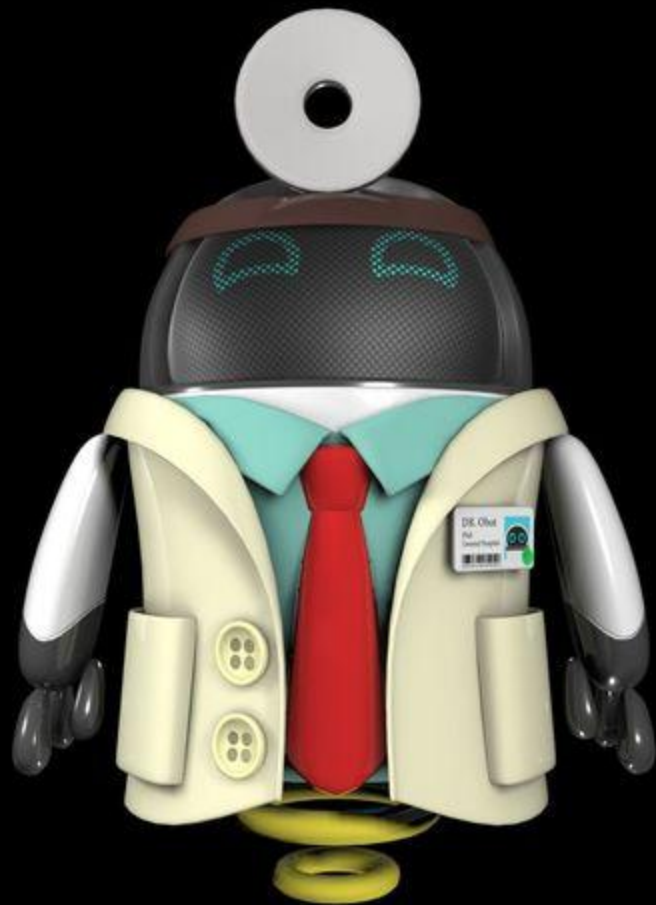
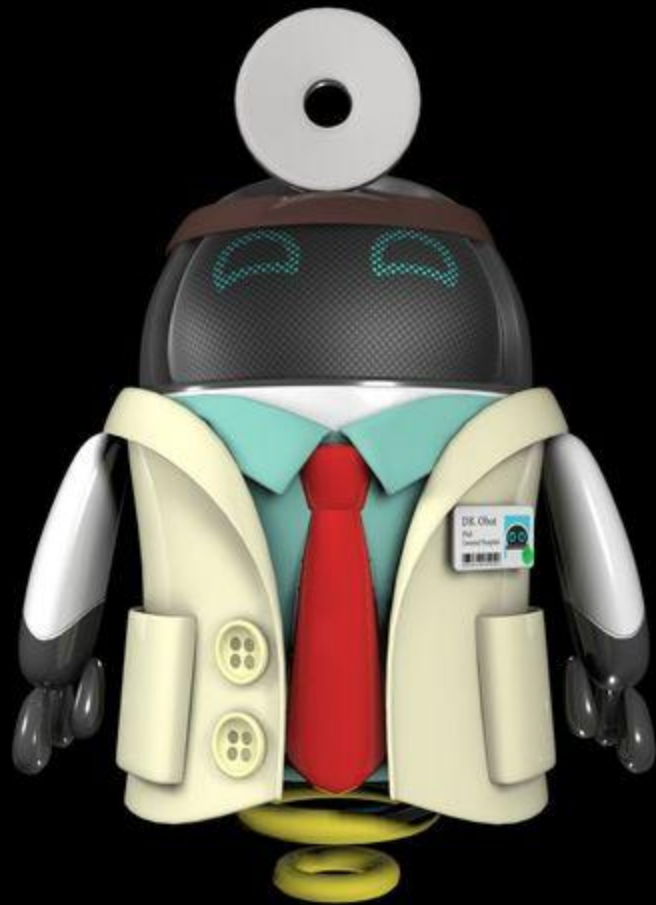
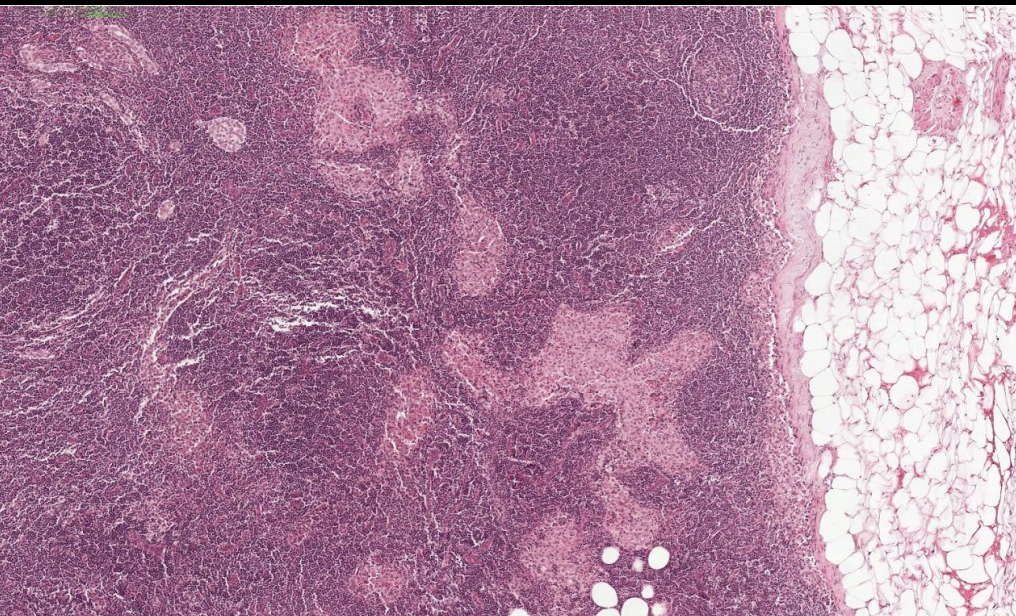
The logo for the children's book series "Where's Waldo?" is displayed. It features the words "WHERE'S" in blue and "WALDO?" in red, both in a bold, stylized font with thick outlines. The text is set against a white background with a decorative, hand-drawn border.

