

```

1 #Spinal Fractures YOLOv11n, 50 epochs, 256x256 images
2 #Jory Hamilton
3 #12/5/25
4 #Gemini-assisted code

```

```

1 #Install Kaggle API, pyyaml, ultralytics enviroment
2 !pip install -q kaggle
3 !pip install pyyaml
4 !pip install ultralytics --upgrade #Enviroment

```

```

Requirement already satisfied: pyyaml in /usr/local/lib/python3.12/dist-packages (6.0.3)
Collecting ultralytics
  Downloading ultralytics-8.3.235-py3-none-any.whl.metadata (37 kB)
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Requirement already satisfied: six>=1.5 in /usr/local/lib/python3.12/dist-packages (from python-dateutil>=2.7->mat
Requirement already satisfied: mpmath<1.4,>=1.1.0 in /usr/local/lib/python3.12/dist-packages (from sympy>=1.13.3->
Requirement already satisfied: MarkupSafe>=2.0 in /usr/local/lib/python3.12/dist-packages (from Jinja2->torch>=1.8
Downloading ultralytics-8.3.235-py3-none-any.whl (1.1 MB)
1.1/1.1 MB 65.8 MB/s eta 0:00:00
Downloading ultralytics_thop-2.0.18-py3-none-any.whl (28 kB)
Installing collected packages: ultralytics-thop, ultralytics

```

```

1 #Import libraries
2 import os
3 from google.colab import files

```

```

4 import sys
5 import h5py
6 import numpy as np
7 import matplotlib.pyplot as plt
8 import matplotlib.patches as patches
9 import random
10 from ultralytics import YOLO
11 import yaml
12 import pandas as pd

```

```

1 #Check if the API key already exists
2 if not os.path.exists('/root/.kaggle/kaggle.json'):
3     print('Please upload your personal kaggle.json file:')
4
5     #Prompt for upload
6     uploaded = files.upload()
7
8     #Move the file to the correct directory
9     !mkdir -p ~/.kaggle
10    !cp kaggle.json ~/.kaggle/
11    !chmod 600 ~/.kaggle/kaggle.json
12    print('\nKaggle API key configured successfully.')
13 else:
14    print('Kaggle API key is already configured.')

```

Please upload your personal kaggle.json file:

No file chosen

Upload widget is only available when the cell has been executed in the current browser session.

Please rerun this cell to enable.

Saving kaggle.json to kaggle.json

Kaggle API key configured successfully.

```

1 #Replace with your dataset id here
2 KAGGLE_DATASET_ID = 'andymalinsky/rsna-2022-hdf5-subset'
3
4 print('Downloading dataset...')
5 !kaggle datasets download -d {KAGGLE_DATASET_ID}

```

Downloading dataset...

Dataset URL: <https://www.kaggle.com/datasets/andymalinsky/rsna-2022-hdf5-subset>

License(s): CC0-1.0

Downloading rsna-2022-hdf5-subset.zip to /content

96% 3.38G/3.51G [00:08<00:00, 286MB/s]

100% 3.51G/3.51G [00:08<00:00, 436MB/s]

```

1 # Unzip and move to the local Colab storage for training
2 zip_name = KAGGLE_DATASET_ID.split('/')[1] + '.zip'
3 !unzip -q '{zip_name}'
4
5 # This is the path the training script will use
6 HDF5_FILE_PATH = '/content/fracture_dataset_subset.h5'

```

Data Prep for YOLOv11n

```

1 #Load HDF5 Dataset
2 print(f"Opening HDF5 file: {HDF5_FILE_PATH}")
3 with h5py.File(HDF5_FILE_PATH, 'r') as hf:
4     print("HDF5 file opened successfully.")
5     print("Keys in the HDF5 file:")
6     for key in hf.keys():
7         print(f"- {key}")

```

Opening HDF5 file: /content/fracture\_dataset\_subset.h5

HDF5 file opened successfully.

Keys in the HDF5 file:

- SliceNumber
- StudyInstanceUID
- bboxes

```
- images
- labels
- split
```

```
1 #Prepare Directory Structure
2
3 YOLO_DATA_DIR = 'yolo_dataset'
4
5 #Create the main output directory
6 os.makedirs(YOLO_DATA_DIR, exist_ok=True)
7 print(f"Main YOLO dataset directory created: {YOLO_DATA_DIR}")
8
9 #Define dataset splits
10 splits = ['train', 'val', 'test']
11
12 #Create subdirectories for each split (images and labels)
13 for split in splits:
14     images_dir = os.path.join(YOLO_DATA_DIR, split, 'images')
15     labels_dir = os.path.join(YOLO_DATA_DIR, split, 'labels')
16     os.makedirs(images_dir, exist_ok=True)
17     os.makedirs(labels_dir, exist_ok=True)
18     print(f"Created directories for {split}: {images_dir} and {labels_dir}")
19
20 print("YOLO dataset directory structure prepared successfully.")
```

```
Main YOLO dataset directory created: yolo_dataset
Created directories for train: yolo_dataset/train/images and yolo_dataset/train/labels
Created directories for val: yolo_dataset/val/images and yolo_dataset/val/labels
Created directories for test: yolo_dataset/test/images and yolo_dataset/test/labels
YOLO dataset directory structure prepared successfully.
```

```
1 #Process and Convert Annotations
2 hf = h5py.File(HDF5_FILE_PATH, 'r')
3 images_dataset = hf['images']
4 bboxes_dataset = hf['bboxes']
5 labels_dataset = hf['labels']
6 split_dataset = hf['split']
7 print("HDF5 datasets loaded successfully.")
```

```
HDF5 datasets loaded successfully.
```

```
1 def convert_to_yolo_coords(image_width, image_height, bbox):
2     #bbox: [x_min, y_min, x_max, y_max]
3     x_min, y_min, x_max, y_max = bbox
4
5     #Clamp coordinates to image boundaries and ensure they are floats for calculations
6     x_min_clamped = max(0.0, float(x_min))
7     y_min_clamped = max(0.0, float(y_min))
8     x_max_clamped = min(float(image_width), float(x_max))
9     y_max_clamped = min(float(image_height), float(y_max))
10
11     #Recalculate width and height after clamping
12     box_width = x_max_clamped - x_min_clamped
13     box_height = y_max_clamped - y_min_clamped
14
15     #If the box is invalid (zero or negative area after clamping), return zeros
16     if box_width <= 0 or box_height <= 0:
17         return [0.0, 0.0, 0.0, 0.0] # Return a zero-area bbox, YOLO will likely ignore it
18
19     #Calculate center coordinates
20     x_center = (x_min_clamped + x_max_clamped) / 2
21     y_center = (y_min_clamped + y_max_clamped) / 2
22
23     #Normalize coordinates
24     x_center_norm = x_center / image_width
25     y_center_norm = y_center / image_height
26     width_norm = box_width / image_width
27     height_norm = box_height / image_height
28
```

```

29     #Ensure normalized coordinates are within [0, 1] range (should be due to clamping)
30     x_center_norm = max(0.0, min(1.0, x_center_norm))
31     y_center_norm = max(0.0, min(1.0, y_center_norm))
32     width_norm = max(0.0, min(1.0, width_norm))
33     height_norm = max(0.0, min(1.0, height_norm))
34
35     return [x_center_norm, y_center_norm, width_norm, height_norm]
36
37 #Initialize a list to store processed data
38 #Each item will be a dictionary containing 'image', 'split', and 'yolo_annotations'
39 processed_data = []
40
41 print("convert_to_yolo_coords function defined and processed_data list initialized.")

