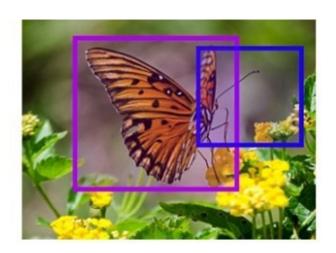


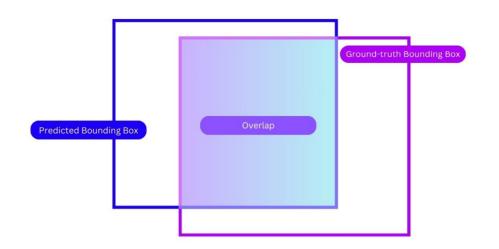
Intersect Over Union (IOU)

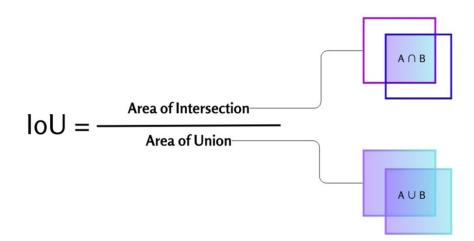






IoU = 0.971 IoU = 0.772 IoU = 0.324







IOU to determine TP and FP

True Positive

False Positive





IoU = 0.922

IoU = 0.258

The model predicted correctly

IoU Threshold = 0.5

The model predicted wrongly.



Confusion Matrix

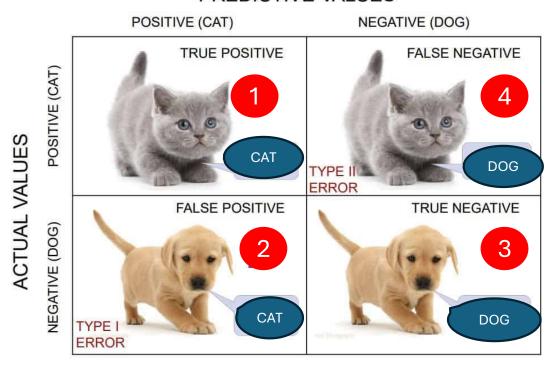
- Confusion matrix is a simple table used to measure how well a classification model is performing.
- True Positive (TP): The model correctly predicted a positive outcome i.e the actual outcome was positive.
- True Negative (TN): The model correctly predicted a negative outcome
 i.e the actual outcome was negative.
- False Positive (FP): The model incorrectly predicted a positive outcome
 i.e the actual outcome was negative. It is also known as a Type I error.
- False Negative (FN): The model incorrectly predicted a negative outcome i.e the actual outcome was positive. It is also known as a Type II error.

	Predicted Positive Predicted Negati		
Actual Positive	True Positive (TP)	False Negative (FN)	
Actual Negative	False Positive (FP)	True Negative (TN)	



Example of Confusion Matrix

PREDICTIVE VALUES



- 1. True Positive (TP): Correctly predicted a positive outcome (cat) and the actual outcome was positive (cat)
- **2. False Positive (FP):** Incorrectly predicted a positive outcome (cat) and the actual outcome was negative (dog)
- **3. True Negative (TN):** Correctly predicted a negative outcome (dog) and the actual outcome was negative (dog)
- 4. False Negative (FN): Incorrectly predicted a negative outcome (dog) and the actual outcome (cat) was positive.



Precision and Recall

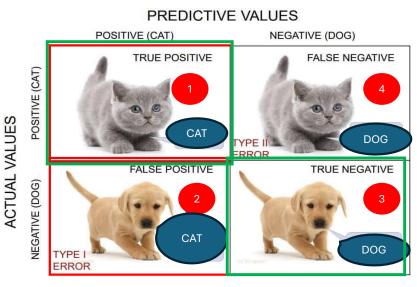
 P (Precision): The accuracy of the detected objects, indicating how many detections were correct. Refers to accuracy of the model's positive predictions.