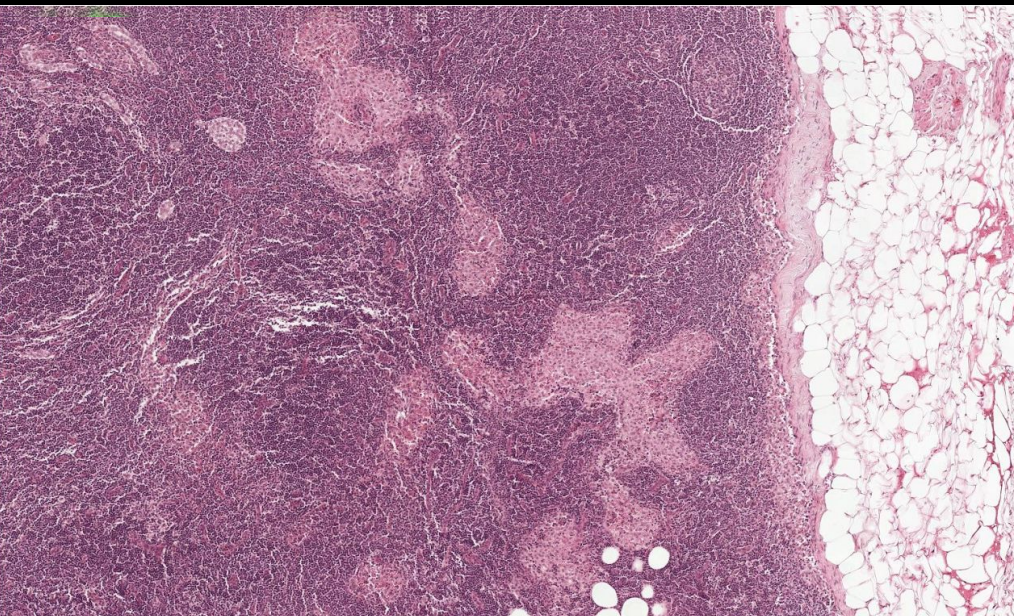
Medical Anomaly Edition



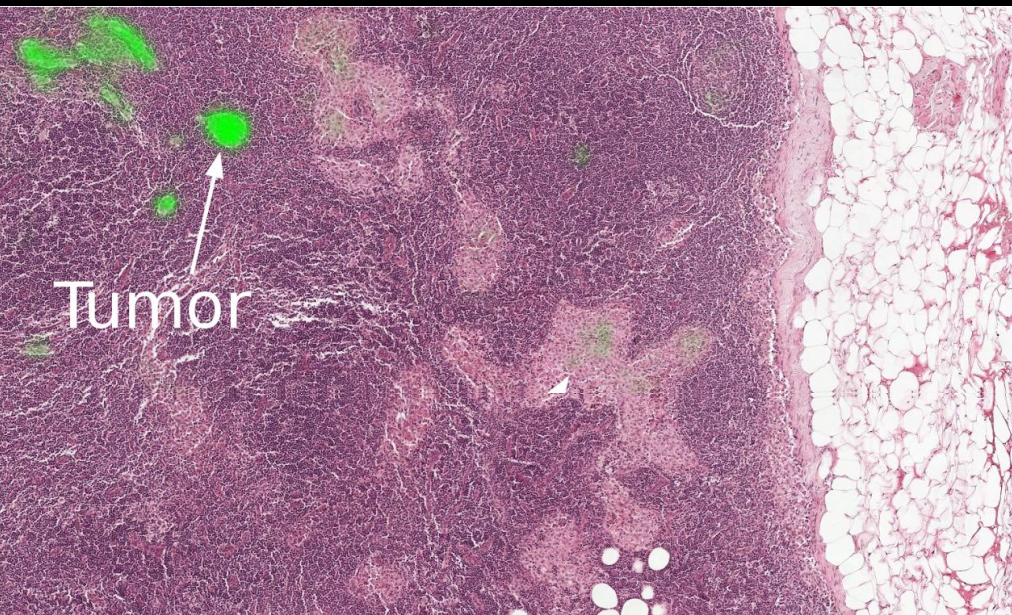
ft. GANs



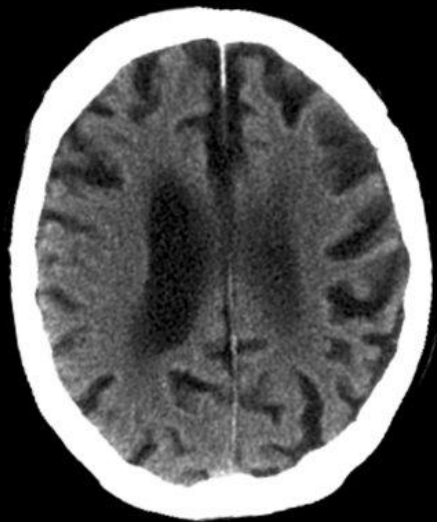




- Blastoma
- Cystic fibrosis
- Appendicitis
- Pneumonia
- Tumor



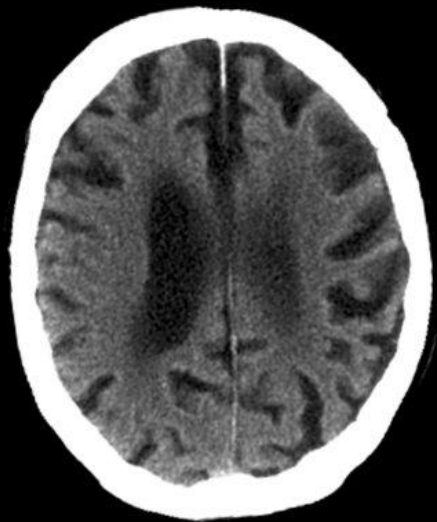
- ✗ Blastoma
- ✗ Cystic fibrosis
- ✗ Appendicitis
- ✗ Pneumonia
- ✓ Tumor



- Blastoma
- Cystic fibrosis
- Appendicitis
- Pneumonia
- Tumor

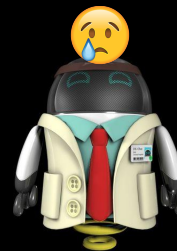


- ☒ Blastoma
- ☒ Cystic fibrosis
- ☒ Appendicitis
- ☒ Pneumonia
- ☒ Tumor



Alzheimer's

- ✗ Blastoma
- ✗ Cystic fibrosis
- ✗ Appendicitis
- ✗ Pneumonia
- ✗ Tumor



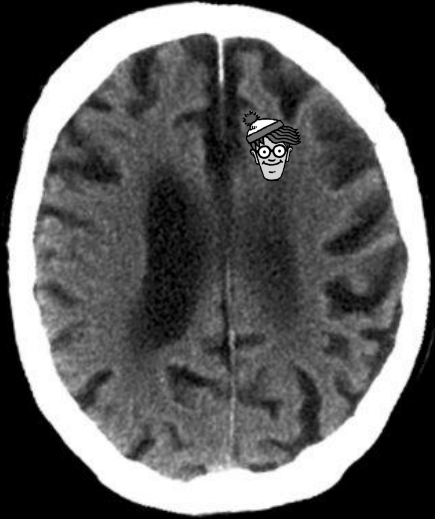


Alzheimer's

- ☐ Blastoma
- ☐ Cystic fibrosis
- ☐ Appendicitis
- ☐ Pneumonia
- ☐ Tumor
- ☒ Alzheimer's