```

convert\_to\_yolo\_coords function defined and processed\_data list initialized.

```

1 #Re-open the HDF5 file and assign datasets inside this cell for robustness
2 with h5py.File(HDF5_FILE_PATH, 'r') as hf:
3     images_dataset = hf['images']
4     bboxes_dataset = hf['bboxes']
5     labels_dataset = hf['labels']
6     split_dataset = hf['split']
7
8     #Clear processed_data list to prevent duplicate entries if re-running
9     processed_data = []
10
11     for i in range(len(images_dataset)):
12         image_data = images_dataset[i]
13         image_height, image_width = image_data.shape[:2]
14
15         # HDF5 stores lists of lists, so we need to iterate if there are multiple bounding boxes
16         raw_bboxes_for_image = bboxes_dataset[i]
17         labels_for_image = labels_dataset[i]
18
19         #Map the numerical split ID to its string representation
20         #Assuming splits_dataset contains integers 0, 1, 2 corresponding to 'train', 'val', 'test'
21         #The 'splits' variable is available from previous steps: splits = ['train', 'val', 'test']
22         split_index = int(split_dataset[i]) # Ensure it's an integer for indexing
23         split_for_image = splits[split_index]
24
25         yolo_annotations = []
26         #Check if there are bounding boxes for the current image
27         if raw_bboxes_for_image.size > 0 and not np.all(raw_bboxes_for_image == -1): # Also check if not all a
28             for j in range(len(raw_bboxes_for_image)):
29                 bbox = raw_bboxes_for_image[j]
30                 if np.all(bbox == -1): # Skip placeholder bounding boxes
31                     continue
32
33                 #Use labels_for_image directly as it appears to be a scalar class ID for the image
34                 class_id = int(labels_for_image)
35
36                 yolo_coords = convert_to_yolo_coords(image_width, image_height, bbox)
37                 #YOLO format: class_id x_center y_center width height
38                 yolo_annotations.append(f"{int(class_id)} {yolo_coords[0]:.6f} {yolo_coords[1]:.6f} {yolo_coor
39
40         processed_data.append({
41             'image_index': i,
42             'image_data': image_data,
43             'split': split_for_image,
44             'yolo_annotations': yolo_annotations
45         })
46
47 print(f"Processed {len(processed_data)} images and their annotations.")
48 print("HDF5 file operations completed.")
49

```

Processed 28812 images and their annotations.  
HDF5 file operations completed.

## Save Images and YOLO Labels

```

1 for item in processed_data:
2     image_index = item['image_index']
3     image_data = item['image_data']
4     split = item['split']
5     yolo_annotations = item['yolo_annotations']
6
7     #Format image_index with leading zeros (e.g., 000001)
8     #Assuming a maximum of 6 digits for image index based on typical dataset sizes
9     formatted_image_index = str(image_index).zfill(6)
10
11    #Construct image file path
12    image_filename = f"{formatted_image_index}.jpeg"
13    image_path = os.path.join(YOLO_DATA_DIR, split, 'images', image_filename)
14
15    #Prepare image data for saving
16    #plt.imshow expects float data to be in [0, 1] range for imshow,
17    #or it will normalize if the dtype is float and min/max are outside [0,1].
18    #Assuming image_data is already normalized or within a reasonable range for direct saving.
19    #If it's a single-channel (grayscale) image, convert to 3-channel for consistent JPEG output
20    if image_data.ndim == 2 or (image_data.ndim == 3 and image_data.shape[-1] == 1):
21        # Stack the single channel to make it 3-channel for JPEG compatibility and consistent visualization
22        image_to_save = np.stack([image_data.squeeze()] * 3, axis=-1)
23    else:
24        image_to_save = image_data
25
26    #Save image as JPEG
27    plt.imshow(image_to_save, cmap='gray') # Use cmap='gray' for potentially grayscale output
28
29    #Construct label file path
30    label_filename = f"{formatted_image_index}.txt"
31    label_path = os.path.join(YOLO_DATA_DIR, split, 'labels', label_filename)
32
33    #Write YOLO annotations to the label file
34    with open(label_path, 'w') as f:
35        if yolo_annotations:
36            for annotation in yolo_annotations:
37                f.write(annotation + '\n')
38
39 print("Images and YOLO labels saved successfully to their respective directories.")

