

CSC 4080: Artificial Intelligence in

Medical Imaging and Health

Homework Set 1

Due date: Feb. 20, 2022

Short Answer questions:

1. Explain the image acquisition process for digital CT Imaging in math. (10%)

For a 2D attenuation function $\mu(x, y)$, describing the X-ray attenuation per unit volume in some slice z in the human body, the Radon transform is given by all line integrals through this function:

$$R(\rho, \varphi)[\mu(x, y)] = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} \mu(x, y) \delta(\rho - x \cos \varphi - y \sin \varphi) dx dy$$

where δ is the Dirac-delta function and s is the line.

When light spreads in the homogeneous (均匀) medium, its intensity attenuation model is: $I = I_0 e^{-\int_L \mu(x) dx}$, where μ is the attenuation coefficient and x is the length. The Radon Transform is $\ln(\frac{I}{I_0}) = - \int_L \mu(x) dx = R_L \mu$.

Two variables (ρ and φ) in radon transform, assume that φ is fixed, only ρ changes. We apply Fourier transformation to $R_L \mu$:

$$\begin{aligned} F_{\rho} R_{\mu}(\rho, \varphi) &= \int_{-\infty}^{\infty} e^{-2\pi i \rho} R_{\mu}(\rho, \varphi) d\rho \\ &= \int_{-\infty}^{\infty} e^{-2\pi i \rho} \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} \mu(x_1, x_2) \delta(\rho - x_1 \cos \varphi - x_2 \sin \varphi) dx_1 dx_2 d\rho \\ &= \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} \mu(x_1, x_2) \int_{-\infty}^{\infty} e^{-2\pi i \rho} \delta(\rho - x_1 \cos \varphi - x_2 \sin \varphi) d\rho dx_1 dx_2 \\ &= \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} \mu(x_1, x_2) e^{-2\pi i (x_1 \cos \varphi + x_2 \sin \varphi)} = F\mu(\xi_1, \xi_2) \end{aligned}$$

Take the inverse of Fourier transform, μ can be calculated. Then the information of the thickness of organs is found. The result is:

$$\begin{aligned} \mu(\xi_1, \xi_2) &= \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} F_{\rho} R_{\mu}(\rho, \varphi) e^{-2\pi i (x_1 \cos \varphi + x_2 \sin \varphi)} dx_1 dx_2 \\ &= \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} \ln\left(\frac{I(\rho, \varphi)}{I_0(\rho, \varphi)}\right) e^{-2\pi i (x_1 \cos \varphi + x_2 \sin \varphi)} e^{-2\pi i \rho} d\rho dx_1 dx_2 \end{aligned}$$

2. What is the purpose of filtering with a Hamming window filter when reconstructing CT images? (10%)

Combined with High pass filters, to reduce the amplification of high-frequencies. It is also a low pass filter, which presents a high degree of smoothing.

3. What is the advantage and disadvantages of X-ray, CT, ultrasound, and MRI? (10%)

Advantages of X-ray: It provides a lucid picture of humans' bones, and the thickness and the shape of organs. It's cheap.

Shortcomings for X-ray: It cannot view the 3D appearance of the human body. The radiation damages humans' health. Short at distinguishing organs with similar density and thickness and adjacent positions.

Advantages of CT: Form images of non-living objects. Create an image of one or more slices through the body.

Shortcomings of CT: Time consuming, expensive and a great amount of radiation.

Advantages of ultrasound: fast and timely, no pain and danger, non-invasive. It has been popularized in clinical applications and it is an important part of medical imaging. Because the equipment is not as expensive as CT or MRI equipment, US diagnosis can obtain arbitrary cross-sectional images of organs, and can also observe the activity of moving organs. It's cheap and convenient.

Shortcomings of ultrasound: the contrast resolution and spatial resolution of the image are not as high as that of CT or MRI.

Advantages of MRI: Examination of nervous system, head, cervical spine (颈椎), thoracic spine, lumbar spine, limbs and other parts has unique advantages. The biggest advantage is that it can not only display diseased tissues, but also reflect the physiological and biochemical information of living tissue functions and metabolic processes.