You only find what you're looking for

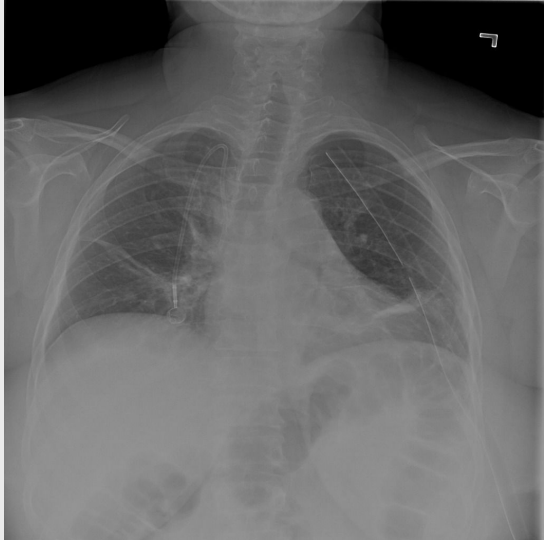
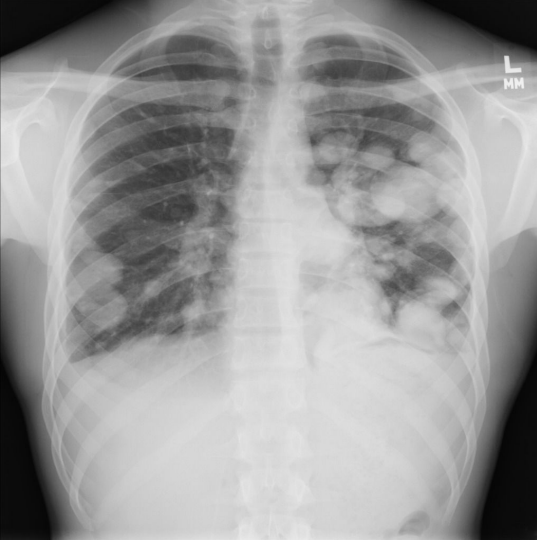
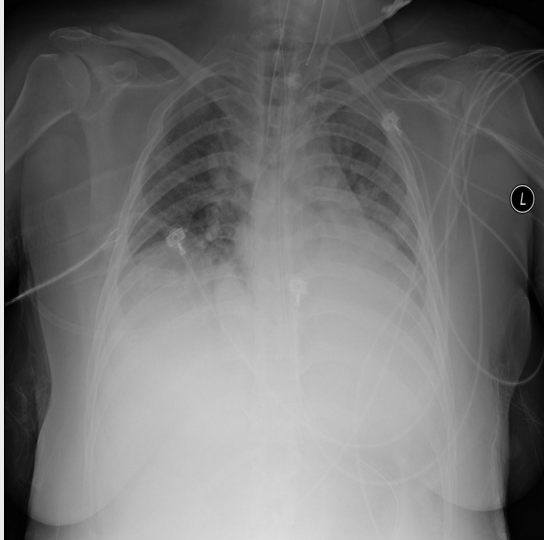


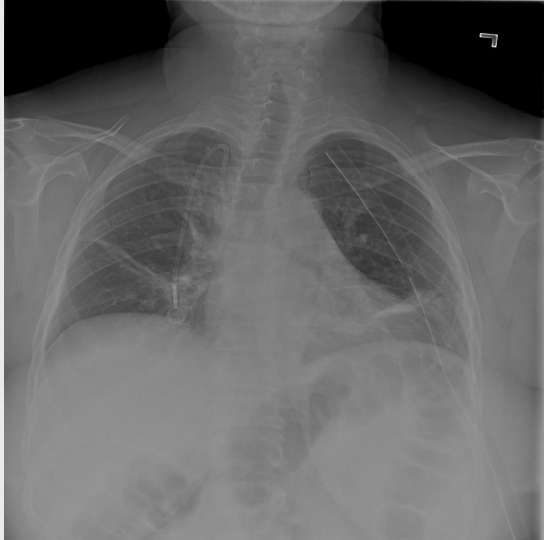
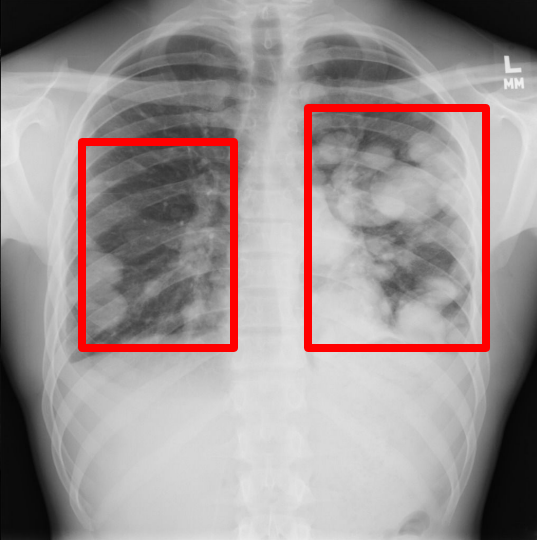
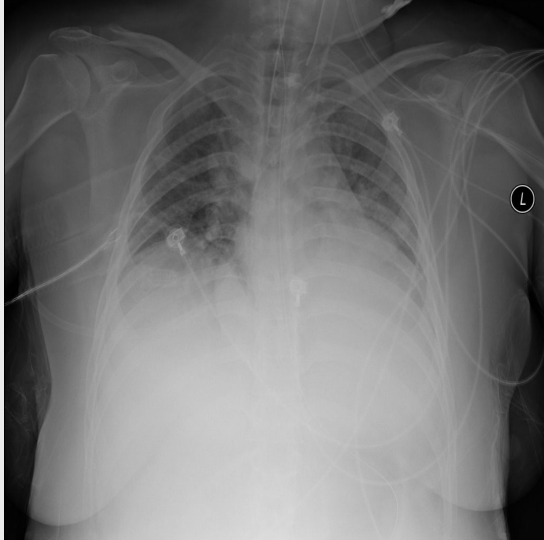


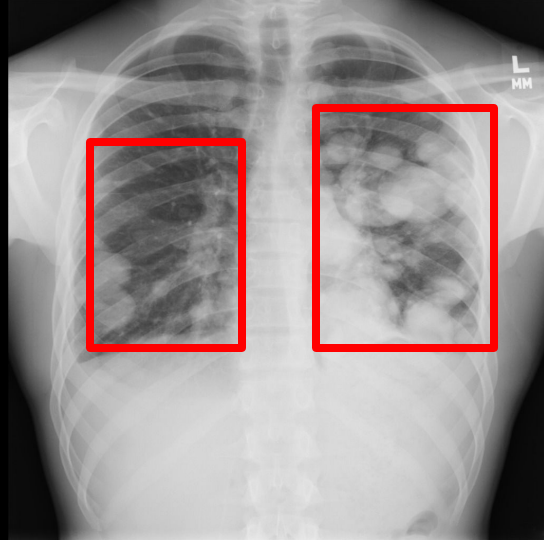
- ✗ Blastoma
- ✗ Cystic fibrosis
- ✗ Appendicitis
- ✗ Pneumonia
- ✗ Tumor
- ✗ Alzheimer's











ZILC





D: Detective



R: Real Data



G: Generator (Forger)





[[[232 234 234]
[232 236 237]
[228 231 235]

...

