

# Creative Chest X-Ray Augmentation Using Cycle GAN

Converting chest X-rays of healthy patients to X-rays with pneumonia

## Task Description & Use Cases

Pneumonia is a common disease of the lung, caused by either bacteria or viruses. (Wiki 2019a)

The goal of this project is to write a tool, which will take an X-Ray image of a person without a disease and alter a marked region to show a diseased lung instead.

Input:



X-Ray of a healthy person with a selected area

Output:



X-Ray of same person with Pneumonia in the marked area

Use Cases:

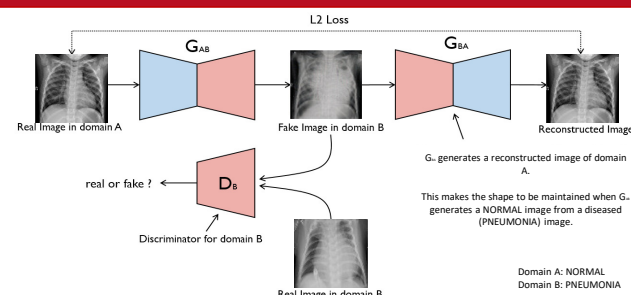
- Simulate diseased lungs for training purposes
- Predict edge cases (how would pneumonia look like in a region no information is known about)
- Training classification models on adversarial examples can prevent attacks
- Work as a POC for similar tasks

## Data Preprocessing

Preprocessing Steps:

- Combine train, validation and test set as no differentiation is needed for the GAN setup
- Remove images smaller than a certain threshold in width and in height
- Remove letters in the image by turning pixels with a high white value to black

## Modeling – Cycle GAN

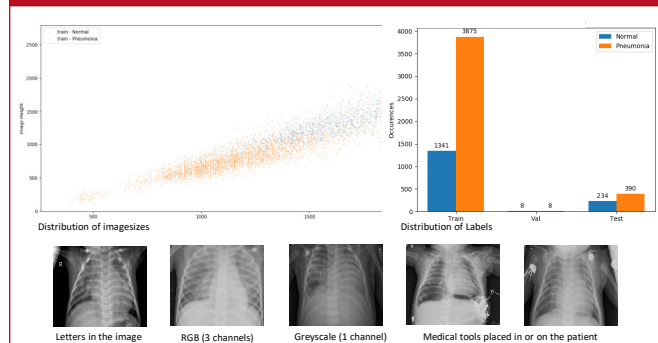


- Transfers an image from one domain into another and back
- Trained over complete network
- Generator: Encoder-Decoder Architecture
- Discriminator: CNN Architecture
- Loss: Cycle consistency loss

## References

- Jason Brownlee, PhD, 2019, <https://machinelearningmastery.com/cyclegan-tutorial-with-keras>
- Paul Mooney, 2018, <https://www.kaggle.com/paultimothymooney/chest-xray-pneumonia>
- Yunjey Choi, 2017, <https://github.com/yunjey/mnist-svhn-transfer> (edited)

## Dataset Exploration

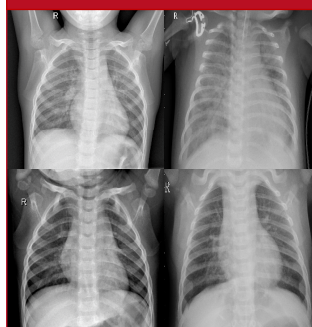


Results:

- Class imbalance towards PNEUMONIA
- Images of healthy lungs tend to be bigger
- Blob of low-resolution images
- Some images show problematic distinguishing factors

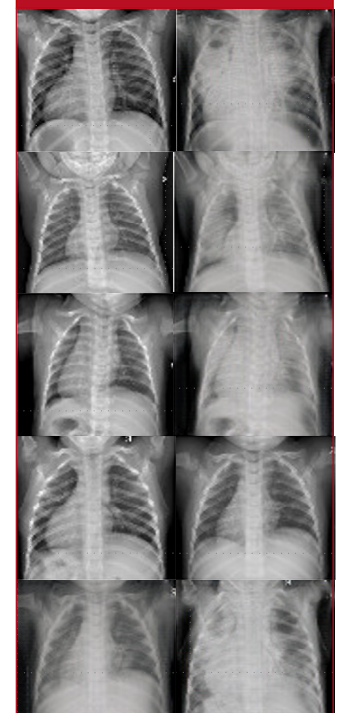
## DATASET

### NORMAL    PNEUMONIA



## GENERATED

### NORMAL    PNEUMONIA



## Evaluation & Results

Evaluation:

- Cycle GAN successfully generates diseased lungs from healthy samples
- Objective analysis shows that only very diseased lungs are generated
- No inclusion of mask because training data was missing
- -> no region for pneumonia can be marked in the generation process.