Shortcomings of MRI: the contrast resolution and spatial resolution of the image are not as high as that of CT or MRI.

Calculation problems:

1. Get the derivatives for the sigmoid function $\sigma(x) = \frac{1}{1+\exp(-x)}$ and softmax function $z_k = \text{softmax}(x_k) = \frac{\exp(x_k)}{\sum_{i=1}^K \exp(x_i)}$ (you should write down the explicit matrix form different from the slides). Talk about the similarities of these two functions and their applications, respectively. (10%)

$$\sigma'(x) = \begin{bmatrix} \frac{d(\frac{1}{1+\exp(-x_1)})}{dx_1} & \frac{d(\frac{1}{1+\exp(-x_1)})}{dx_2} & \dots & \frac{d(\frac{1}{1+\exp(-x_1)})}{dx_n} \\ \frac{d(\frac{1}{1+\exp(-x_2)})}{dx_1} & \frac{d(\frac{1}{1+\exp(-x_2)})}{dx_2} & \dots & \frac{d(\frac{1}{1+\exp(-x_2)})}{dx_n} \\ \dots & \dots & \dots & \dots \\ \frac{d(\frac{1}{1+\exp(-x_n)})}{dx_1} & \frac{d(\frac{1}{1+\exp(-x_n)})}{dx_2} & \dots & \frac{d(\frac{1}{1+\exp(-x_n)})}{dx_n} \end{bmatrix} = \begin{bmatrix} \frac{\exp(-x_1)}{(1+\exp(-x_1))^2} & 0 & \dots & 0 \\ 0 & \frac{\exp(-x_2)}{(1+\exp(-x_2))^2} & \dots & 0 \\ \dots & \dots & \dots & \dots \\ 0 & 0 & 0 & \frac{\exp(-x_n)}{(1+\exp(-x_n))^2} \end{bmatrix}$$

$$z_k'(x) = \begin{bmatrix} \frac{d(\frac{\exp(x_1)}{\sum_{i=1}^K \exp(x_i)})}{dx_1} & \frac{d(\frac{\exp(x_1)}{\sum_{i=1}^K \exp(x_i)})}{dx_2} & \dots & \frac{d(\frac{\exp(x_1)}{\sum_{i=1}^K \exp(x_i)})}{dx_K} \\ \frac{d(\frac{\exp(x_2)}{\sum_{i=1}^K \exp(x_i)})}{dx_1} & \frac{d(\frac{\exp(x_2)}{\sum_{i=1}^K \exp(x_i)})}{dx_2} & \dots & \frac{d(\frac{\exp(x_2)}{\sum_{i=1}^K \exp(x_i)})}{dx_K} \\ \dots & \dots & \dots & \dots \\ \frac{d(\frac{\exp(x_K)}{\sum_{i=1}^K \exp(x_i)})}{dx_1} & \frac{d(\frac{\exp(x_K)}{\sum_{i=1}^K \exp(x_i)})}{dx_2} & \dots & \frac{d(\frac{\exp(x_K)}{\sum_{i=1}^K \exp(x_i)})}{dx_K} \end{bmatrix} = \begin{bmatrix} \frac{\sum_{i=1, i \neq 1}^K \exp(x_i + x_1)}{(\sum_{i=1}^K \exp(x_i))^2} & \frac{-\exp(x_2 + x_1)}{(\sum_{i=1}^K \exp(x_i))^2} & \dots & \frac{-\exp(x_K + x_1)}{(\sum_{i=1}^K \exp(x_i))^2} \\ \frac{-\exp(x_1 + x_2)}{(\sum_{i=1}^K \exp(x_i))^2} & \frac{\sum_{i=1, i \neq 2}^K \exp(x_i + x_2)}{(\sum_{i=1}^K \exp(x_i))^2} & \dots & \frac{-\exp(x_K + x_2)}{(\sum_{i=1}^K \exp(x_i))^2} \\ \dots & \dots & \dots & \dots \\ \frac{-\exp(x_1 + x_K)}{(\sum_{i=1}^K \exp(x_i))^2} & \frac{-\exp(x_2 + x_K)}{(\sum_{i=1}^K \exp(x_i))^2} & \frac{-\exp(x_3 + x_K)}{(\sum_{i=1}^K \exp(x_i))^2} & \frac{\sum_{i=1, i \neq K}^K \exp(x_i + x_K)}{(\sum_{i=1}^K \exp(x_i))^2} \end{bmatrix}$$

