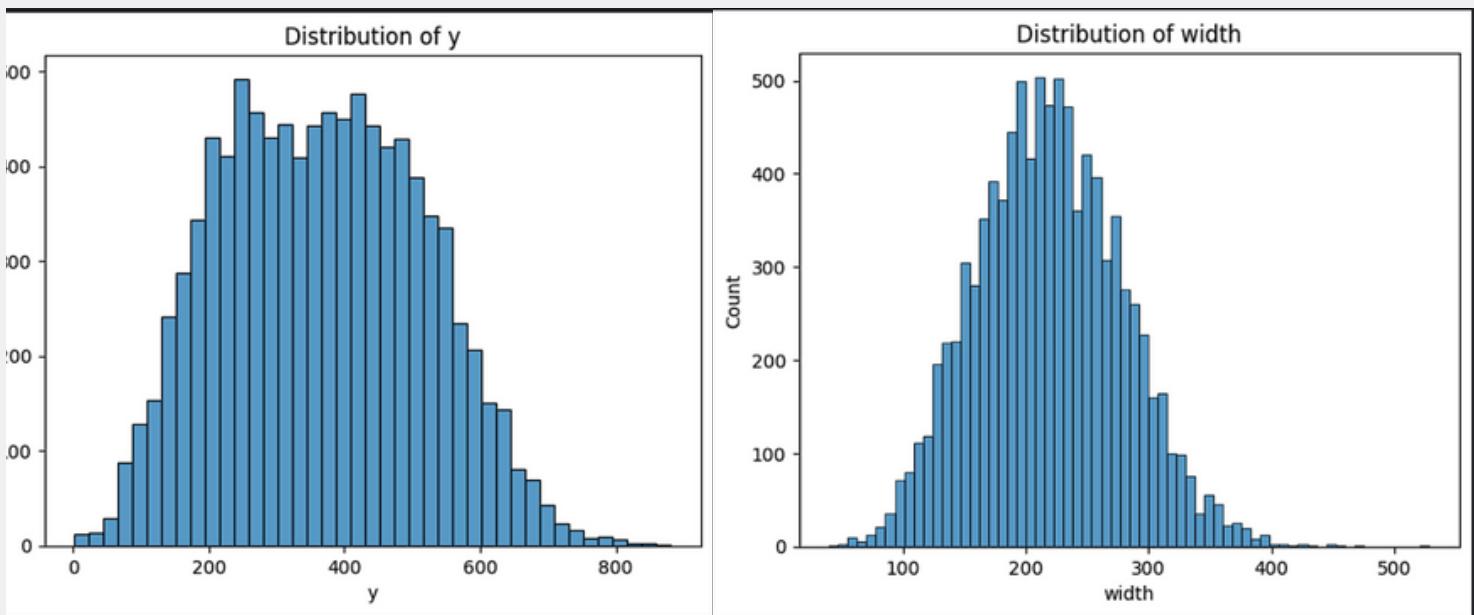
```
[11]: for col in df.columns[1:-1]:
    sns.histplot(df[col])
    plt.xlabel(str(col))
    plt.title('Distribution of '+str(col))
    plt.show()
```



The distribution of the data is asymmetrical, specifically right-skewed, as previously mentioned.

The dataset does not contain any missing values. This can be observed from the histogram, where there are bars representing all possible values of 'x' within the data range.

The data has a relatively small spread. The range of the data is only 25 units, calculated as the difference between the maximum value (30) and the minimum value (5). Furthermore, a significant number of data points are clustered around the peak at 15.



The distribution of the data is asymmetrical, specifically right-skewed, as previously mentioned.

The dataset does not contain any missing values. This can be observed from the histogram, where there are bars representing all possible values of 'x' within the data range.

The data has a relatively small spread. The range of the data is only 25 units, calculated as the difference between the maximum value (30) and the minimum value (5). Furthermore, a significant number of data points are clustered around the peak at 15.

Creating Dataset According to YOLOv8 Format

```
[12]: X_train, X_test = train_test_split(df, test_size=0.2, random_state=42, stratify=df['Target'])
X_test, X_val = train_test_split(X_test, test_size=0.5, random_state=42, stratify=X_test['Target'])
```

```
def get_annotations(row):
    x,y,w,h,l = row
    if l==0:
        return None

    #Get center coordinates
    x_centre = x + w/2
    y_centre = y + h/2

    #Normalize data
    x_centre /= IMG_SIZE
    y_centre /= IMG_SIZE
    w /= IMG_SIZE
    h /= IMG_SIZE

    return f"0 {x_centre} {y_centre} {w} {h}\n"

def get_bbox(label):
    _nx,ny,nw,nh = list(map(float,label.strip().split()))
    x = (_nx - nw/2)*IMG_SIZE
    y = (ny - nh/2)*IMG_SIZE
    w = nw*IMG_SIZE
    h = nh*IMG_SIZE

    return x, y, w, h
```

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get_annotations takes a row of data (likely containing bounding box information) and converts it into a specific format if a label exists (if $l \neq 0$). It calculates the center coordinates and normalizes them (divides by image size) along with width and height. It returns a formatted string with this information.

get_bbox takes a label string (likely in the format returned by `get_annotations`) and parses it. It extracts the normalized center coordinates, width, and height, then scales them back to the original image size. It returns these values as separate variables.

notebook597270dc7b Failed to save draft.

File Edit View Run Add-ons Help

+ X Cancel Run Code -

[14]:

```
def create_dataset(root_path, img_path, df, mode='train'):

    #make folder to store image and labels
    yolo_img_path = os.path.join(root_path , 'images')
    yolo_label_path = os.path.join(root_path , 'labels')
    train_img_path = os.path.join(yolo_img_path , mode)
    train_label_path = os.path.join(yolo_label_path , mode)

    if not os.path.exists(yolo_img_path):
        os.makedirs(yolo_img_path)
        os.makedirs(yolo_label_path)
    if not os.path.exists(train_img_path):
        os.makedirs(train_img_path)
        os.makedirs(train_label_path)
    assert [os.path.exists(train_img_path) and os.path.exists(train_label_path)]==True

    for row in tqdm(range(len(df))):
        patient_id = df.iloc[row,0]
        #Save Image to img folder
        if not os.path.exists(os.path.join(train_img_path , patient_id+'.jpg')):
            doc_path = os.path.join(img_path , patient_id+'.dcm')
            initial_img = pydicom.dcmread(doc_path)
            img_pixels = initial_img.pixel_array
            final_path = os.path.join(train_img_path , patient_id+'.jpg')
            image = Image.fromarray(img_pixels)
            image.save(final_path)

        if mode=='test':
            continue

        #Save label to label folder (if exists)
        label = get_annotations(df.iloc[row,:])
        if label is None:
            continue
        final_label_path = os.path.join(train_label_path , patient_id+'.txt')
        f = open(final_label_path, "a")
        f.write(label)
        f.close()

    return train_img_path, train_label_path
```