```

```
ERROR:root:Unexpected exception finding object shape
Traceback (most recent call last):
  File "/usr/local/lib/python3.12/dist-packages/google/colab/_debugpy_repr.py", line 54, in get_shape
    shape = getattr(obj, 'shape', None)
            ^^^^^^^^^^^^^^^^^^^^^^^^^^^^^
  File "/usr/local/lib/python3.12/dist-packages/h5py/_hl/dataset.py", line 539, in shape
    shape = self.id.shape
            ^^^^^^^^^^^^^
  File "h5py/h5d.pyx", line 205, in h5py.h5d.DatasetID.shape.__get__
  File "h5py/h5d.pyx", line 206, in h5py.h5d.DatasetID.shape.__get__
  File "h5py/_objects.pyx", line 54, in h5py._objects.with_phil.wrapper
  File "h5py/_objects.pyx", line 55, in h5py._objects.with_phil.wrapper
  File "h5py/h5d.pyx", line 372, in h5py.h5d.DatasetID.get_space
RuntimeError: Unable to synchronously get dataspace (identifier is not of specified type)
ERROR:root:Unexpected exception finding object shape
Traceback (most recent call last):
  File "/usr/local/lib/python3.12/dist-packages/google/colab/_debugpy_repr.py", line 54, in get_shape
    shape = getattr(obj, 'shape', None)
            ^^^^^^^^^^^^^^^^^^^^^^^^^^^^^
  File "/usr/local/lib/python3.12/dist-packages/h5py/_hl/dataset.py", line 539, in shape
    shape = self.id.shape
            ^^^^^^^^^^^^^
  File "h5py/h5d.pyx", line 205, in h5py.h5d.DatasetID.shape.__get__
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RuntimeError: Unable to synchronously get dataspace (identifier is not of specified type)
ERROR:root:Unexpected exception finding object shape
```

```

Traceback (most recent call last):
  File "/usr/local/lib/python3.12/dist-packages/google/colab/_debugpy_repr.py", line 54, in get_shape
    shape = getattr(obj, 'shape', None)
            ^^^^^^^^^^^^^^^^^^^^^^^^^^^^^
  File "/usr/local/lib/python3.12/dist-packages/h5py/_hl/dataset.py", line 539, in shape
    shape = self.id.shape
            ^^^^^^^^^^^^^
  File "h5py/h5d.pyx", line 205, in h5py.h5d.DatasetID.shape.__get__
  File "h5py/h5d.pyx", line 206, in h5py.h5d.DatasetID.shape.__get__
  File "h5py/_objects.pyx", line 54, in h5py._objects.with_phil.wrapper
  File "h5py/_objects.pyx", line 55, in h5py._objects.with_phil.wrapper
  File "h5py/h5d.pyx", line 372, in h5py.h5d.DatasetID.get_space
RuntimeError: Unable to synchronously get dataspace (identifier is not of specified type)
Images and YOLO labels saved successfully to their respective directories.

```

```

1 #Define the full path where the data.yaml file will be created
2 yaml_file_path = os.path.join(YOLO_DATA_DIR, 'data.yaml')
3
4 #Determine the number of classes (nc) and a list of corresponding class names (names)
5 nc = 2 # Assuming 0 for 'no fracture' and 1 for 'fracture' based on common dataset labeling for binary classif
6 names = ['no_fracture', 'fracture'] # Adjust based on actual class IDs present in HDF5 labels_dataset if needed
7
8 #Construct a Python dictionary containing the YOLO dataset configuration
9 data_yaml_content = {
10     'path': os.path.abspath(YOLO_DATA_DIR), # Absolute path to the dataset root
11     'train': 'train/images',
12     'val': 'val/images',
13     'test': 'test/images',
14     'nc': nc,
15     'names': names
16 }
17
18 #Use yaml.dump() to write the created dictionary to the data.yaml file
19 with open(yaml_file_path, 'w') as f:
20     yaml.dump(data_yaml_content, f)
21
22 #Print a confirmation message
23 print(f"data.yaml has been successfully generated at: {yaml_file_path}")
24
25 #For verification, print the entire content of the newly created data.yaml file
26 print("\nContent of data.yaml:")
27 with open(yaml_file_path, 'r') as f:
28     print(f.read())

```

data.yaml has been successfully generated at: yolo\_dataset/data.yaml

Content of data.yaml:

```

names:
- no_fracture
- fracture
nc: 2
path: /content/yolo_dataset
test: test/images
train: train/images
val: val/images

```

```

1 #Load the model
2 #yolo11n.pt is the 'nano' version (fastest).
3 model = YOLO("yolo11n.pt")
4
5 #Train the model
6 results = model.train(
7     data=yaml_file_path, # Path to your config file, corrected to use the variable
8     epochs=50,           # 50-100 is usually a good start
9     imgsz=256,           # X-ray resolution
10    batch=16,             # Adjust based on GPU VRAM
11    name="fracture_v11_run4", # Name of the run (re-using previous name for consistency)
12    augment=True,          # Data augmentation (vital for small datasets)
13    optimizer='auto',     # Automatically selects AdamW or SGD
14    patience=30            # Early stopping patience

```

```

15 )
16
17 #Validation
18 #Evaluate performance on the validation set
19 metrics = model.val()
20 print(f"mAP@50: {metrics.box.map50}")

```

Downloading <https://github.com/ultralytics/assets/releases/download/v8.3.0/yolo11n.pt> to 'yolo11n.pt': 100% —  
 Ultralytics 8.3.235 Python-3.12.12 torch-2.9.0+cu126 CUDA:0 (NVIDIA L4, 22693MiB)  
**engine/trainer:** agnostic\_nms=False, amp=True, augment=True, auto\_augment=randaugument, batch=16, bgr=0.0, box=7.5,  
 Downloading <https://ultralytics.com/assets/Arial.ttf> to '/root/.config/Ultralytics/Arial.ttf': 100% — 7  
 Overriding model.yaml nc=80 with nc=2

|    | from         | n | params | module                               | arguments                 |
|----|--------------|---|--------|--------------------------------------|---------------------------|
| 0  | -1           | 1 | 464    | ultralytics.nn.modules.conv.Conv     | [3, 16, 3, 2]             |
| 1  | -1           | 1 | 4672   | ultralytics.nn.modules.conv.Conv     | [16, 32, 3, 2]            |
| 2  | -1           | 1 | 6640   | ultralytics.nn.modules.block.C3k2    | [32, 64, 1, False, 0.25]  |
| 3  | -1           | 1 | 36992  | ultralytics.nn.modules.conv.Conv     | [64, 64, 3, 2]            |
| 4  | -1           | 1 | 26080  | ultralytics.nn.modules.block.C3k2    | [64, 128, 1, False, 0.25] |
| 5  | -1           | 1 | 147712 | ultralytics.nn.modules.conv.Conv     | [128, 128, 3, 2]          |
| 6  | -1           | 1 | 87040  | ultralytics.nn.modules.block.C3k2    | [128, 128, 1, True]       |
| 7  | -1           | 1 | 295424 | ultralytics.nn.modules.conv.Conv     | [128, 256, 3, 2]          |
| 8  | -1           | 1 | 346112 | ultralytics.nn.modules.block.C3k2    | [256, 256, 1, True]       |
| 9  | -1           | 1 | 164608 | ultralytics.nn.modules.block.SPPF    | [256, 256, 5]             |
| 10 | -1           | 1 | 249728 | ultralytics.nn.modules.block.C2PSA   | [256, 256, 1]             |
| 11 | -1           | 1 | 0      | torch.nn.modules.upsampling.Upsample | [None, 2, 'nearest']      |
| 12 | [-1, 6]      | 1 | 0      | ultralytics.nn.modules.conv.Concat   | [1]                       |
| 13 | -1           | 1 | 111296 | ultralytics.nn.modules.block.C3k2    | [384, 128, 1, False]      |
| 14 | -1           | 1 | 0      | torch.nn.modules.upsampling.Upsample | [None, 2, 'nearest']      |
| 15 | [-1, 4]      | 1 | 0      | ultralytics.nn.modules.conv.Concat   | [1]                       |
| 16 | -1           | 1 | 32096  | ultralytics.nn.modules.block.C3k2    | [256, 64, 1, False]       |
| 17 | -1           | 1 | 36992  | ultralytics.nn.modules.conv.Conv     | [64, 64, 3, 2]            |
| 18 | [-1, 13]     | 1 | 0      | ultralytics.nn.modules.conv.Concat   | [1]                       |
| 19 | -1           | 1 | 86720  | ultralytics.nn.modules.block.C3k2    | [192, 128, 1, False]      |
| 20 | -1           | 1 | 147712 | ultralytics.nn.modules.conv.Conv     | [128, 128, 3, 2]          |
| 21 | [-1, 10]     | 1 | 0      | ultralytics.nn.modules.conv.Concat   | [1]                       |
| 22 | -1           | 1 | 378880 | ultralytics.nn.modules.block.C3k2    | [384, 256, 1, True]       |
| 23 | [16, 19, 22] | 1 | 431062 | ultralytics.nn.modules.head.Detect   | [2, [64, 128, 256]]       |