Similarities:

Both of functions have a range (0, 1). The outputs can represent probabilities in the output layer of CNN and NN in classification problems. However, sigmoid functions can only be used in the binary classification. Softmax functions is an extension of sigmoid functions at some extend.

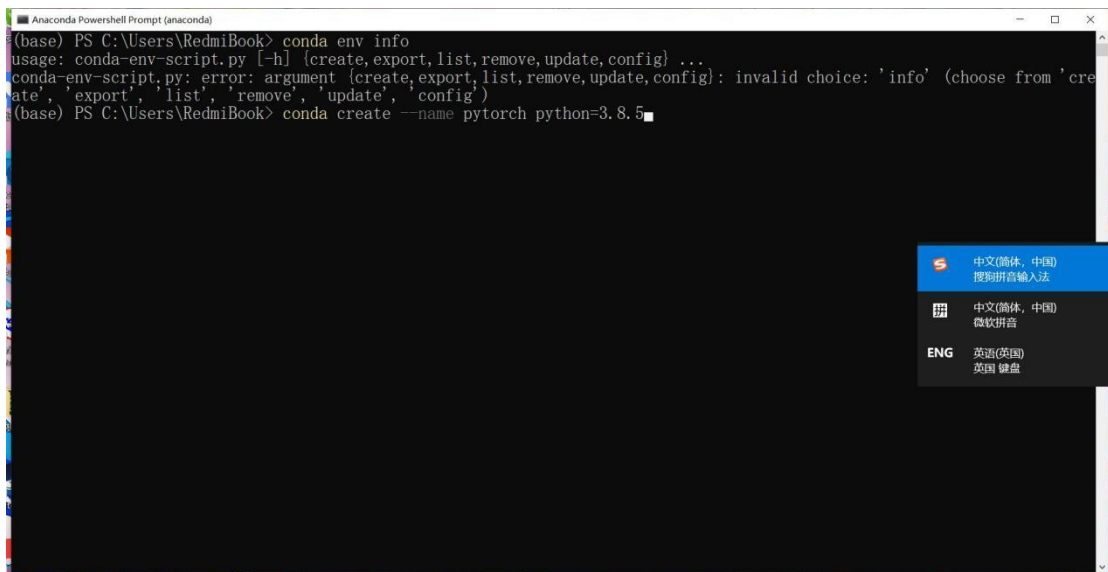
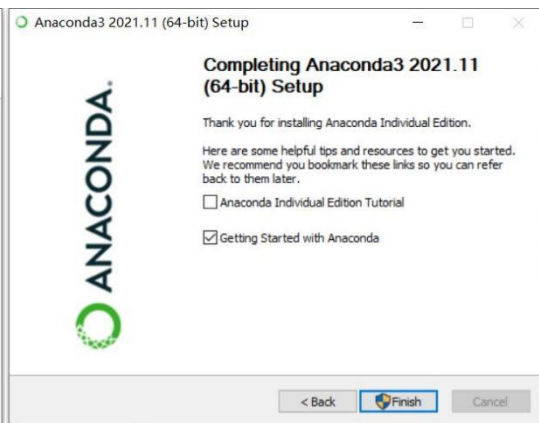
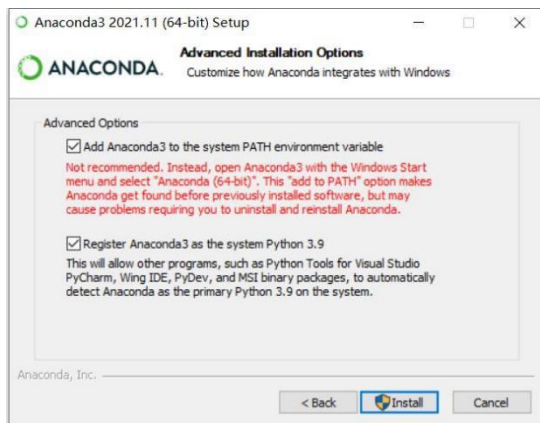
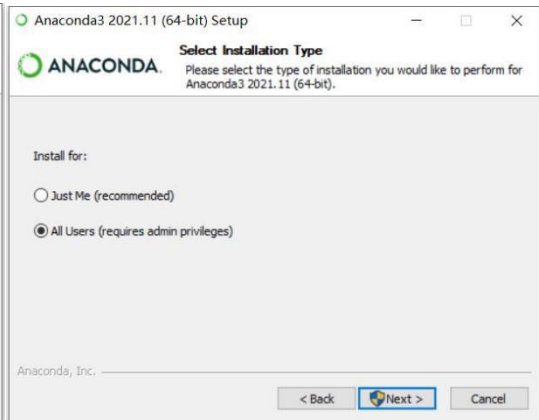
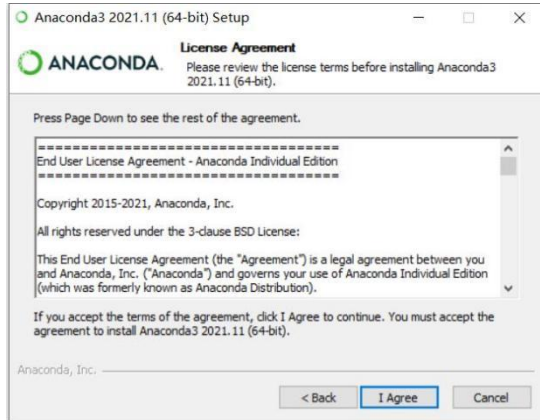
2. Write down the cross-entropy loss function, and its derivatives. Discuss how it works for classification tasks. Is there any disadvantage of using naïve cross-entropy for classification? (10%)