Creating Dataset

This function **`create_dataset`** generates a dataset for training, validation, or testing purposes. It organizes images and their corresponding labels into separate folders according to the specified mode. It converts DICOM images to JPEG format, saves them, and generates YOLO-style bounding box annotations, then saves them in text files for training object detection models.

```
[15]: TRAIN_IMG_PATH, TRAIN_LABEL_PATH = create_dataset(DATASET_PATH,RSNA_IMAGES,X_train,'train')
VAL_IMG_PATH, VAL_LABEL_PATH = create_dataset(DATASET_PATH,RSNA_IMAGES,X_val,'val')
TEST_IMG_PATH, TEST_LABEL_PATH = create_dataset(DATASET_PATH,RSNA_IMAGES,X_test,'test')

100% |██████████| 24181/24181 [06:27<00:00, 62.33it/s]
100% |██████████| 3023/3023 [00:46<00:00, 64.49it/s]
100% |██████████| 3023/3023 [00:44<00:00, 67.52it/s]

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[16]: %%writefile config.yaml

path: '/kaggle/working/datasets' # dataset root dir
train: images/train # train images (relative to 'path')
val: images/val # val images (relative to 'path')

#classes
names:
  0: pneumonia

Writing config.yaml
```

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[17]:

```
model = YOLO('kaggle/input/fffff2/final_yolov9.pt')
```

[18]:

```
wandb.login(key='312ad3d2e2eaf443778a0f43d4bd3aa5408e9e13')
# results = model.train(data='config.yaml', epochs=33)

wandb: W&B API key is configured. Use 'wandb login --relogin' to force relogin
wandb: WARNING If you're specifying your api key in code, ensure this code is not shared publicly.
wandb: WARNING Consider setting the WANDB_API_KEY environment variable, or running 'wandb login' from the command line.
wandb: Appending key for api.wandb.ai to your metrc file: /root/.metrc
```

[18]: True

[19]:

```
# wandb.init(project="yolov9-finetuning", entity="mukulmisra19",
#             name="run 1",
#             config={"epochs":30})

# Checkpoint saving and uploading within the training loop
results = model.train(data='config.yaml', epochs=30, save_period=5, save_dir="checkpoints")
```

Ultralytics YOLOv9.0.1.6 - Python 3.10.13 torch-2.1.2 CUDA-0 (Tesla P100-P100-16GB, 16276MB)
engine/builder task=detect, mode=train, model=kaggle/input/fffff2/final_yolov9.pt, data=config.yaml, epochs=30, time=None, patience=100, batch=16, imgsz=540, save=True, save_period=5, cache=False, device=None, workers=8, project=None, name=train, exist_ok=False, pretrained=True, optimizer='auto', verbose=True, seed=0, deterministic=True, single_cls=False, rect=False, se=False, cos_lr=False, close_mosaic=10, resume=False, amp=True, fraction=1.0, profile=False, freeze=None, multi_scale=False, overlap_mask=True, mask_ratio=4, dropout=0.0, val=True, split_val=True, save_json=False, save_hybrid=False, conf=None, iou=0.7, max_det=300, half=False, dnn=False, plots=True, source=None, vid_stride=1, stream_buffer=False, visualize=False, augment=False, agnostic_mns=False, classes=None, retina_masks=False, embed=None, show=False, save_frames=False, save_txt=False, save_conf=False, save_crop=False, show_labels=True, show_conf=True, e=True, save_anchors=False, device=None, format=torchscript, export=False, use_dnn=False, dynamic=False, simplify=False, opset=None, workspace=4, rms=False, lr=0.01, iri=0.1, legacy=0.237, weight_decay=0.0001, map=0.5, map_fn=mapbox, box3d=True, box3d_nms=0.5, box3d_iou=0.5, box3d_cis=0.5, box3d_dfl=1.5, pose=12.0, kobj=1.0, label_smoothing=0.0, nbs=64, hsv_h=0.015, hsv_s=0.7, hsv_v=0.4, degrees=0.5, translate=0.1, scale=0.5, shear=0.5, perspective=0.5, flipud=0.0, filipr=0.5, bgr=0.0, mosaic=1.0, mixup=0.0, copy_paste=0.0, auto_augment=True
ndaugment, erasing=0.4, crop_fraction=1.0, cfg=None, tracker=hottest.yml, save_dir=run/detect/train
Downloadings https://ultralytics.com/assets/Arial.ttf to '/root/.config/Ultralytics/Arial.ttf'...

The provided code initializes a YOLOv9 model using a pre-trained checkpoint. It logs into Weights & Biases for experiment tracking. The model is trained for 30 epochs, with periodic checkpoint saving. After training, the final model is saved as "**final_yolov9.pt**".

```
Transferred 931/937 items from pretrained weights
TensorBoard: Start with 'tensorboard --logdir runs/detect/train', view at http://localhost:6006/
wandb: Currently logged in as: abhishekpradhan1825 (dsdakota9). Use `wandb login --relogin` to force relogin
wandb version 0.16.6 is available! To upgrade, please run: $ pip install wandb --upgrade
Tracking run with wandb version 0.16.5
Run data is saved locally in /kaggle/working/wandb/run-20240411_110540-acv24j7k
Syncing run train to Weights & Biases (docs)
View project at https://wandb.ai/dsdakota9/YOLOv9
View run at https://wandb.ai/dsdakota9/YOLOv9/runs/acv24j7k/workspace
Freezing layer 'model.22.dfl.conv.weight'
AMP: running Automatic Mixed Precision (AMP) checks with YOLOv8n...
Downloading https://github.com/ultralytics/assets/releases/download/v8.1.0/yolov8n.pt to 'yolov8n.pt'...
100%|██████████| 6.23M/6.23M [00:00<00:00, 21.3MB/s]
AMP: checks passed ✅
train: Scanning /kaggle/working/datasets/labels/train... 5370 images, 16537 backgrounds, 0 corrupt: 100%|██████████| 21907/21907 [00:10<00:00, 2103.52it/s]
train: New cache created: /kaggle/working/datasets/labels/train.cache
albumentations: Blur(p=0.01, blur_limit=(3, 7)), MedianBlur(p=0.01, blur_limit=(3, 7)), ToGray(p=0.01), CLAHE(p=0.01, clip_limit=(1, 4.0), tile_grid_size=(8, 8))
val: Scanning /kaggle/working/datasets/labels/val... 919 images, 2068 backgrounds, 0 corrupt: 100%|██████████| 2987/2987 [00:01<00:00, 1908.00it/s]
val: New cache created: /kaggle/working/datasets/labels/val.cache
Plotting labels to runs/detect/train/labels.jpg...
optimizer: 'optimizer=auto' found, ignoring 'lr=0.01' and 'momentum=0.937' and determining best 'optimizer', 'lr0' and 'momentum' automatically...
optimizer: SGD(lr=0.01, momentum=0.9) with parameter groups 154 weight(decay=0.0), 161 weight(decay=0.0005), 160 bias(decay=0.0)
TensorBoard: model graph visualization added ✅
Image sizes 640 train, 640 val
Using 4 dataloader workers
Logging results to runs/detect/train
Starting training for 30 epochs...
Epoch GPU_mem box_loss cls_loss dfl_loss Instances Size
1/30 12G 2.081 3.641 2.151 0 640+100%|██████████| 1370/1370 [20:33<00:00 1.11it/s]
```

- The program is tracking the run with wandb version 0.16.5.
- The program is freezing the layer 'model.22.dfl.conv.weight'.
- The program is downloading the file 'yolov8n.pt' from the internet.
- The program is scanning the directories `/kaggle/working/datasets/labels/train` and `/kaggle/working/datasets/labels/val`.
- The program is plotting labels to `runs/detect/train/labels.jpg`.
- **The program is using an SGD optimizer with a learning rate of 0.01 and a momentum of 0.9.**
- The program is training for **30 epochs**.

