

# Code for Sam Model Development And Integration with YOLOV11 Model

**By Shehryar Khan**

```
from ultralytics import YOLO
from PIL import Image
import matplotlib.pyplot as plt

# Load the trained model from your training directory
model = YOLO("runs/detect/train/weights/best.pt") # Load the best
trained weights

# Upload an image
from google.colab import files
uploaded = files.upload() # Upload an image from your local system

# Get the filename of the uploaded image
image_path = list(uploaded.keys())[0] # This gives the uploaded file's
name

# Perform object detection
results = model(image_path)

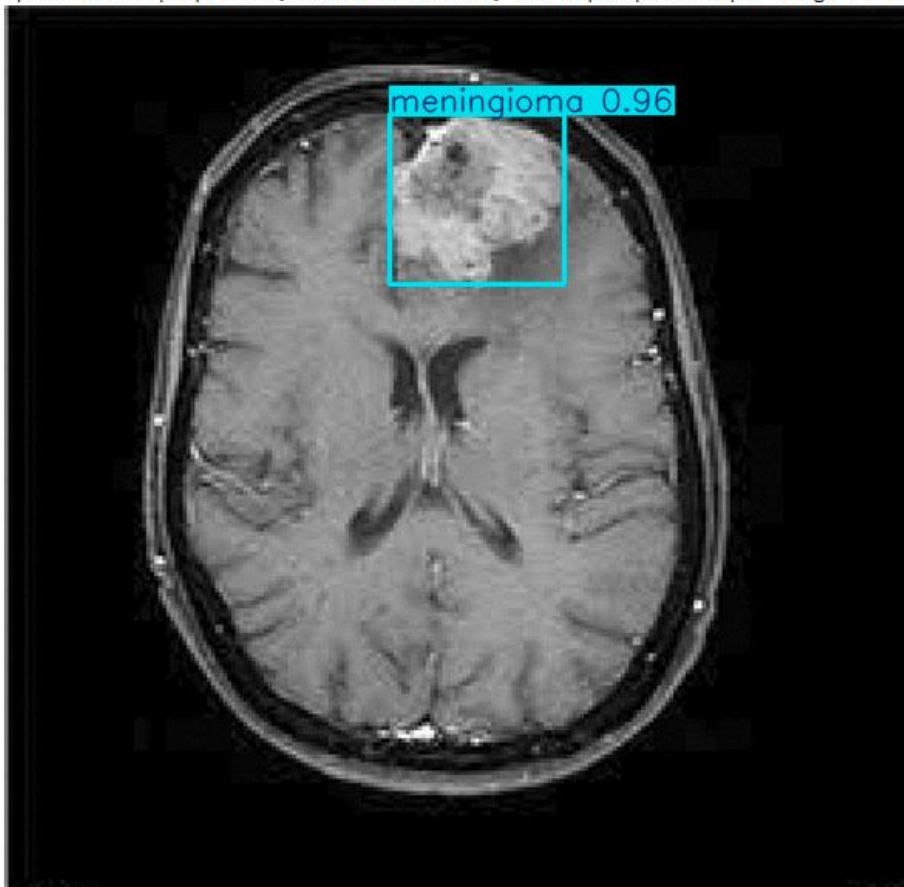
# Show the image with bounding boxes
results[0].show() # This displays the image with predictions

# Optional: Save the result to file
results[0].save(filename="detected_image.jpg")

# Show in notebook using matplotlib
img = Image.open("detected_image.jpg")
plt.imshow(img)
plt.axis('off')
plt.title("Detected Image")
plt.show()
```

Output

image 1/1 /content/41598\_2023\_41576\_Fig1\_HTML.jpg: 640x640 1 meningioma, 21.7ms  
Speed: 4.2ms preprocess, 21.7ms inference, 2.7ms postprocess per image at shape (1, 3, 640, 640)



SAM Model Integrations;

```
import torch
import numpy as np
import cv2
import matplotlib.pyplot as plt
from segment_anything import sam_model_registry, SamPredictor
from PIL import Image
from ultralytics import YOLO
```

```
# Load trained YOLO model
```

```
yolo_model = YOLO("runs/detect/train/weights/best.pt")
```

```
# Upload your image (or provide path)
```

```
from google.colab import files
```

```
uploaded = files.upload()
```

```
image_path = list(uploaded.keys())[0]
```

```
# Run detection
```

```
results = yolo_model(image_path)
```

```

detections = results[0].boxes.xyxy.cpu().numpy().astype(int) # xyxy
format (x1, y1, x2, y2)
# Download SAM weights if not already present
!wget
https://dl.fbaipublicfiles.com/segment_anything/sam_vit_b_01ec64.pth

# Load the SAM model
sam_checkpoint = "sam_vit_b_01ec64.pth"
sam = sam_model_registry["vit_b"](checkpoint=sam_checkpoint)
sam.to("cuda" if torch.cuda.is_available() else "cpu")
predictor = SamPredictor(sam)
# Load image and set it for SAM
image_bgr = cv2.imread(image_path)
image_rgb = cv2.cvtColor(image_bgr, cv2.COLOR_BGR2RGB)
predictor.set_image(image_rgb)

# Loop through YOLO bounding boxes and apply SAM
for i, box in enumerate(detections):
    input_box = np.array(box) # x1, y1, x2, y2
    masks, scores, logits = predictor.predict(box=input_box,
multimask_output=True)

    # Plot the best mask (highest confidence)
    plt.figure(figsize=(5, 5))
    plt.imshow(image_rgb)
    plt.imshow(masks[0], alpha=0.6) # semi-transparent mask
    plt.title(f"Object {i+1} (Confidence: {scores[0]:.2f})")
    plt.axis("off")
    plt.show()

```

Output



Choose files 41598\_202...g1\_HTML.jpg

• 41598\_2023\_41576\_Fig1\_HTML.jpg(image/jpeg) - 54666 bytes, last modified: 22/04/2025 - 100% done  
saving 41598\_2023\_41576\_Fig1\_HTML.jpg to 41598\_2023\_41576\_Fig1\_HTML (1).jpg

```
image 1/1 /content/41598_2023_41576_Fig1_HTML (1).jpg: 640x640 1 meningioma, 25.5ms
Speed: 8.9ms preprocess, 25.5ms inference, 3.1ms postprocess per image at shape (1, 3, 640, 640)
--2025-04-22 11:07:33-- https://dl.fbaipublicfiles.com/segment_anything/sam_vit_b_01ec64.pth
Resolving dl.fbaipublicfiles.com (dl.fbaipublicfiles.com)... 108.157.254.124, 108.157.254.121, 108.157.254.102, ...
Connecting to dl.fbaipublicfiles.com (dl.fbaipublicfiles.com)|108.157.254.124|:443... connected.
HTTP request sent, awaiting response... 200 OK
Length: 375042383 (358M) [binary/octet-stream]
Saving to: 'sam_vit_b_01ec64.pth'
```

```
sam_vit_b_01ec64.pt 100%[=====>] 357.67M 34.4MB/s in 5.3s
```

```
2025-04-22 11:07:38 (67.7 MB/s) - 'sam_vit_b_01ec64.pth' saved [375042383/375042383]
```

Object 1 (Confidence: 0.96)