[221 226 227]
[214 224 224]
[209 222 220]]

[[231 233 233]
[229 233 234]
[227 231 236]

...

[217 224 227]
[216 226 226]
[211 222 220]]



Color: Pink
Shape: Circle
Radius: 12cm
Icing?: Yes
Hole?: Yes
Hole Size: 2.5cm
Sprinkles?: Yes
Sprinkles Size: 1cm x 0.2cm



Color: (200, 130, 150)
Shape: 0
Radius: 12cm
Icing?: 1
Hole?: 1
Hole Size: 2.5cm
Sprinkles?: 1
Sprinkles Size: 1cm x 0.2cm



20
78
96
...
83
92
48
63



R: Real Data



D: Detective



G: Generator (Forger)





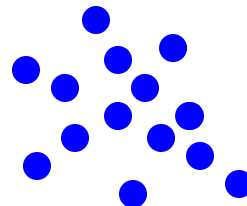
R: Real Data

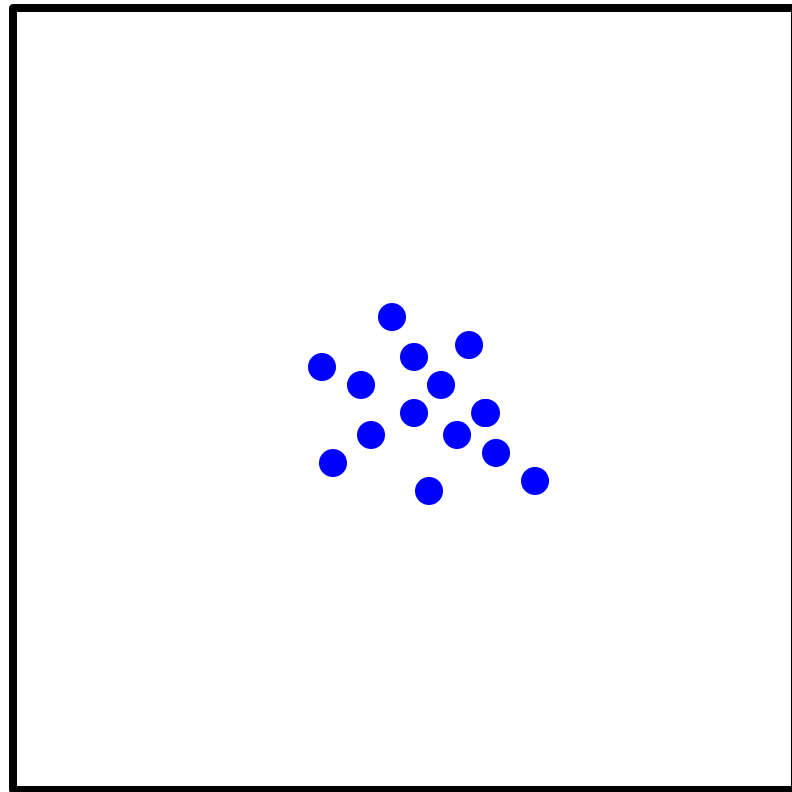
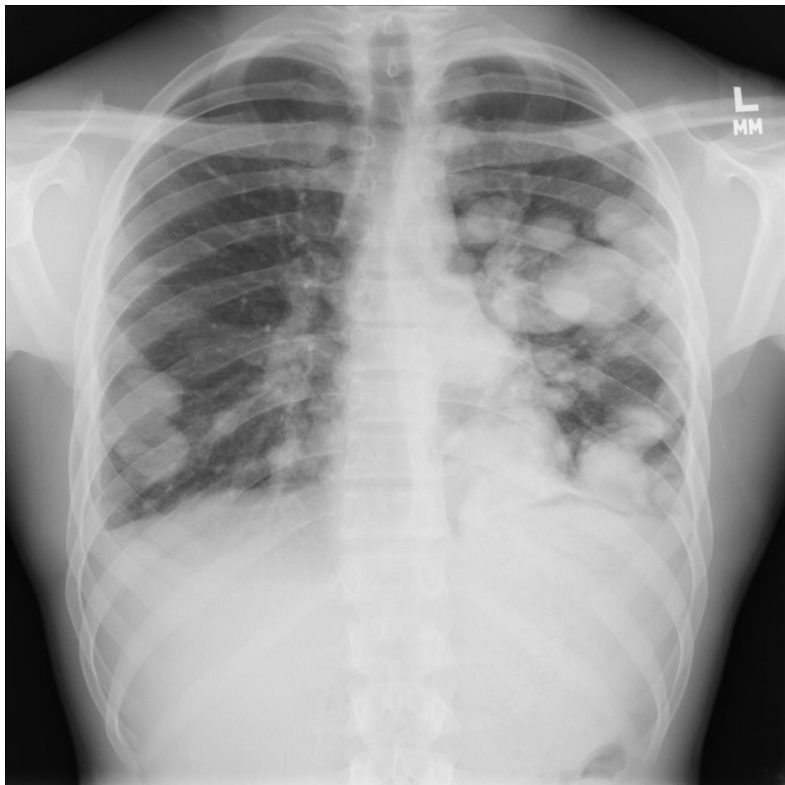


D: Detective

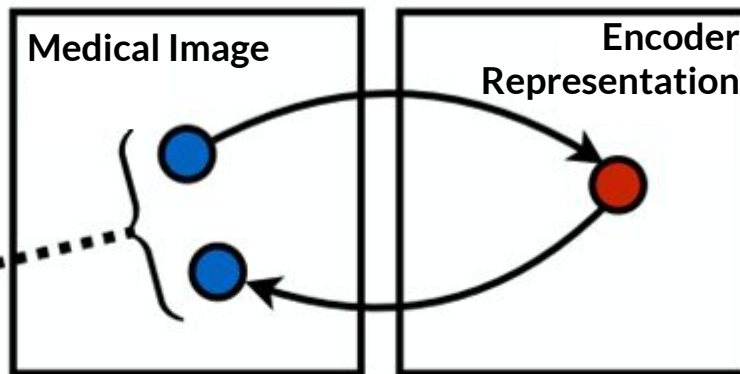


G: Generator (Forger)