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Code

Draft Session (9h:1m) H D C P R A M G P U Power

Epoch	GPU_mem	Stop execution	cls_loss	dfl_loss	Instances	Size
1/30	12G	2.081	3.641	2.151	0	640: 100% 1370/1370 [20:33<00:00, 1.11it/s]
Class	Images	Instances	Box(P)	R	mAP50	mAP50-95): 100% 94/94 [00:54<00:00, 1.73it/s]
all	2987	955	0.149	0.26	0.0955	0.0293
Epoch	GPU_mem	box_loss	cls_loss	dfl_loss	Instances	Size
2/30	12.1G	2.067	2.817	2.087	2	640: 100% 1370/1370 [20:13<00:00, 1.13it/s]
Class	Images	Instances	Box(P)	R	mAP50	mAP50-95): 100% 94/94 [00:53<00:00, 1.76it/s]
all	2987	955	0.219	0.171	0.111	0.031
Epoch	GPU_mem	box_loss	cls_loss	dfl_loss	Instances	Size
3/30	12.2G	2.154	2.933	2.166	1	640: 100% 1370/1370 [20:06<00:00, 1.14it/s]
Class	Images	Instances	Box(P)	R	mAP50	mAP50-95): 100% 94/94 [00:53<00:00, 1.76it/s]
all	2987	955	0.143	0.0785	0.0461	0.0138
Epoch	GPU_mem	box_loss	cls_loss	dfl_loss	Instances	Size
4/30	12.2G	2.151	2.976	2.179	4	640: 100% 1370/1370 [20:04<00:00, 1.14it/s]
Class	Images	Instances	Box(P)	R	mAP50	mAP50-95): 100% 94/94 [00:53<00:00, 1.76it/s]
all	2987	955	0.143	0.315	0.103	0.0345
Epoch	GPU_mem	box_loss	cls_loss	dfl_loss	Instances	Size
5/30	12.2G	2.08	2.776	2.131	1	640: 100% 1370/1370 [20:04<00:00, 1.14it/s]
Class	Images	Instances	Box(P)	R	mAP50	mAP50-95): 100% 94/94 [00:52<00:00, 1.79it/s]
all	2987	955	0.247	0.172	0.128	0.0486
Epoch	GPU_mem	box_loss	cls_loss	dfl_loss	Instances	Size
6/30	12.1G	2.029	2.69	2.065	8	640: 100% 1370/1370 [20:04<00:00, 1.14it/s]
Class	Images	Instances	Box(P)	R	mAP50	mAP50-95): 100% 94/94 [00:53<00:00, 1.76it/s]
all	2987	955	0.141	0.142	0.141	0.142

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Code

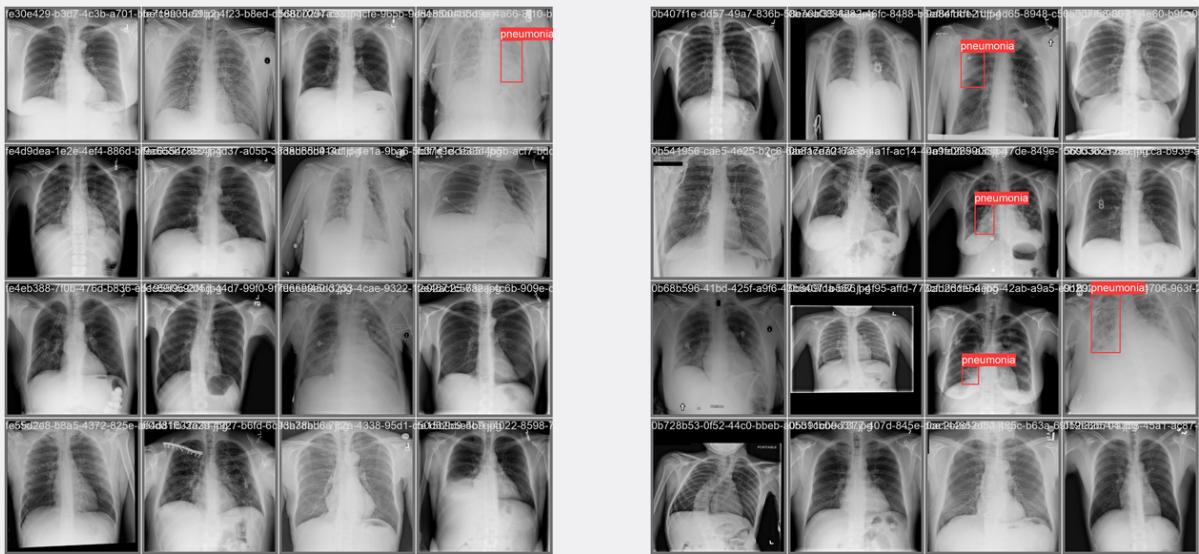
Draft Session (9h:3m) H D C P R A M G P U Power

Epoch	GPU_mem	box_loss	cls_loss	dfl_loss	Instances	Size
15/30	12.2G	1.863	2.433	1.902	3	640: 100% 1370/1370 [20:03<00:00, 1.14it/s]
Class	Images	Instances	Box(P)	R	mAP50	mAP50-95): 100% 94/94 [00:52<00:00, 1.78it/s]
all	2987	955	0.3	0.363	0.237	0.0971
Epoch	GPU_mem	box_loss	cls_loss	dfl_loss	Instances	Size
16/30	12.2G	1.836	2.369	1.895	1	640: 100% 1370/1370 [20:03<00:00, 1.14it/s]
Class	Images	Instances	Box(P)	R	mAP50	mAP50-95): 100% 94/94 [00:52<00:00, 1.78it/s]
all	2987	955	0.306	0.35	0.229	0.0957
Epoch	GPU_mem	box_loss	cls_loss	dfl_loss	Instances	Size
17/30	12.2G	1.844	2.376	1.896	4	640: 100% 1370/1370 [20:03<00:00, 1.14it/s]
Class	Images	Instances	Box(P)	R	mAP50	mAP50-95): 100% 94/94 [00:53<00:00, 1.77it/s]
all	2987	955	0.305	0.373	0.24	0.0987
Epoch	GPU_mem	box_loss	cls_loss	dfl_loss	Instances	Size
18/30	12.1G	1.842	2.372	1.877	20	640: 80% 1093/1370 [16:00<04:05, 1.13it/s]