YOLO11n summary: 181 layers, 2,590,230 parameters, 2,590,214 gradients, 6.4 GFLOPs

Transferred 448/499 items from pretrained weights

Freezing layer 'model.23.dfl.conv.weight'

**AMP:** running Automatic Mixed Precision (AMP) checks...

**AMP:** checks passed ✓

**train:** Fast image access ✓ (ping: 0.0±0.0 ms, read: 308.9±155.2 MB/s, size: 6.7 KB)

**train:** Scanning /content/yolo\_dataset/train/labels... 20744 images, 15558 backgrounds, 0 corrupt: 100% —

**train:** New cache created: /content/yolo\_dataset/train/labels.cache

**albumentations:** Blur(p=0.01, blur\_limit=(3, 7)), MedianBlur(p=0.01, blur\_limit=(3, 7)), ToGray(p=0.01, method='wei

**val:** Fast image access ✓ (ping: 0.0±0.0 ms, read: 349.3±197.7 MB/s, size: 7.9 KB)

**val:** Scanning /content/yolo\_dataset/val/labels... 4180 images, 3135 backgrounds, 0 corrupt: 100% — 4180

**val:** New cache created: /content/yolo\_dataset/val/labels.cache

Plotting labels to /content/runs/detect/fracture\_v11\_run4/labels.jpg...

**optimizer:** 'optimizer=auto' found, ignoring 'lr0=0.01' and 'momentum=0.937' and determining best 'optimizer', 'lr0

**optimizer:** SGD(lr=0.01, momentum=0.9) with parameter groups 81 weight(decay=0.0), 88 weight(decay=0.0005), 87 bias

Image sizes 256 train, 256 val

Using 8 dataloader workers

Logging results to /content/runs/detect/fracture\_v11\_run4

Starting training for 50 epochs...

| Epoch | GPU_mem | box_loss | cls_loss  | dfl_loss | Instances | Size                           |
|-------|---------|----------|-----------|----------|-----------|--------------------------------|
| 1/50  | 0.439G  | 0.493    | 3.347     | 0.3006   | 1         | 256: 100% — 1297/1297 10.0it/  |
|       | Class   | Images   | Instances | Box(P    | R         | mAP50 mAP50-95): 100% — 131/13 |
|       | all     | 4180     | 1045      | 0        | 0         | 0 0                            |

```

1
2
3 #Load custom/trained model
4 #Path points to the 'best.pt' file generated during your training run
5 model = YOLO("/content/runs/detect/fracture_v11_run4/weights/best.pt") #Adjusted path based on previous run na
6
7 #Get a sample image from the test set for inference
8 test_image_dir = os.path.join(YOLO_DATA_DIR, 'test', 'images')
9 #List all files in the test image directory

```

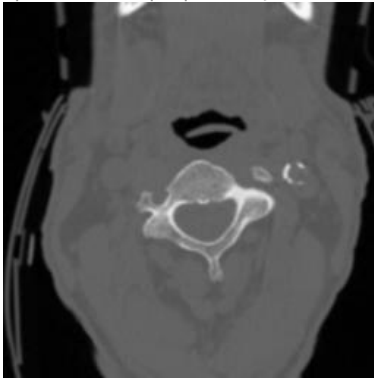
```

10 image_files = os.listdir(test_image_dir)
11
12 #Filter for image files (e.g., .jpeg, .jpg, .png)
13 image_files = [f for f in image_files if f.lower().endswith(('.jpeg', '.jpg', '.png'))]
14
15 if image_files:
16     #Pick the first image found in the test directory
17     sample_image_path = os.path.join(test_image_dir, image_files[0])
18     print(f"Using sample image for inference: {sample_image_path}")
19
20     #Predict on the sample image
21     results = model(sample_image_path)
22
23     #Show results
24     results[0].show()
25 else:
26     print(f"No image files found in {test_image_dir} to perform inference on.")
27