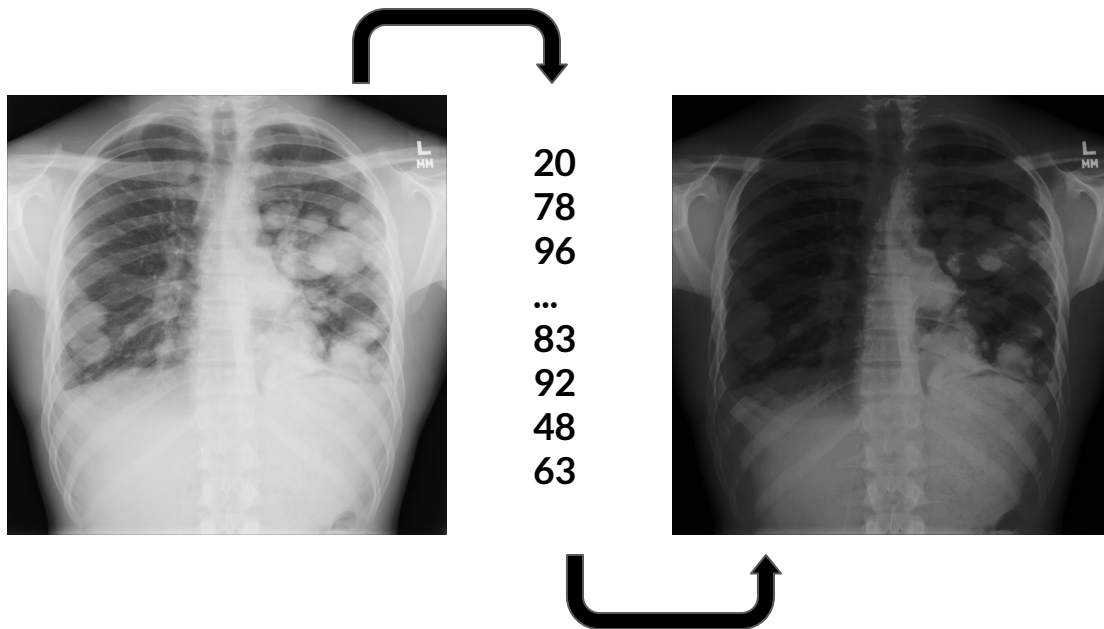
cycle-consistency
loss



Reconstruction Loss

Encoder Matching

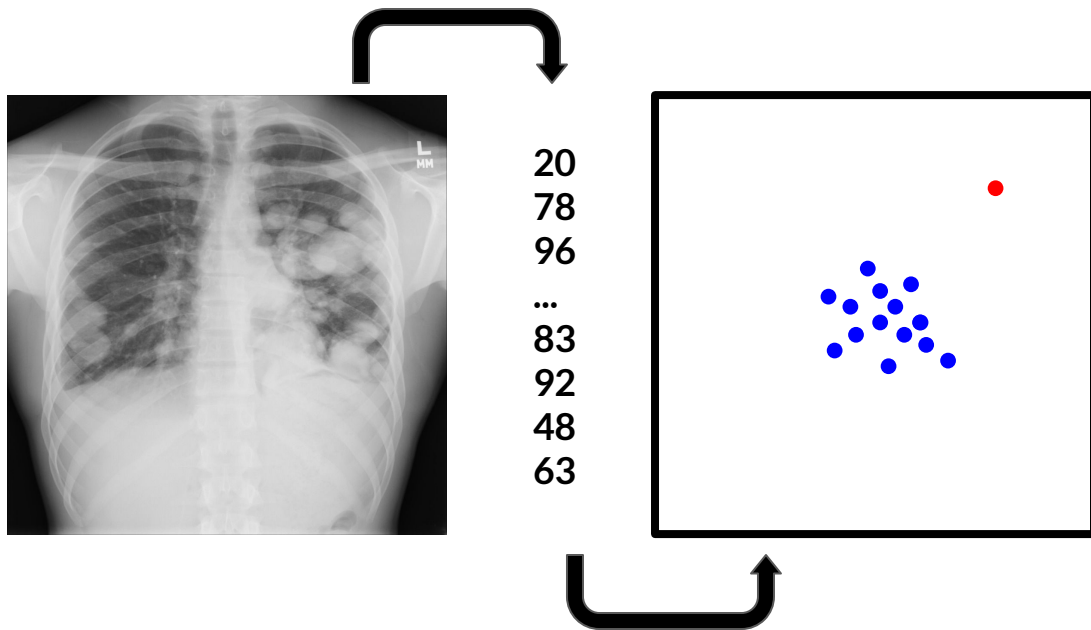
Discrimination



Reconstruction Loss

Encoder Matching

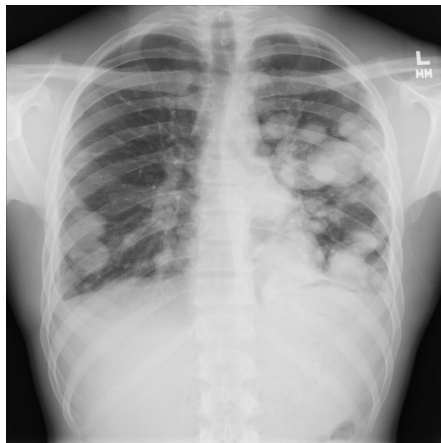
Discrimination



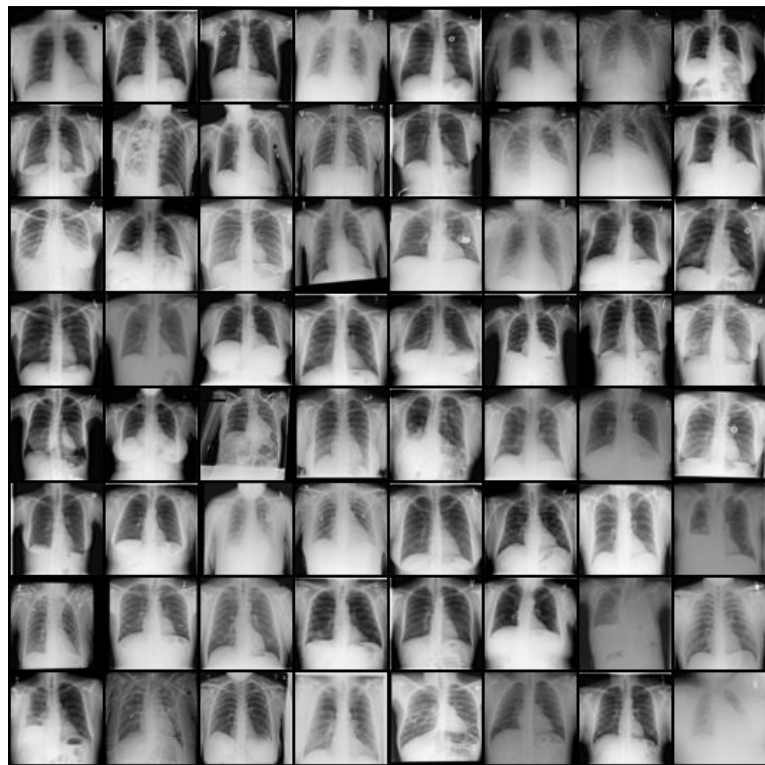
Reconstruction Loss