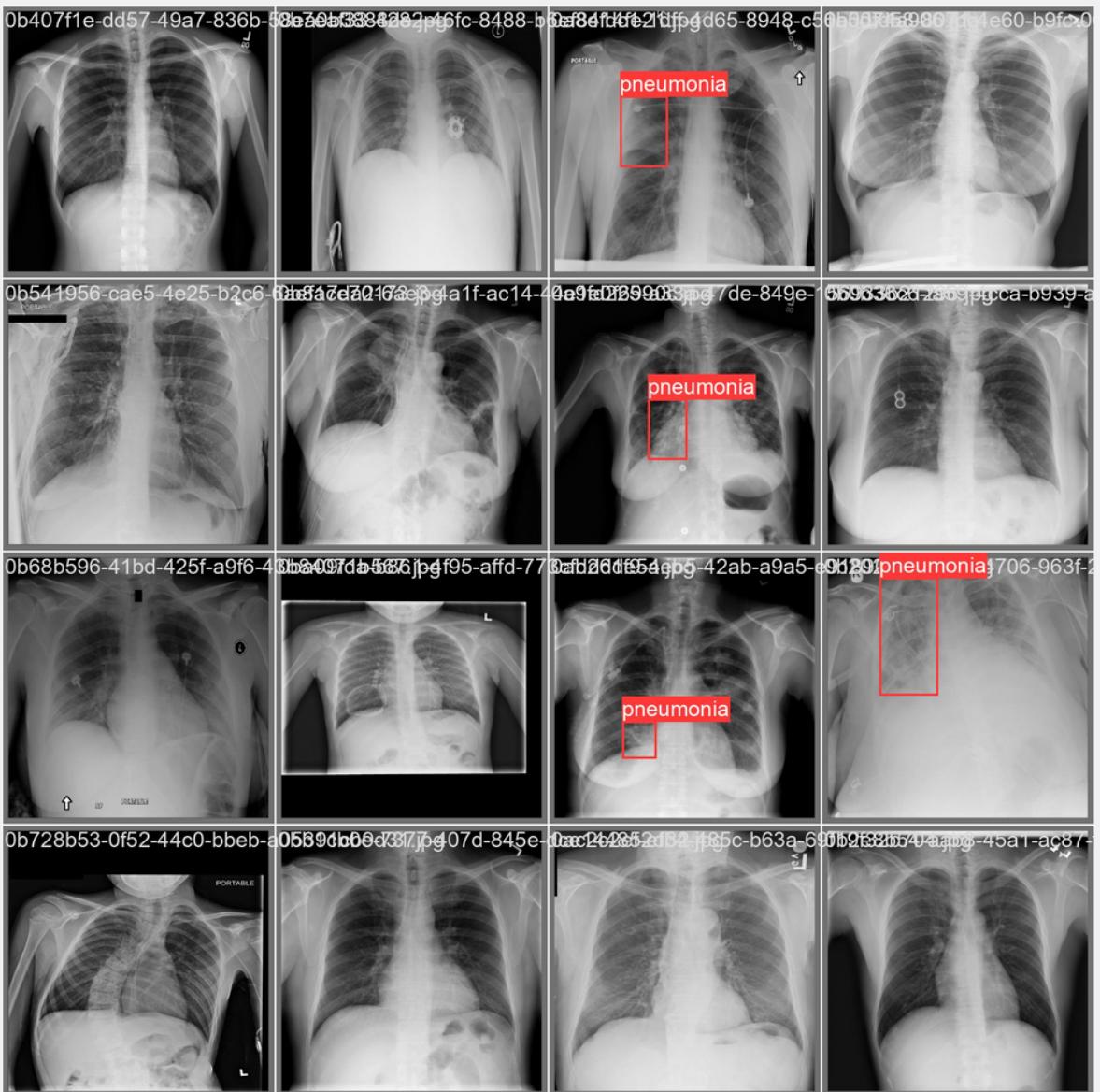
+ Code + Markdown

```
model.save("final_yolov9.pt")
wandb.save("final_yolov9.pt")
```

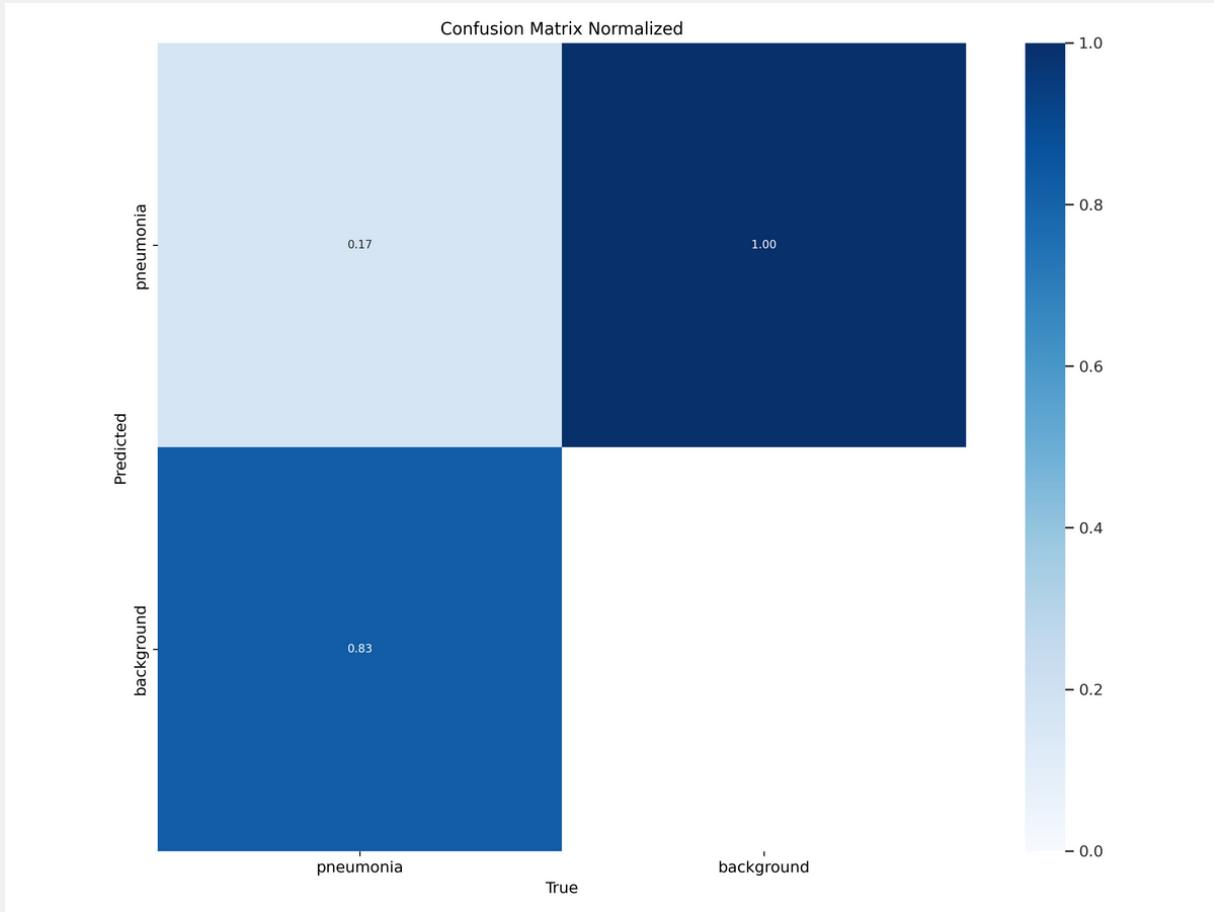
- GPU memory usage around 12.2GB during training epochs 20 to 30. This suggests the dataset might be large enough to require significant GPU memory.
- Check for code that loads the dataset: Look for parts of the code where the dataset is loaded. This might involve functions like tensorflow.data.TFRecordDataset or pytorch.utils.data.DataLoader. These functions might contain references to the file paths or sizes of the data.