```

Using sample image for inference: yolo\_dataset/test/images/006280.jpeg

image 1/1 /content/yolo\_dataset/test/images/006280.jpeg: 256x256 (no detections), 10.7ms  
Speed: 0.5ms preprocess, 10.7ms inference, 0.7ms postprocess per image at shape (1, 3, 256, 256)



```

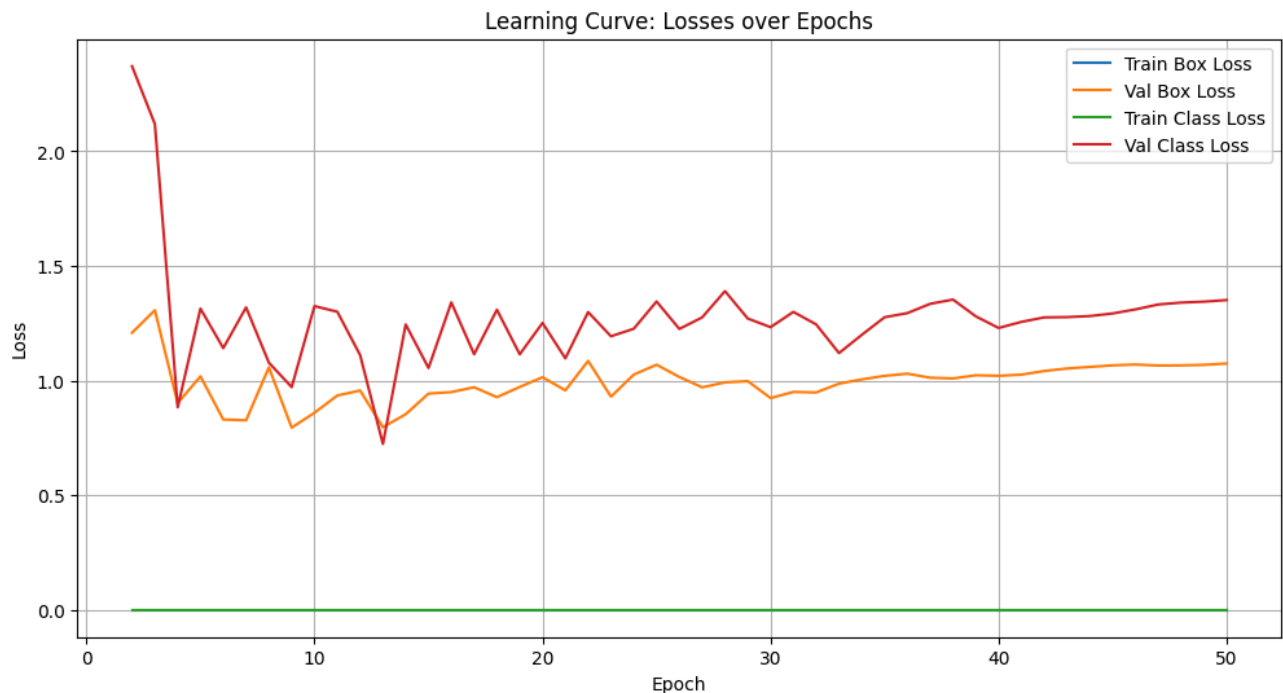
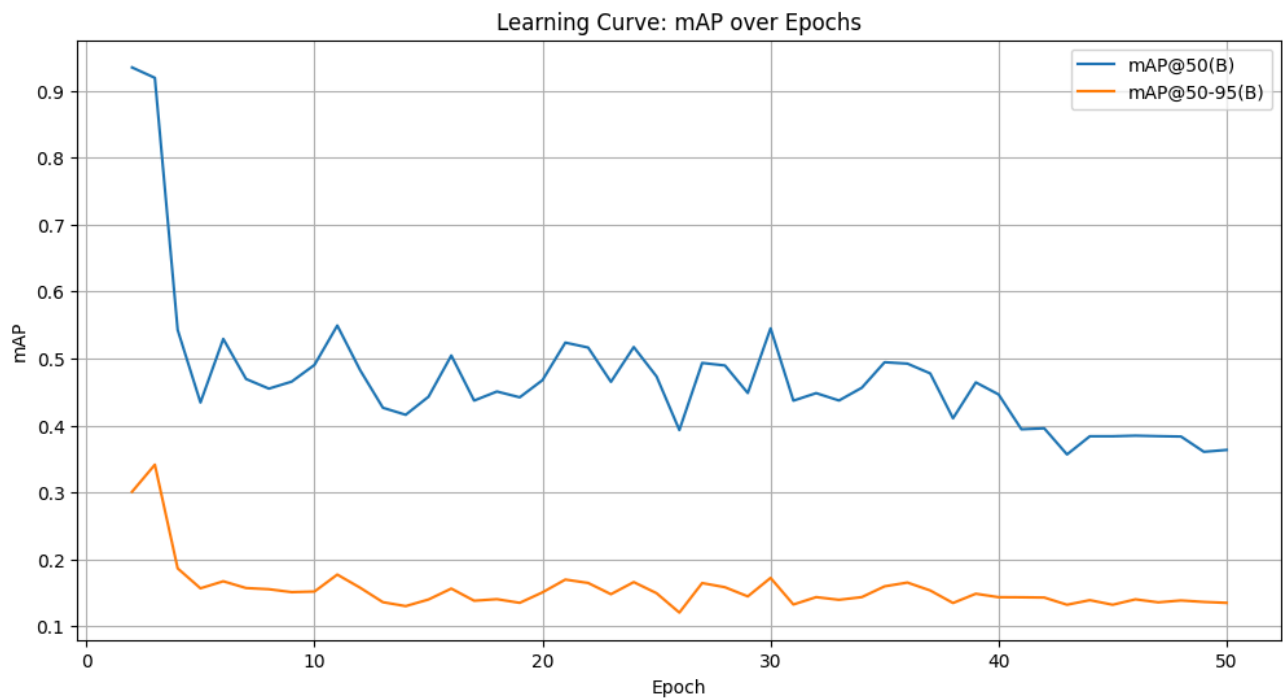
1 #Define the path to the training results CSV file
2 results_path = '/content/runs/detect/fracture_v11_run4/results.csv'
3
4 #Check if the file exists
5 if os.path.exists(results_path):
6     print(f"Loading training results from: {results_path}")
7     # Load the results CSV, skipping the first row which contains headers with special characters
8     # and might cause parsing issues. We'll manually set column names.
9     results_df = pd.read_csv(results_path, skiprows=1)
10
11 #Assign meaningful column names based on Ultralytics output structure
12 #Common columns are 'epoch', 'metrics/precision(B)', 'metrics/recall(B)', 'metrics/mAP50(B)', 'metrics/mAP
13 #Adjust these column names if your results.csv has a different structure after skipping the first row.
14 results_df.columns = [
15     'epoch',
16     'metrics/precision(B)',
17     'metrics/recall(B)',
18     'metrics/mAP50(B)',
19     'metrics/mAP50-95(B)',
20     'train/box_loss',
21     'train/cls_loss',
22     'train/dfl_loss',
23     'train/obj_loss', # Added missing column
24     'val/box_loss',
25     'val/cls_loss',
26     'val/dfl_loss',
27     'val/obj_loss', # Added missing column
28     'labels',
29     'lr'
30 ]
31
32 #Plotting mAP@50 over epochs

```



```
33 plt.figure(figsize=(12, 6))
34 plt.plot(results_df['epoch'], results_df['metrics/mAP50(B)'], label='mAP@50(B)')
35 plt.plot(results_df['epoch'], results_df['metrics/mAP50-95(B)'], label='mAP@50-95(B)')
36 plt.xlabel('Epoch')
37 plt.ylabel('mAP')
38 plt.title('Learning Curve: mAP over Epochs')
39 plt.legend()
40 plt.grid(True)
41 plt.show()
42
43 #Plotting Loss Curves
44 plt.figure(figsize=(12, 6))
45 plt.plot(results_df['epoch'], results_df['train/box_loss'], label='Train Box Loss')
46 plt.plot(results_df['epoch'], results_df['val/box_loss'], label='Val Box Loss')
47 plt.plot(results_df['epoch'], results_df['train/cls_loss'], label='Train Class Loss')
48 plt.plot(results_df['epoch'], results_df['val/cls_loss'], label='Val Class Loss')
49 plt.xlabel('Epoch')
50 plt.ylabel('Loss')
51 plt.title('Learning Curve: Losses over Epochs')
52 plt.legend()
53 plt.grid(True)
54 plt.show()
55
56 else:
57     print(f"Error: Training results file not found at {results_path}")
58     print("Please ensure the training run completed and generated a 'results.csv' file.")
```