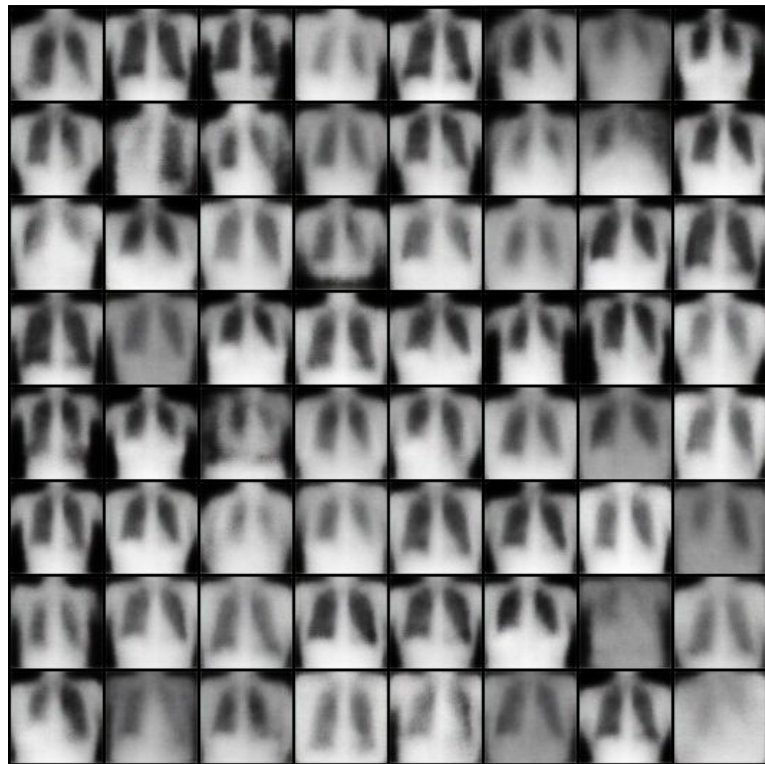
Encoder Matching

Discrimination



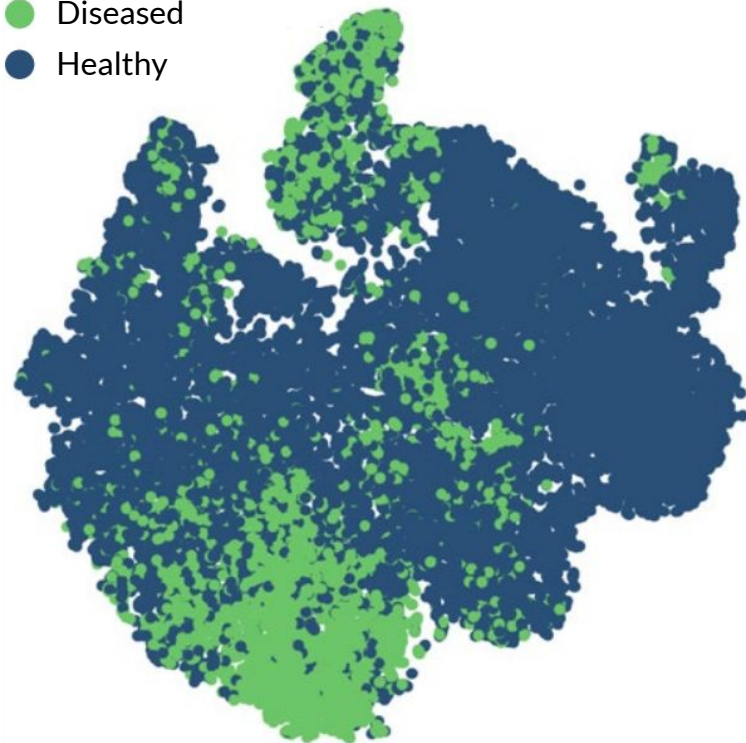
0.8

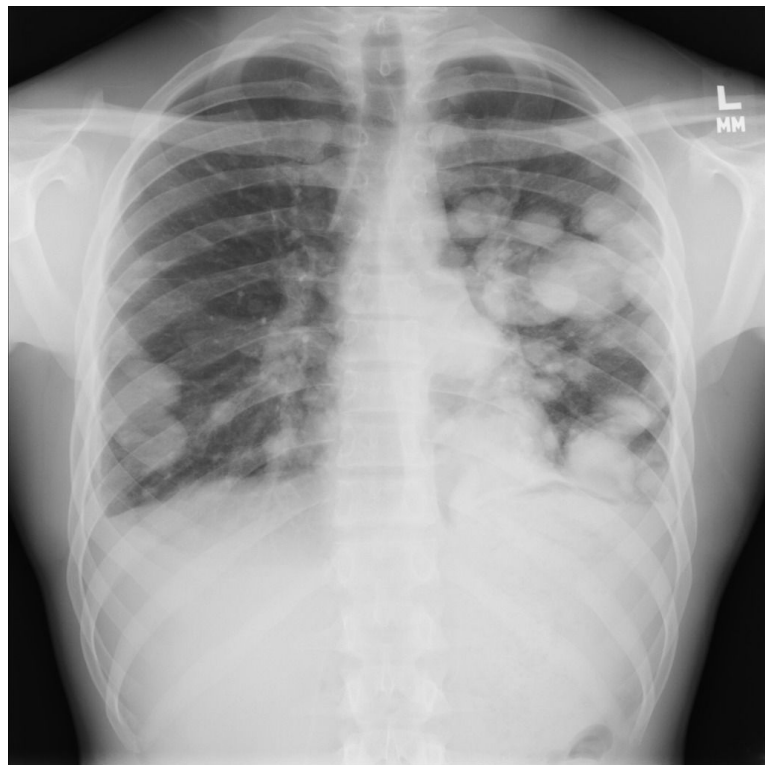


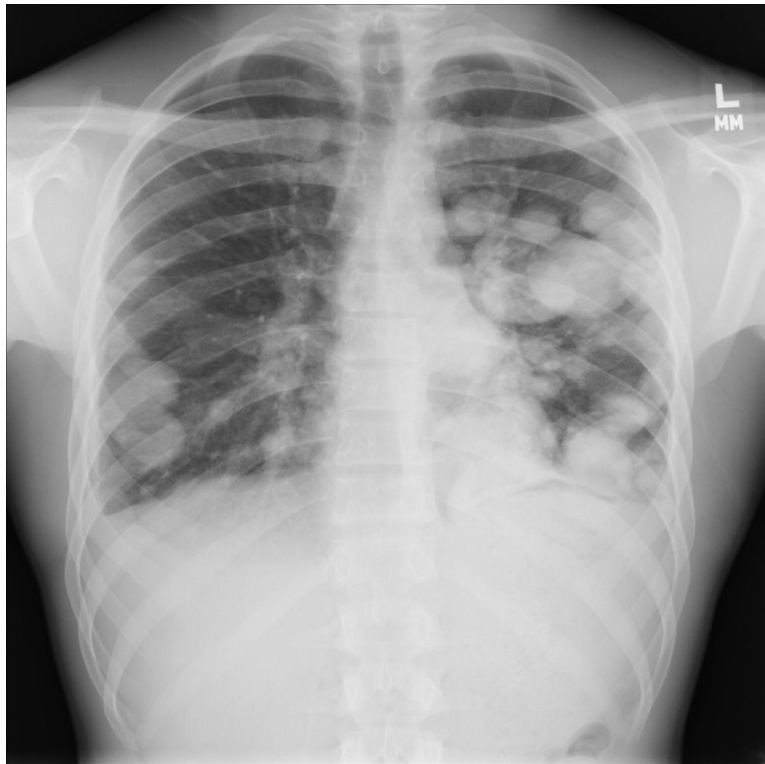


● Diseased

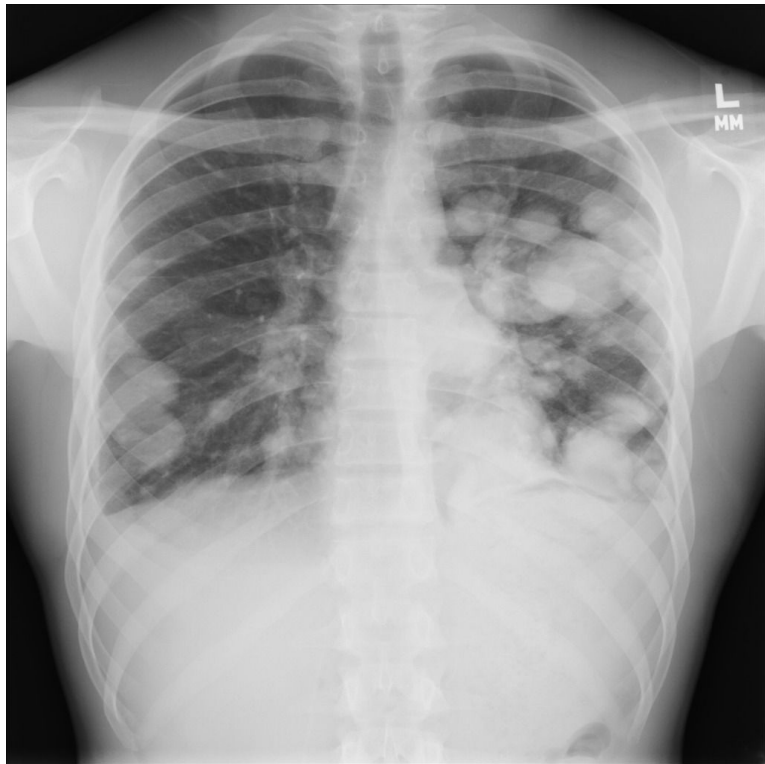
● Healthy





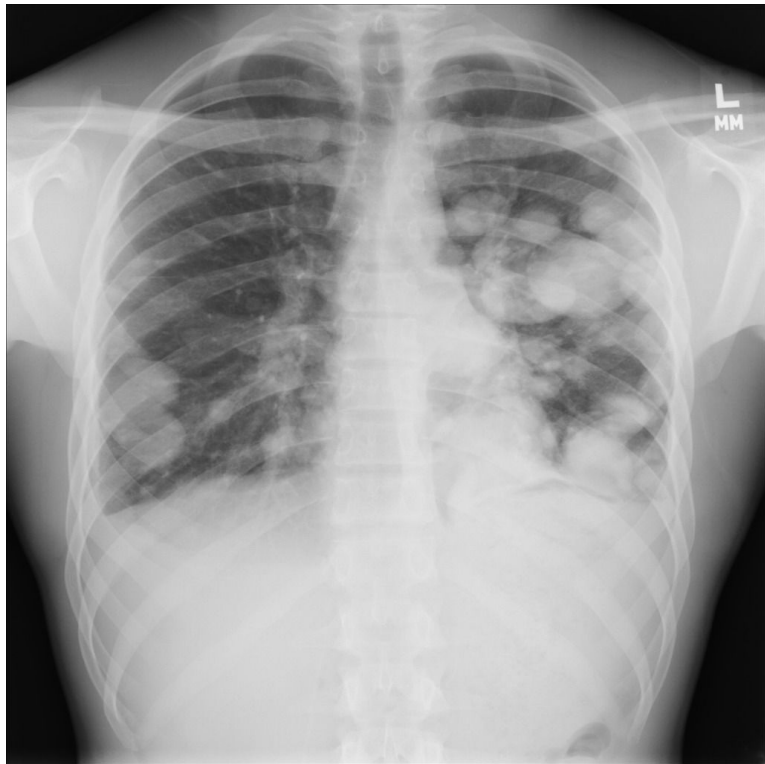


```
def evaluate(image):  
    reconstruction_loss = image_diff(image, reconstruct(image))  
    embedding           = in_dist(healthy_embed, encode(image))  
    discriminator       = netD(image)  
  
    total_loss = reconstruction_loss + embedding + discriminator  
  
    threshold = 0.30  
  
    return total_loss > threshold
```