Lungs Image

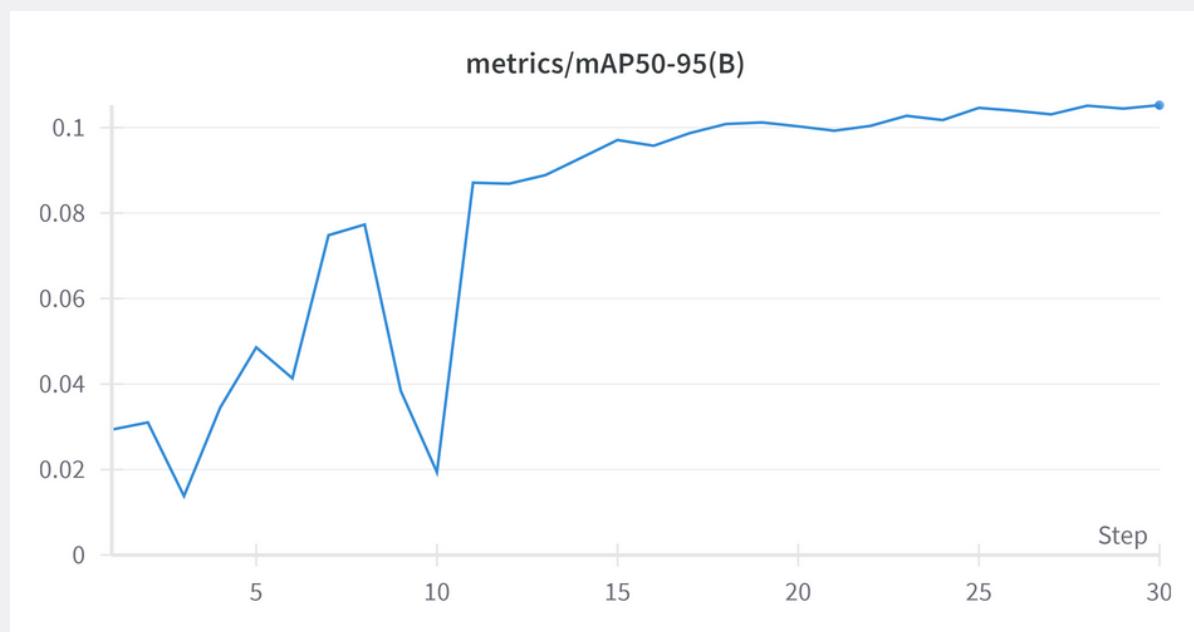
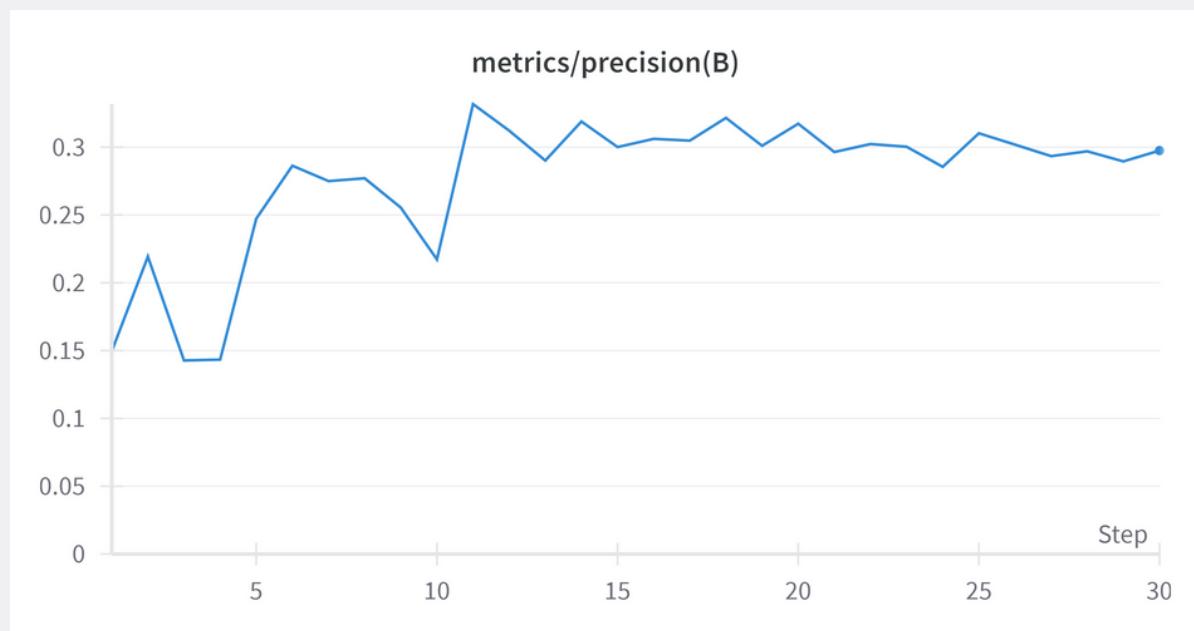
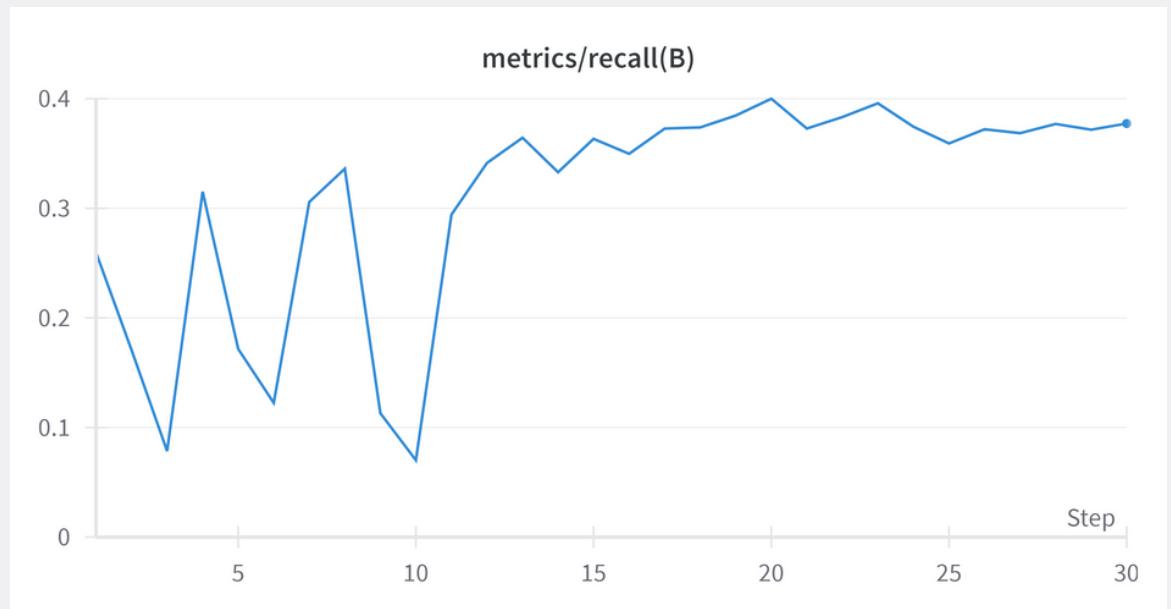