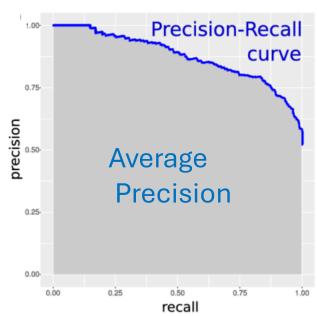
$$Precision (P) = \frac{TP}{TP + FP}$$

 R (Recall): The ability of the model to identify all relevant instances of objects in the images. (True positives are correctly identified cancerous cells, and false negatives are cancerous cells that the model missed.)

Recall (R) =
$$\frac{TP}{TP + FN}$$

 Average precision (AP) is the area under the precision-recall curve (PR curve). A higher AP indicates better performance







Mean Average Precision (mAP)

- Mean Average Precision
 - The average of AP calculated across all classes in an object detection task
- mAP50 (easy detection):
 - Mean average precision calculated at an intersection over union (IoU) threshold of 0.50 across all classes.
- mAP50-95 (more accurate)
 - The average of the mean average precision calculated at varying IoU thresholds, ranging from 0.50 to 0.95 across all classes.
 - The thresholds are typically incremented by 0.05 and mAP is calculated for each IoU.

$$\mathsf{mAP} = \frac{1}{\mathsf{N}} \sum_{i=1}^{\mathsf{N}} \mathsf{AP}_{\mathsf{IoU}(\mathsf{k})}$$

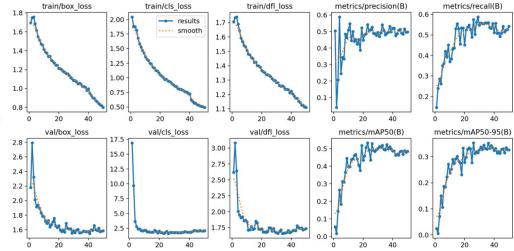
 $\mathsf{AP}_{IoU(k)}$ is the Average Precision at the (k)-th IoU threshold and N is the number of class.



Precision vs Recall

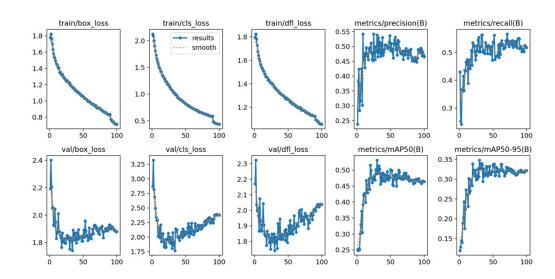
• Epoch50, batch16, yolov8l.pt

Class	Images	Instances	Box(P	R	mAP50	mAP50-95):
all	244	726	0.511	0.595	0.521	0.35
corner-cast	190	399	0.207	0.206	0.0934	0.0386
roof	154	202	0.654	0.698	0.634	0.394
shipping-container	28	125	0.672	0.88	0.836	0.616



• Epoch100, batch16, yolov8l.pt

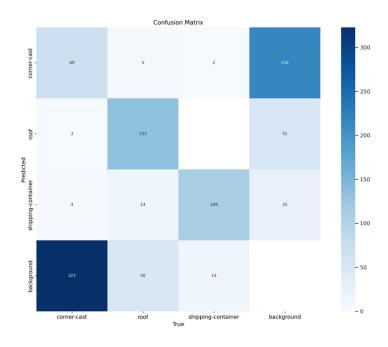
Class	Images	Instances	Box(P	R	mAP50	mAP50-95)
all	244	726	0.539	0.502	0.531	0.348
corner-cast	190	399	0.204	0.138	0.119	0.0467
roof	154	202	0.761	0.504	0.664	0.413
hipping-container	28	125	0.651	0.864	0.809	0.584

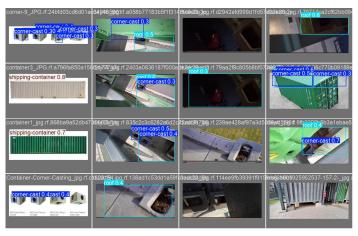




Confusion Matrix

• Epoch50, batch16, yolov8l.pt





• Epoch100, batch16, yolov8l.pt