Anomaly

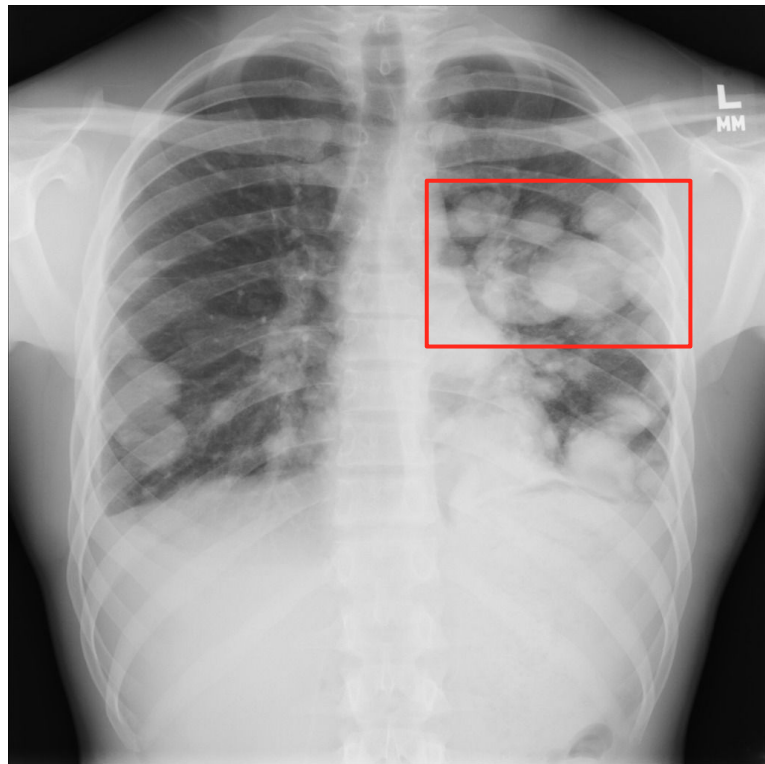




Anomaly



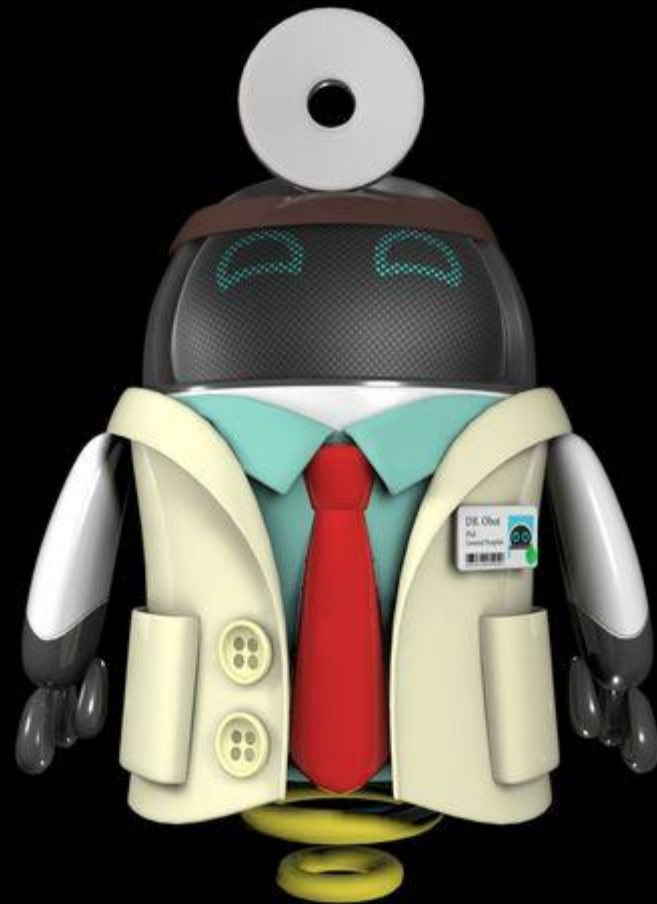
```
def localize_anomaly(image):  
    reconstruction = reconstruct_image(image)  
  
    image_similarity = similar(image, reconstruction)  
    anomaly_location = local_min(image_similarity)  
  
    return anomaly_location
```

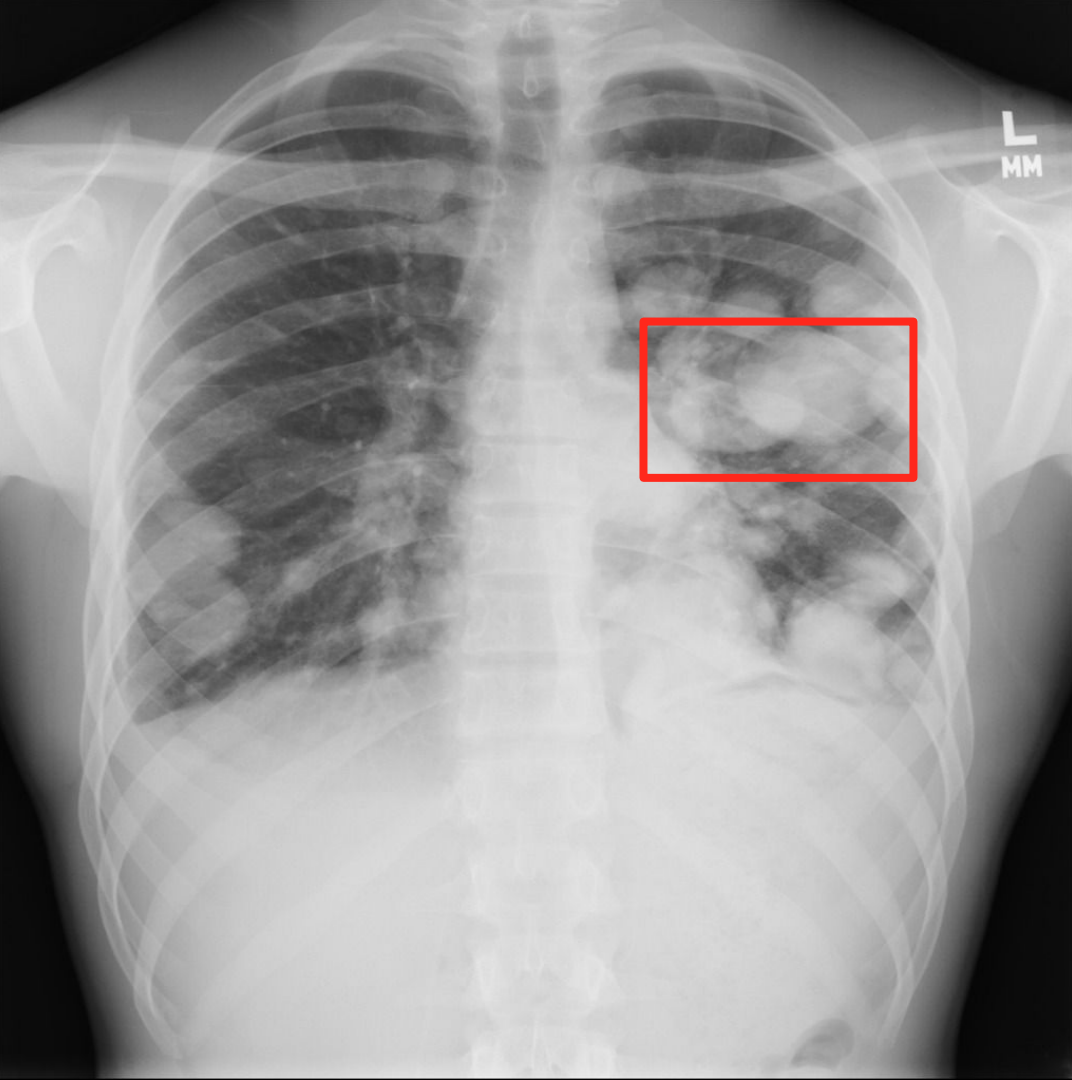



Anomaly

ZILC







liamhinzman.com

liamhinzman@gmail.com