Loading training results from: /content/runs/detect/fracture\_v11\_run4/results.csv



```

1 #Construct the full path to the confusion matrix image
2 confusion_matrix_path = os.path.join('/content/runs/detect/fracture_v11_run4', 'confusion_matrix.png')
3
4 #Check if the file exists before attempting to load
5 if os.path.exists(confusion_matrix_path):
6     print(f>Loading confusion matrix from: {confusion_matrix_path}")
7     #Use matplotlib.pyplot.imread() to load the image
8     confusion_matrix_img = plt.imread(confusion_matrix_path)
9
10    #Display the image using matplotlib.pyplot.imshow()
11    plt.figure(figsize=(10, 10))
12    plt.imshow(confusion_matrix_img)
13
14    #Add a title to the plot
15    plt.title("Confusion Matrix for Fracture Detection Model")
16

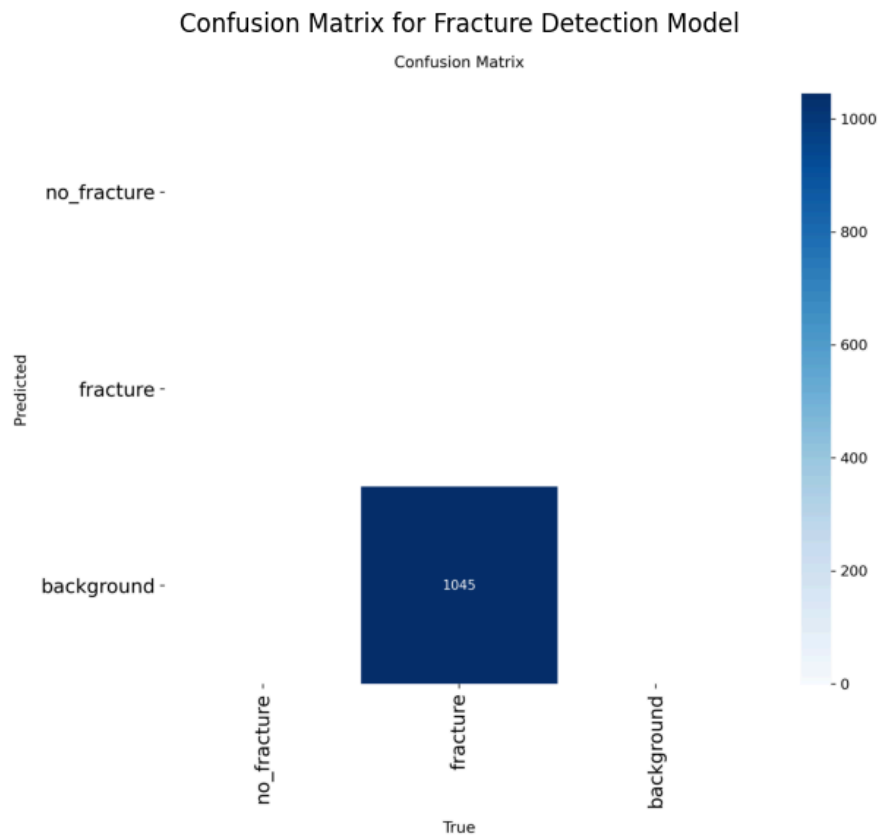
```

```

17 #Use matplotlib.pyplot.axis('off') to hide the axes
18 plt.axis('off')
19
20 #Show the plot
21 plt.show()
22 else:
23     print(f"Error: Confusion matrix not found at {confusion_matrix_path}")

```

Loading confusion matrix from: /content/runs/detect/fracture\_v11\_run4/confusion\_matrix.png



Additional diagnostics and analysis if needed

```

1 #Identify sample images with fractures
2 #Part of the process to verifying YOLO annotations
3
4 #Ensure HDF5_FILE_PATH and splits are defined from previous cells
5 #HDF5_FILE_PATH = '/content/fracture_dataset_subset.h5'
6 #splits = ['train', 'val', 'test']
7
8 #Open the HDF5 file in read mode
9 with h5py.File(HDF5_FILE_PATH, 'r') as hf:
10     print(f"Opened HDF5 file: {HDF5_FILE_PATH}")
11
12 #Access the labels_dataset and split_dataset
13 labels_dataset = hf['labels']
14 split_dataset = hf['split']
15
16 #Initialize an empty list to store fracture image samples
17 fracture_image_samples = []
18 print("Starting to search for fracture images...")
19
20 #Iterate through the labels_dataset
21 for i in range(len(labels_dataset)):
22     # Check if the class_id for the current image is 1 (fracture)
23     class_id = int(labels_dataset[i])
24

```

```

25     #If the class_id is 1, process the image
26     if class_id == 1:
27         # a. Determine its dataset split
28         split_index = int(split_dataset[i])
29         split_for_image = splits[split_index]
30
31         #Add a dictionary containing the image_index and its split to the list
32         fracture_image_samples.append({
33             'image_index': i,
34             'split': split_for_image
35         })
36
37         #Continue until at least 5 images with fractures have been identified
38         if len(fracture_image_samples) >= 5:
39             break
40
41     #Print the fracture_image_samples list
42     print("Identified fracture image samples:")
43     for sample in fracture_image_samples:
44         print(sample)
45

```

```

Opened HDF5 file: /content/fracture_dataset_subset.h5
Starting to search for fracture images...
Identified fracture image samples:
{'image_index': 7, 'split': 'train'}
{'image_index': 15, 'split': 'train'}
{'image_index': 19, 'split': 'train'}
{'image_index': 22, 'split': 'test'}
{'image_index': 25, 'split': 'train'}

