

MedCLIP Zero-Shot Disease Classification  
6-class setting: CheXpert-5 + Others

Tag / image variant : clahe

Preds CSV : D:\MedVLMBench\phase1\results\medclip\_chexpert\_sanity\_clahe\_preds.csv

Classes:

- CheXpert-5: Atelectasis, Cardiomegaly, Consolidation, Edema, Pleural Effusion
- Others : any chexpert\_label outside these five

How predictions are made:

- MedCLIP outputs probabilities only for the 5 CheXpert diseases.
- We take argmax over these 5 scores to get the predicted class.
- MedCLIP never predicts 'Others' (this is by design).

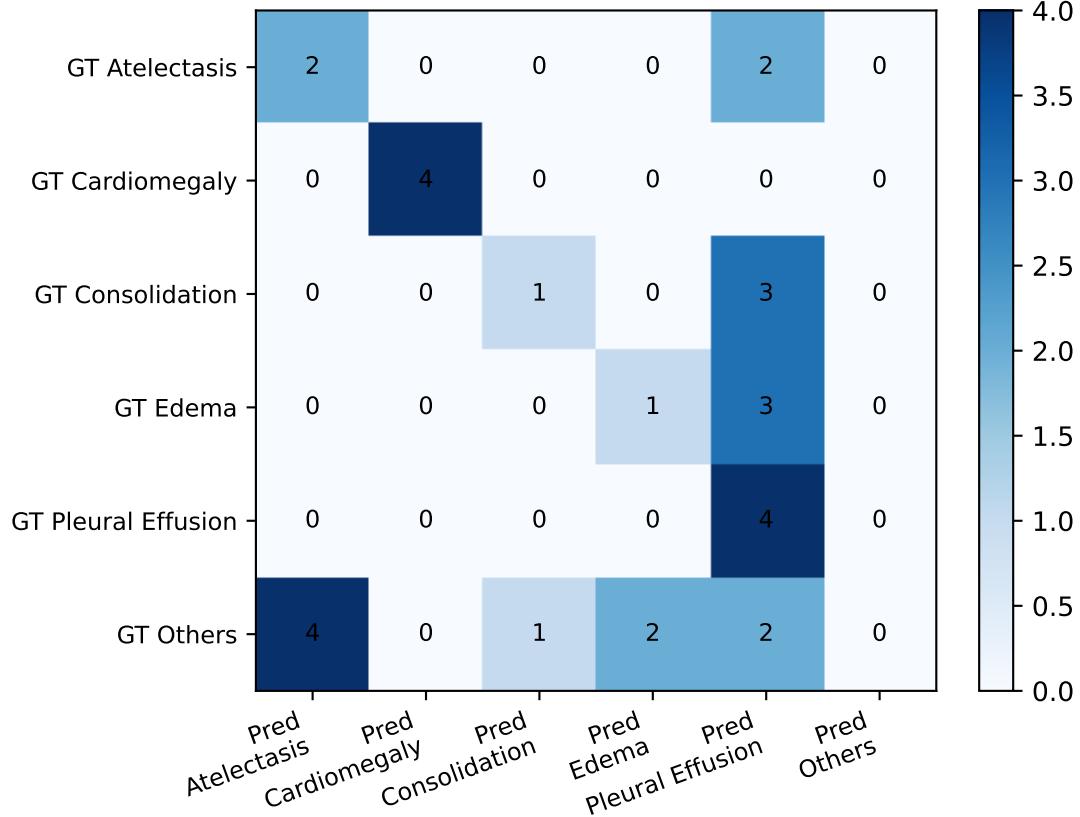
Interpretation:

- If GT = Others, any prediction must be one of the 5 diseases.  
This reveals how often MedCLIP mis-classifies non-CheXpert pathologies into the limited disease set.

Overall performance on this subset:

- Samples used : 29
- Overall accuracy: 0.414
- Macro F1 : 0.411

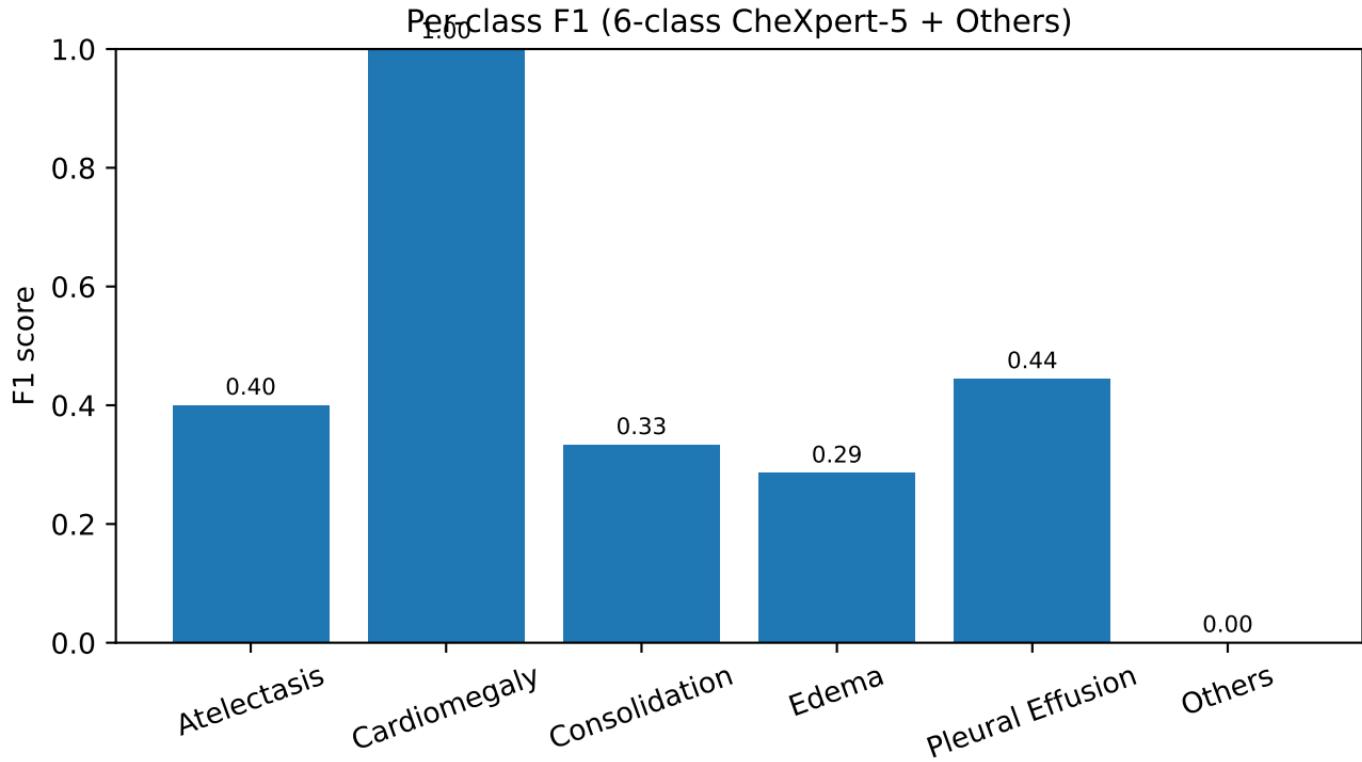
6-class confusion matrix (rows = GT, cols = Pred)



Confusion matrix for the 6-class task (CheXpert-5 + Others).

Diagonal cells show correct predictions. Off-diagonal cells show confusions between specific classes.

In particular, the GT 'Others' row highlights how non-CheXpert conditions are forced into the 5 disease classes.



$F1 = 2 * (\text{Precision} * \text{Recall}) / (\text{Precision} + \text{Recall})$ .

For 'Others', F1 shows how well MedCLIP avoids over-assigning non-CheXpert pathologies to the 5 target diseases.

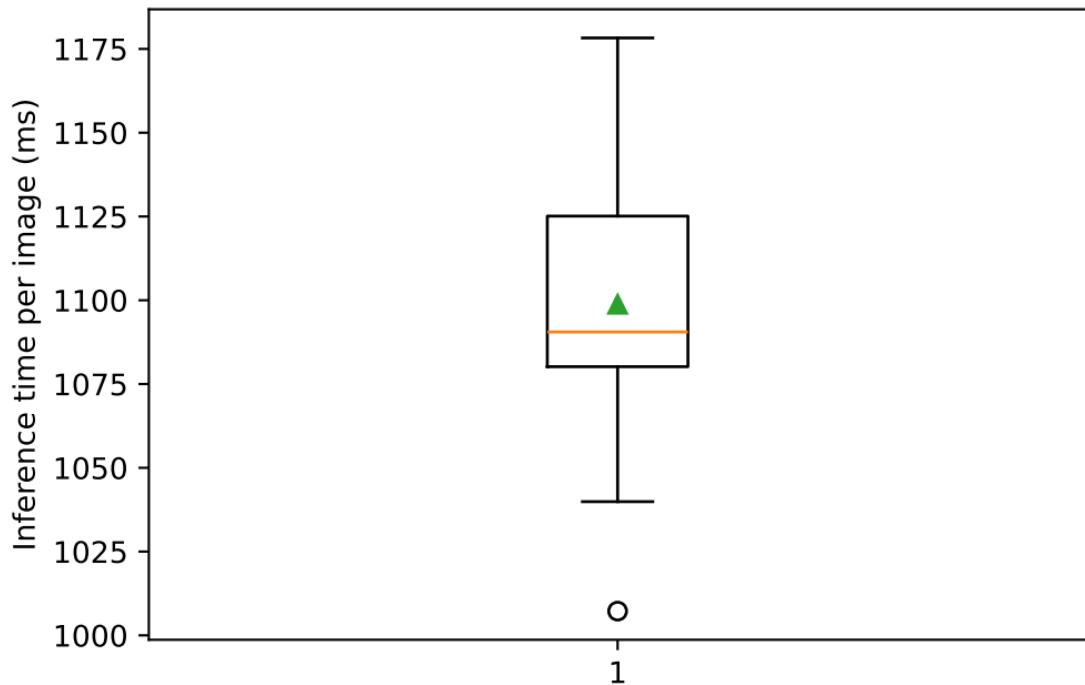
## Per-class metrics (6-class CheXpert-5 + Others)

Class	Support	Accuracy	Precision	Recall	F1
Atelectasis	4	0.500	0.333	0.500	0.400
Cardiomegaly	4	1.000	1.000	1.000	1.000
Consolidation	4	0.250	0.500	0.250	0.333
Edema	4	0.250	0.333	0.250	0.286
Pleural Effusion	4	1.000	0.286	1.000	0.444
Others	9	0.000	0.000	0.000	0.000

Support = number of ground-truth samples for each class.

Accuracy here is class-wise: fraction of samples for that class that are correctly predicted.

### MedCLIP per-image inference latency (same CSV)



Boxplot summarizing median, interquartile range, mean and outliers for per-image inference time. Useful when comparing different image variants (raw vs CLAHE vs Gauss+CLAHE).