Assume \mathbf{x} is the result vector of the last layer, and the probability generate by softmax function. Then $\mathbf{p}=\text{softmax}(\mathbf{x})$, Assume the target one is the i th vector, then the indicator function \mathbf{y} with $y_i = 1, y_j = 0$ if $j \neq i$. $L=-\sum_i y_i \log(p_i)$, $\frac{dL}{dx_i}=-\sum_i y_i \frac{d\log(p_i)}{dp_i} \frac{dp_i}{dx_i} = \sum_i y_i \frac{1}{p_i} \frac{dp_i}{dx_i}$. In the task, the softmax function transform the result into probability. Cross entropy function provides a judgement for the efficiency of the model. And the cross entropy function is differentiable, which means it can update the model by gradient descending. Yes, the disadvantage is that if p_i is very small, then the gradient will be very large (gradient explode) since it's in the denominator. And softmax function is similar to an exponential function. The repeating calculation of exponential and log functions.

Programming problems:

1. Install Anaconda (a very good Python version that can help you easily implement data science applications).

Based on Anaconda, install some commonly used packages for Python, including Numpy, Pandas, Matplotlib, Seaborn, PyTorch (CPU version is OK if you don't have GPU on your device). Record your installation process in detail, screenshots for every step are needed. (10%)



```

Anaconda Powershell Prompt (anaconda)

package                                     build
-----
ca-certificates-2021.10.26                haa95532_4      116 KB
certifi-2021.10.8                         py38haa95532_2  152 KB
openssl-1.1.1m                           h2bbff1b_0      4.8 MB
pip-21.2.2                               py38haa95532_0  1.9 MB
python-3.8.5                             h5fd99ec_1      15.7 MB
setuptools-58.0.4                       py38haa95532_0  779 KB
sqlite-3.37.2                            h2bbff1b_0      799 KB
wheel-0.37.1                             pyhd3eb1b0_0    33 KB
wincertstore-0.2                         py38haa95532_2  15 KB

Total:                                     24.2 MB

The following NEW packages will be INSTALLED:

ca-certificates      pkgs/main/win-64::ca-certificates-2021.10.26-haa95532_4
certifi              pkgs/main/win-64::certifi-2021.10.8-py38haa95532_2
openssl              pkgs/main/win-64::openssl-1.1.1m-h2bbff1b_0
pip                  pkgs/main/win-64::pip-21.2.2-py38haa95532_0
python               pkgs/main/win-64::python-3.8.5-h5fd99ec_1
setuptools            pkgs/main/win-64::setuptools-58.0.4-py38haa95532_0
sqlite                pkgs/main/win-64::sqlite-3.37.2-h2bbff1b_0
vc                    pkgs/main/win-64::vc-14.2-h21ff451_1
vs2015_runtime        pkgs/main/win-64::vs2015_runtime-14.27.29016-h5e58377_2
wheel                 pkgs/main/noarch::wheel-0.37.1-pyhd3eb1b0_0