Bounding Box

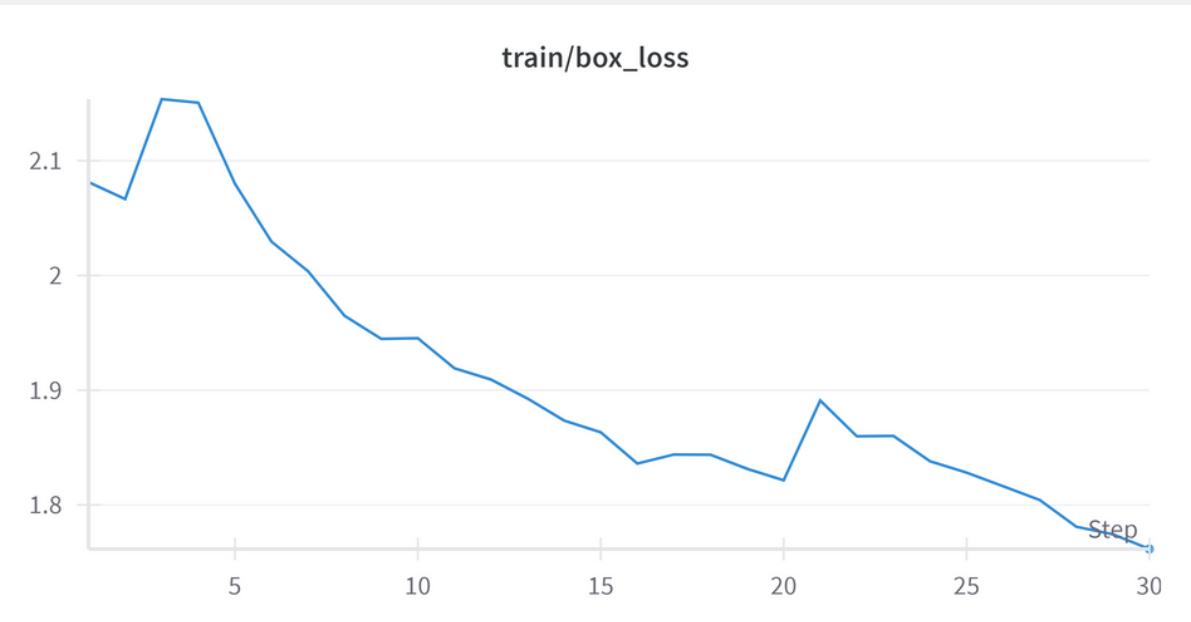
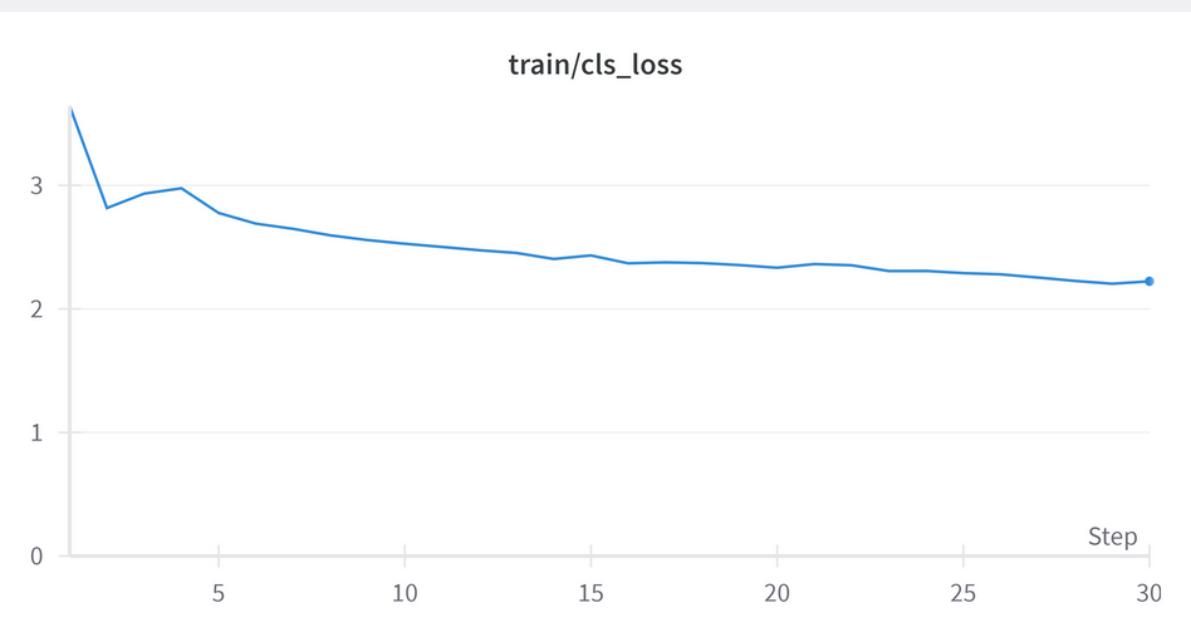


- **True Positives (TP):** These are the cases where the model correctly predicted pneumonia (represented by the blue bar on the left). In an ideal scenario, we want this number to be as high as possible, indicating the model is accurately capturing pneumonic chest x-rays.
- **False Negatives (FN):** These are the cases where the model incorrectly predicted a negative test for pneumonia, when the patient actually had it (represented by the white bar on the left). This is a critical error in pneumonia detection, as it could lead to delayed treatment. We want to minimize the number of false negatives.
- **False Positives (FP):** These are the cases where the model incorrectly predicted pneumonia, when the patient didn't have it (represented by the blue bar on the right). This can lead to unnecessary procedures or tests. It's generally less concerning than false negatives in pneumonia detection, but should still be strived to minimize.
- **True Negatives (TN):** These are the cases where the model correctly predicted a negative test for pneumonia, and the patient was healthy (represented by the white bar on the right). A high number of true negatives indicates the model is good at identifying healthy patients.