```

```

1 #Ensure YOLO_DATA_DIR and names (class labels) are accessible
2 #YOLO_DATA_DIR = 'yolo_dataset'
3 #names = ['no_fracture', 'fracture']
4
5 print(f"Visualizing YOLO annotations for {len(fracture_image_samples)} fracture images...")
6
7 for i, sample in enumerate(fracture_image_samples):
8     image_index = sample['image_index']
9     split = sample['split']
10
11     #Format image_index with leading zeros (e.g., 000001)
12     formatted_image_index = str(image_index).zfill(6)
13
14     #Construct image and label file paths
15     image_filename = f"{formatted_image_index}.jpeg"
16     label_filename = f"{formatted_image_index}.txt"
17
18     image_path = os.path.join(YOLO_DATA_DIR, split, 'images', image_filename)
19     label_path = os.path.join(YOLO_DATA_DIR, split, 'labels', label_filename)
20
21     if not os.path.exists(image_path):
22         print(f"Image not found at {image_path}. Skipping.")
23         continue
24     if not os.path.exists(label_path):
25         print(f"Label file not found at {label_path}. Skipping.")
26         continue
27
28     #Load the JPEG image
29     image = plt.imread(image_path)
30     image_height, image_width = image.shape[:2]
31
32     #Create plot for the image
33     fig, ax = plt.subplots(1, figsize=(8, 8))
34     ax.imshow(image)
35
36     #Read YOLO annotations from the .txt file
37     with open(label_path, 'r') as f:
38         annotations = f.readlines()
39

```

```

40 #Parse and draw bounding boxes
41 if annotations:
42     for annotation_line in annotations:
43         parts = annotation_line.strip().split()
44         if len(parts) == 5:
45             class_id_yolo, x_center_norm, y_center_norm, width_norm, height_norm = map(float, parts)
46             class_id_yolo = int(class_id_yolo)
47
48             # Convert normalized YOLO coordinates to pixel coordinates
49             center_x_px = x_center_norm * image_width
50             center_y_px = y_center_norm * image_height
51             width_px = width_norm * image_width
52             height_px = height_norm * image_height
53
54             x_min_px = center_x_px - (width_px / 2)
55             y_min_px = center_y_px - (height_px / 2)
56
57             # Create a rectangle patch
58             rect = patches.Rectangle(
59                 (x_min_px, y_min_px),
60                 width_px,
61                 height_px,
62                 linewidth=2,
63                 edgecolor='r',
64                 facecolor='none'
65             )
66
67             # Add the rectangle to the plot
68             ax.add_patch(rect)
69
70             # Add class label
71             if class_id_yolo < len(names):
72                 class_name = names[class_id_yolo]
73                 ax.text(
74                     x_min_px,
75                     y_min_px - 5, # Position above the box
76                     f'{class_name}',
77                     color='r',
78                     fontsize=12,
79                     bbox=dict(facecolor='white', alpha=0.7, edgecolor='none', pad=0)
80                 )
81             else:
82                 print(f"Warning: Malformed annotation line in {label_filename}: {annotation_line.strip()}")
83         else:
84             print(f"No annotations found in {label_filename} for image index {image_index}.")
85
86 # Add a title to the plot
87 ax.set_title(f"Image Index: {image_index} (Split: {split}) with YOLO Annotations")
88
89 # Hide axes
90 ax.axis('off')
91
92 # Display the plot
93 plt.show()
94
95 print("Finished visualizing YOLO annotations.")

```

Visualizing YOLO annotations for 5 fracture images...

**fracture** Image Index: 7 (Split: train) with YOLO Annotations

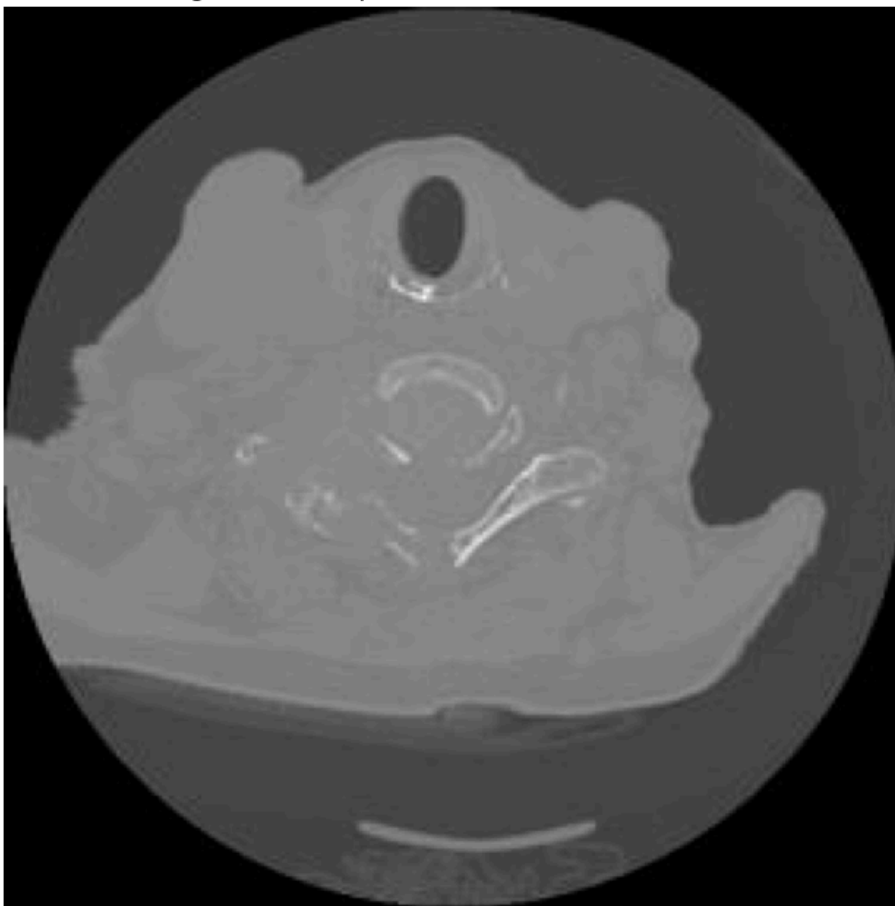
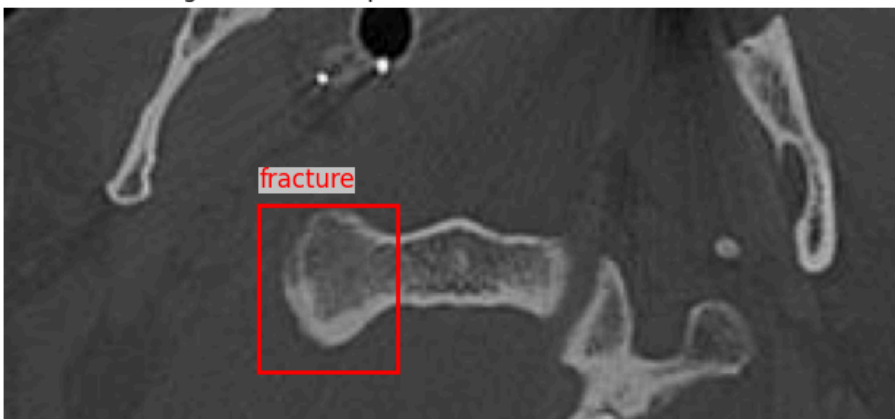