wincertstore          pkgs/main/win-64::wincertstore-0.2-py38haa95532_2

Proceed ([y]/n)?

```

```

Anaconda Powershell Prompt (anaconda)

done
# To activate this environment, use
#
#     $ conda activate pytorch
#
# To deactivate an active environment, use
#
#     $ conda deactivate
#

(base) PS C:\Users\RedmiBook>

```

```

Anaconda Powershell Prompt (anaconda)

done
# To activate this environment, use
#
#     $ conda activate pytorch
#
# To deactivate an active environment, use
#
#     $ conda deactivate
#

(base) PS C:\Users\RedmiBook> conda activate pytorch
(pytorch) PS C:\Users\RedmiBook> conda list
# packages in environment at D:\anaconda\envs\pytorch:
#
# Name                  Version      Build      Channel
ca-certificates         2021.10.26   haa95532_4
certifi                 2021.10.8    py38haa95532_2
openssl                 1.1.1m       h2bbff1b_0
pip                     21.2.2       py38haa95532_0
python                  3.8.5        h5fd99ec_1
setuptools               58.0.4       py38haa95532_0
sqlite                  3.37.2       h2bbff1b_0
vc                       14.2         h21ff451_1
vs2015_runtime          14.27.29016  h5e58377_2
wheel                   0.37.1       pyhd3eb1b0_0
wincertstore            0.2          py38haa95532_2
(pytorch) PS C:\Users\RedmiBook> conda install pytorch torchvision torchaudio cudatoolkit=10.2 -c pytorch

```

```

Anaconda Prompt (anaconda) - conda install pytorch torchvision torchaudio cudatoolkit=10.2 -c pytorch
done
(pytorch) C:\Users\RedmiBook>

Anaconda Prompt (anaconda) - conda install pytorch torchvision torchaudio cudatoolkit=10.2 -c pytorch
blas 1.0 mkl
ca-certificates 2021.10.26 haa95332_4
certifi 2021.10.8 py38haa95332_2
cudatoolkit 10.2.89 h74a9793_1
cython 0.11.0 pypi_0 pypi
fairscale 4.59.1 pypi_0 pypi
intel-openmp 2.10.4 hd328e21_0
jpeg 9d h2bbff1b_0
mkl 2021.4.0 haa95332_3556
mkl-service 2.4.0 pypi_0 pypi
mkl_fft 1.3.1 py38h2bbff1b_0
mkl_random 1.2.2 py38hf11a4ad_0
numpy 1.21.5 py38haa95332_0
numpy-base 1.21.5 py38hc2deb75_0
olefile 0.46 pyhd3eb1b0_0
opencv 1.1.1m h2bbff1b_0
packaging 21.3 pypi_0 pypi
pandas 1.4.1 pypi_0 pypi
pillow 8.4.0 py38hd45dc43_0
pip 21.2.2 py38haa95332_0
pyparsing 3.0.7 pypi_0 pypi
python 3.8.5 h5fd99cc_1
python-dateutil 2.8.2 pypi_0 pypi
pytorch 1.10.2 py3.8_cuda10.2_cudnn7.0 pytorch
pytorch-mutex 1.0 cuda pytorch
pytz 2021.3 pypi_0 pypi
scipy 1.8.0 pypi_0 pypi
seaborn 0.11.2 pypi_0 pypi
setuptools 58.0.4 py38haa95332_0
six 1.16.0 pyhd3eb1b0_0
tqdm 4.62.3 h2bbff1b_0
torchaudio 0.10.2 py38_cu102 pytorch
torchvision 0.11.3 py38_cu102 pytorch
tqdm 0.0.0 pypi_0 pypi
typing_extensions 3.10.0.2 pyh06a4308_0
wcwidth 1.0.2 h2f4451_1
wheel 0.37.1 pyhd3eb1b0_0
winertstore 0.2 py38haa95332_2
xz 5.2.5 h624d977_0
zlib 1.2.11 h8cc25b3_4