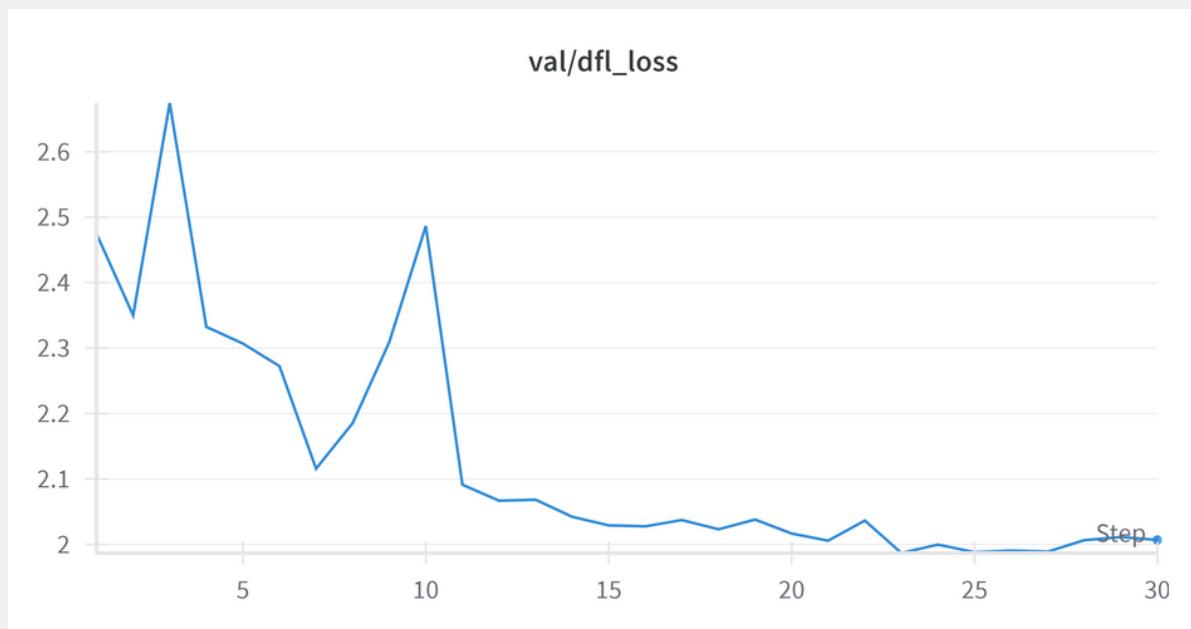
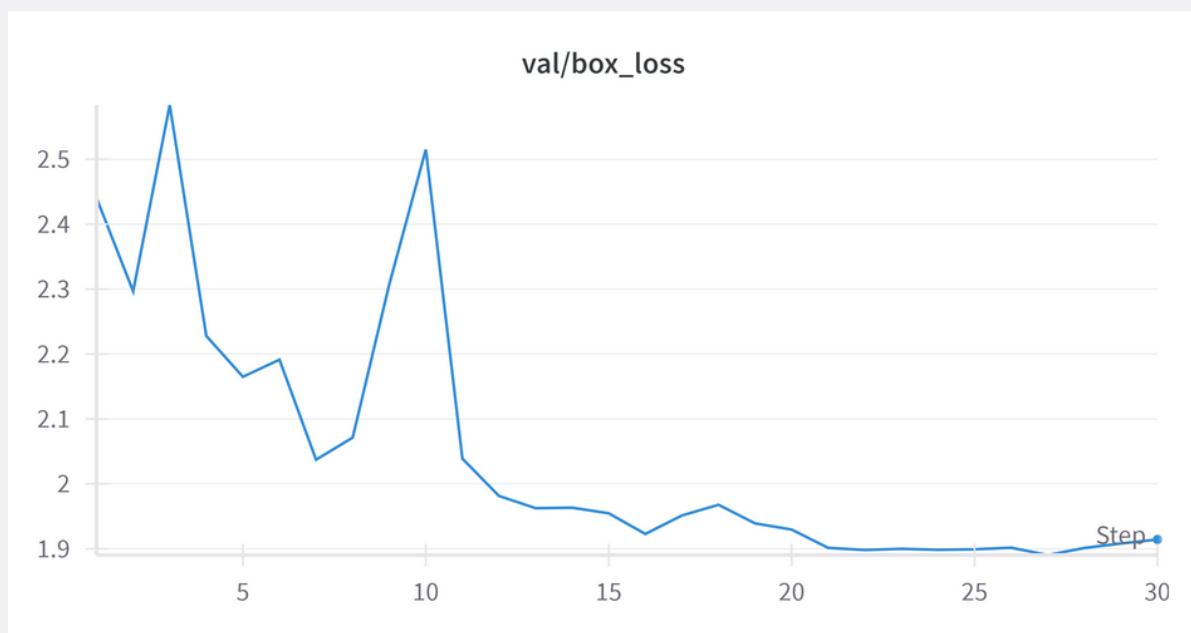
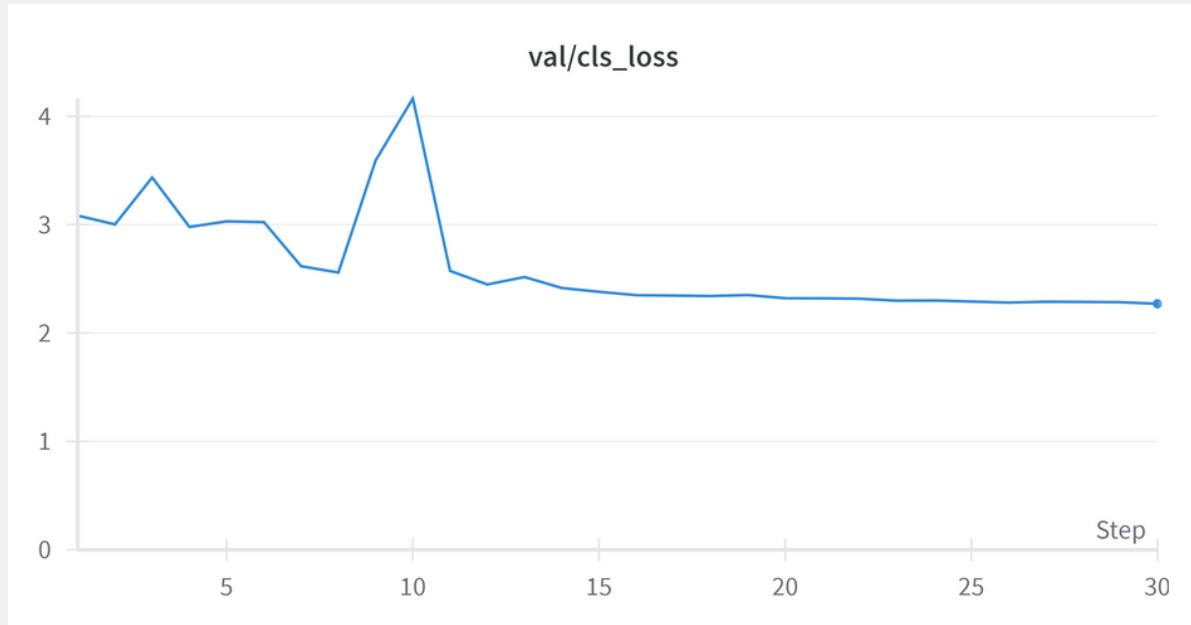
Testing Standard