Image Index: 15 (Split: train) with YOLO Annotations



```

1 #Initialize an empty dictionary to store fracture counts for each split
2 fracture_counts = {'train': 0, 'val': 0, 'test': 0}
3
4 print("Counting fracture instances in YOLO dataset...")
5
6 #Ensure YOLO_DATA_DIR and splits are accessible from previous cells
7 #YOLO_DATA_DIR = 'yolo_dataset'
8 #splits = ['train', 'val', 'test']
9
10 #Loop through each split
11 for split in splits:
12     #Construct the full path to its 'labels' directory
13     labels_dir = os.path.join(YOLO_DATA_DIR, split, 'labels')
14
15     #Check if the labels directory exists
16     if not os.path.exists(labels_dir):
17         print(f"Warning: Labels directory not found for split '{split}' at {labels_dir}")

```

```

18         continue
19
20     #Iterate through each .txt file in the current split's labels directory
21     for filename in os.listdir(labels_dir):
22         if filename.endswith('.txt'):
23             label_filepath = os.path.join(labels_dir, filename)
24
25             #Flag to check if fracture is found in the current image
26             found_fracture = False
27
28             #Open and read the label file
29             with open(label_filepath, 'r') as f:
30                 for line in f:
31                     # 6. Check if any line in the file begins with '1' (class_id 1 for 'fracture')
32                     if line.strip().startswith('1 '):
33                         found_fracture = True
34                         break # Found a fracture, no need to check further lines in this file
35
36             #If a line starting with '1' is found, increment the count
37             if found_fracture:
38                 fracture_counts[split] += 1
39
40 #Print the fracture_counts dictionary
41 print("\nSummary of images with fracture annotations per split:")
42 for split, count in fracture_counts.items():
43     print(f" {split.capitalize()}: {count} images with fracture annotations")
44

```

Counting fracture instances in YOLO dataset...

Summary of images with fracture annotations per split:

Train: 5186 images with fracture annotations

Val: 1045 images with fracture annotations

Test: 972 images with fracture annotations

```

1 #Initialize a dictionary to store fracture bounding box counts per image for each split
2 fracture_bbox_counts_per_image = {
3     'train': [],
4     'val': [],
5     'test': []
6 }
7
8 print("Counting fracture bounding boxes per image for each split...")
9
10 #Ensure YOLO_DATA_DIR and splits are accessible from previous cells
11 YOLO_DATA_DIR = 'yolo_dataset'
12 #splits = ['train', 'val', 'test']
13
14 #Loop through each split
15 for split in splits:
16     #Construct the full path to its 'labels' directory
17     labels_dir = os.path.join(YOLO_DATA_DIR, split, 'labels')
18
19     #Check if the labels directory exists
20     if not os.path.exists(labels_dir):
21         print(f"Warning: Labels directory not found for split '{split}' at {labels_dir}")
22         continue
23
24     #Iterate through each .txt file in the current split's labels directory
25     for filename in os.listdir(labels_dir):
26         if filename.endswith('.txt'):
27             label_filepath = os.path.join(labels_dir, filename)
28
29             #Initialize a counter for bounding boxes in the current image
30             current_image_bbox_count = 0
31
32             #Open and read the label file
33             with open(label_filepath, 'r') as f:
34                 for line in f:
35                     #For each line, check if it starts with '1 ' (class_id 1 for 'fracture')

```

```

36         if line.strip().startswith('1 '):
37             #If it does, increment current_image_bbox_count
38             current_image_bbox_count += 1
39
40     #After processing all lines in the file, if current_image_bbox_count is > 0,
41     #    append this count to the corresponding list in the dictionary for the current split.
42     if current_image_bbox_count > 0:
43         fracture_bbox_counts_per_image[split].append(current_image_bbox_count)
44
45 #After processing all files for all splits, print the dictionary
46 print("\nSummary of fracture bounding box counts per image:")
47 for split, counts in fracture_bbox_counts_per_image.items():
48     print(f"    {split.capitalize()} split: {len(counts)} images with fractures, total bounding boxes: {sum(counts)}")
49     #Optional: print the list of counts for more detail
50 print(f"    Counts: {counts}")

```

Counting fracture bounding boxes per image for each split

Summary of fracture bounding box counts per image:

Train split: 10 images with fractures, total bounding boxes: 10

Val split: 10 images with fractures, total bounding boxes: 10

Test split: 10 images with fractures, total bounding boxes: 10

Counts: Train: [1, 1, 1, 1, 1, 1, 1, 1, 1, 1], Val: [1, 1, 1, 1, 1, 1, 1, 1, 1, 1], Test: [1, 1, 1, 1, 1, 1, 1, 1, 1, 1]

```

1 #Calculate descriptive statistics for fracture bounding box counts per image for each split
2 print("Calculating descriptive statistics for fracture bounding box counts:")
3 descriptive_stats = {}
4 for split, counts in fracture_bbox_counts_per_image.items():
5     if counts:
6         descriptive_stats[split] = {
7             'mean': np.mean(counts),
8             'median': np.median(counts),
9             'min': np.min(counts),
10            'max': np.max(counts)
11        }
12     else:
13         descriptive_stats[split] = {
14             'mean': 0,
15             'median': 0,
16             'min': 0,
17             'max': 0
18        }
19
20 #Print the descriptive statistics
21 for split, stats in descriptive_stats.items():
22     print(f"\nDescriptive Statistics for {split.capitalize()} Split (Fracture BBoxes per Image):")
23     for stat, value in stats.items():
24         print(f"    {stat.capitalize()}: {value:.2f}")
25

```

Calculating descriptive statistics for fracture bounding box counts:

Descriptive Statistics for Train Split (Fracture BBoxes per Image):

Mean: 1.00  
Median: 1.00  
Min: 1.00  
Max: 1.00

Descriptive Statistics for Val Split (Fracture BBoxes per Image):

Mean: 1.00  
Median: 1.00  
Min: 1.00  
Max: 1.00

Descriptive Statistics for Test Split (Fracture BBoxes per Image):

Mean: 1.00  
Median: 1.00  
Min: 1.00  
Max: 1.00



```
1 #Visualize the distribution of fracture bounding box counts per image for all elites using histograms
```