```

2. Here's two basic Python programming problems: (10%)

1) Write a program to sum all the numbers in a list;

```

Q2(1)

list=[i for i in range(100)]
s=0
for i in list:
    s+=i
s

[3]
... 4950

```

2) Write a program to find all prime numbers in a list.

```
def isprime(M):
    for i in range(2,int(np.sqrt(M))+1):
        if M%i==0:
            return True
    return False
list= [i for i in range(100)]

ans=[]
for i in list:
    if isprime(i)==False:
        ans.append(i)
ans
```

3. Here's two basic Numpy programming problems: (10%)

1) Given an array, calculate its softmax result;

```
def softmax(x):
    return np.exp(x)/np.sum(np.exp(x))
x=np.array([1,2,3])
softmax(x)
```

array([0.09003057, 0.24472847, 0.66524096])

2) use the softmax result to calculate the cross-entropy loss (you are free to construct the input x, and ground truth label y by yourself).

```
x=np.array([1,2,3])
y=softmax(x)
-np.log(y[0])
```

2.40760596444438

4. Medical data exploration. You are going to work with a real medical image dataset, including exploring data labels, visualizing and observing the data, along with processing the data. Please fill in the blanks in “data_exploration.ipynb” and successfully run all the cells. (20%)


```
# Plot a histogram of the distribution of the pixels
### Your code here ###
raw_image=raw_image.reshape(1,1048576)
sns.distplot(raw_image,bins=len(np.unique(raw_image)),kde=False,color='b')
import matplotlib.patches as mp
info = mp.Patch(label='Pixel Mean: %.4f & Standard Deviation %.4f'%(raw_image.std(),raw_image.mean()))
### End your code ###
plt.legend(handles=[info],loc='upper center')
plt.title('Distribution of Pixel Intensities in the Image')
plt.xlabel('Pixel Intensity')
plt.ylabel('# Pixels in Image')
```

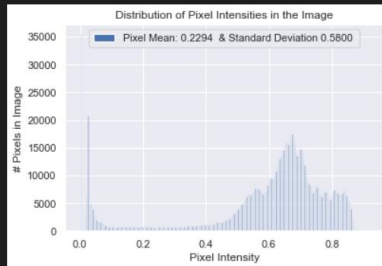
✓ 0.6s

Python

D:\anaconda\envs\pytorch\lib\site-packages\seaborn\distributions.py:2619: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

warnings.warn(msg, FutureWarning)

Text(0, 0.5, '# Pixels in Image')



```
# Create transformation for data standardization

norm_mean = [0.6, 0.6, 0.6]
norm_std = [0.225, 0.225, 0.225]
data_transform = transforms.Compose([
    transforms.Resize((320, 320)),
    transforms.ToTensor(),
    ### Your code here, use API like transforms.XXX() ###
    transforms.Normalize(norm_mean, norm_std)
    ### End your code ###
])
dataset = XRayDataset('nih/images-small', data_transform)
```