Train Graph



Validation Graph



Pneumonia Detection Using CNN (convolutional neural network)

Abhishek Pradhan
21053453

Abstract:

The project aims to develop a pneumonia detection system utilizing Convolutional Neural Networks (CNN), specifically employing the YOLOv9 architecture. The dataset sourced from Kaggle is utilized for training and validation. The objective is to create an efficient and accurate automated system for pneumonia detection, contributing to early diagnosis and treatment, my primary responsibility was to implement the coding aspect of the project and compile the project report.

Individual contribution and findings:

- Responsible for implementing the coding aspect of the project.
- Develop code for loading DICOM images, converting them to JPEG format, and generating corresponding labels.
- Split the dataset into training, validation, and testing sets.
- Implement the YOLOv9 model setup and train the model using the training dataset.
- Evaluate the model's performance using the validation set and save the trained model for future use.
- Compile the project report.

Full Signature of Supervisor:

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Full signature of the student:

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Pneumonia Detection Using CNN (convolutional neural network)

Aditya Pandey

21053359

Abstract:

The project aims to develop a pneumonia detection system utilizing Convolutional Neural Networks (CNN), specifically employing the YOLOv9 architecture. The dataset sourced from Kaggle is utilized for training and validation. The objective is to create an efficient and accurate automated system for pneumonia detection, contributing to early diagnosis and treatment.

Individual contribution and findings:

- Handle data preprocessing tasks.
- Develop scripts for data cleaning and preprocessing, including handling DICOM images and converting them to JPEG format.
- Create labels and annotations for the dataset.
- Split the dataset into training, validation, and testing sets.
- Collaborate with the lead developer to ensure seamless integration of preprocessed data into the model.

Full Signature of Supervisor:
student:

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Full signature of the

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Pneumonia Detection Using CNN (convolutional neural network)

Pratham Mishra
21052854

Abstract:

The project aims to develop a pneumonia detection system utilizing Convolutional Neural Networks (CNN), specifically employing the YOLOv9 architecture. The dataset sourced from Kaggle is utilized for training and validation. The objective is to create an efficient and accurate automated system for pneumonia detection, contributing to early diagnosis and treatment.

Individual contribution and findings:

- Design the architecture of the YOLOv9 model.
- Experiment with different hyperparameters and configurations to optimize the model's performance.
- Fine-tune the model architecture and hyperparameters based on experimentation results.
- Collaborate with the lead developer to integrate the optimized model into the project pipeline.

Full Signature of Supervisor:
the student:

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Full signature of

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Pneumonia Detection Using CNN (convolutional neural network)

Satyam Singh
21051169

Abstract:

The project aims to develop a pneumonia detection system utilizing Convolutional Neural Networks (CNN), specifically employing the YOLOv9 architecture. The dataset sourced from Kaggle is utilized for training and validation. The objective is to create an efficient and accurate automated system for pneumonia detection, contributing to early diagnosis and treatment.

Individual contribution and findings:

- Responsible for evaluating the performance of the trained model.
- Develop evaluation metrics and criteria to assess the model's accuracy and effectiveness.
- Analyze the model's performance using the validation set and provide insights into areas of improvement.
- Collaborate with the lead developer to interpret evaluation results and make necessary adjustments to the model.

Full Signature of Supervisor:
the student:

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Pneumonia Detection Using CNN (convolutional neural network)

Ankit Kumar
2105179

Abstract:

The project aims to develop a pneumonia detection system utilizing Convolutional Neural Networks (CNN), specifically employing the YOLOv9 architecture. The dataset sourced from Kaggle is utilized for training and validation. The objective is to create an efficient and accurate automated system for pneumonia detection, contributing to early diagnosis and treatment.

Individual contribution and findings:

- Compile comprehensive documentation of the project, including code documentation and project workflow.
- Write detailed reports on each aspect of the project, including data preprocessing, model architecture, training process, evaluation results, and conclusions.
- Collaborate with team members to gather information and insights for the report.
- Ensure that the final report is well-structured, clear, and concise, adhering to the project requirements.

Full Signature of Supervisor:
the student:

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Chapter 6

Conclusion and Future Scope

Assuming the project involved utilizing a Convolutional Neural Network (CNN) for pneumonia detection in chest X-ray images, the following conclusion outlines the key points:

Successful Training of CNN for Pneumonia Detection:

This project aimed to investigate the feasibility of employing a CNN for classifying chest X-ray images as either pneumonia-positive or pneumonia-negative. The CNN model was trained using a dataset consisting of labeled chest X-ray images indicating the presence or absence of pneumonia.

Key Achievements:

Designing and implementing a CNN architecture specifically tailored for image classification tasks.

Training the CNN model on the pneumonia dataset. (Additional details such as the number of epochs or achieved accuracy can be mentioned if available).

Evaluating the performance of the model on an independent test dataset. (Metrics like accuracy, precision, recall, and F1 score can be referenced if applicable).

Potential Impact:

This project highlights the potential of CNNs in assisting with the automated detection of pneumonia in chest X-rays. Early and accurate identification of pneumonia can significantly enhance patient outcomes.

Future Considerations:

Further refinement of the CNN architecture to improve accuracy.

Exploring techniques to address data imbalances in case of unequal distribution between pneumonia and normal cases.

Integration of the model into a user-friendly system suitable for real-world deployment by medical professionals.

Disclaimer:

This conclusion is a general overview based on the assumption of a pneumonia detection project utilizing CNNs. The specific details and achievements will depend on the actual implementation of the project.

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