✓ 0.8s

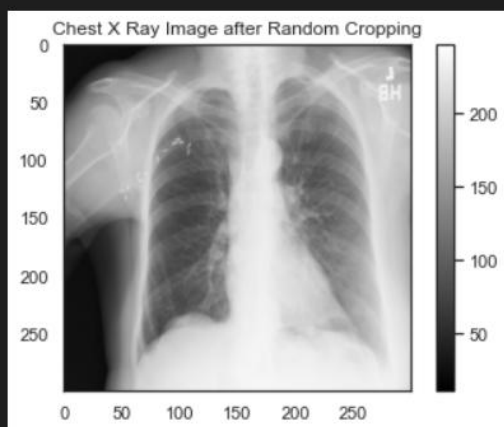
```
# Random cropping

data (module) transforms ns.Compose([
    transforms.Resize((320, 320)),
    ### Your code here, use API like transforms.XXX() ###
    transforms.RandomCrop(300),
    ### End your code ###
    transforms.ToTensor()
])
dataset = XRayDataset('nih/images-small', data_transform)

sns.set_style("white")
generated_image = dataset.__getitem__(0)
inverted_image = transform_invert(copy.deepcopy(generated_image), data_transform)
plt.imshow(inverted_image, cmap='gray')
plt.colorbar()
plt.title('Chest X Ray Image after Random Cropping')
```

✓ 0.3s

Text(0.5, 1.0, 'Chest X Ray Image after Random Cropping')



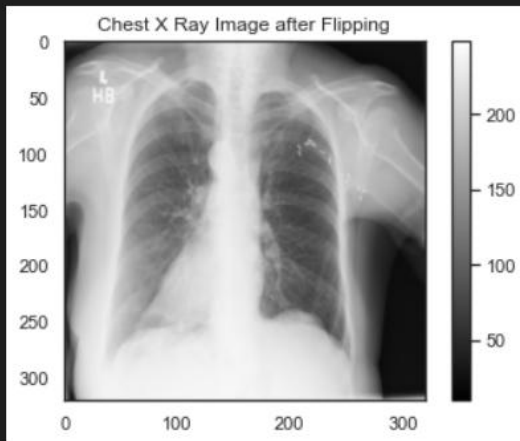
```
# Horizontal flip

data_transform = transforms.Compose([
    transforms.Resize((320, 320)),
    ### Your code here, use API like transforms.XXX() ###
    transforms.RandomHorizontalFlip(p=1),
    ### End your code ###
    transforms.ToTensor()
])
dataset = XRayDataset('nih/images-small', data_transform)

sns.set_style("white")
generated_image = dataset.__getitem__(0)
inverted_image = transform_invert(copy.deepcopy(generated_image), data_transform)
plt.imshow(inverted_image, cmap='gray')
plt.colorbar()
plt.title('Chest X Ray Image after Flipping')
```

✓ 0.2s

Text(0.5, 1.0, 'Chest X Ray Image after Flipping')



```
# Random rotation

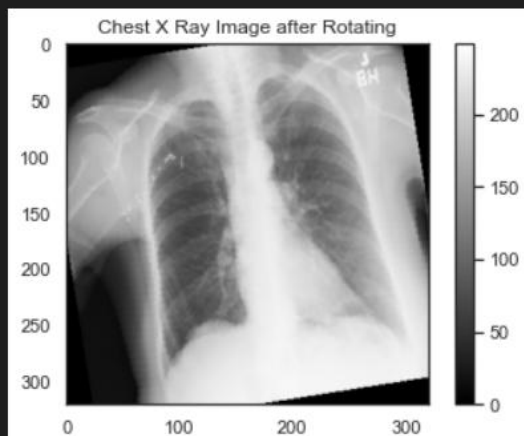
data_transform = transforms.Compose([
    transforms.Resize((320, 320)),
    ### Your code here, use API like transforms.XXX() ###
    transforms.RandomRotation(20),
    ### End your code ###
    transforms.ToTensor()
])

dataset = XRayDataset('nih/images-small', data_transform)

sns.set_style("white")
generated_image = dataset.__getitem__(0)
inverted_image = transform_invert(copy.deepcopy(generated_image), data_transform)
plt.imshow(inverted_image, cmap='gray')
plt.colorbar()
plt.title('Chest X Ray Image after Rotating')
```

✓ 0.2s

Text(0.5, 1.0, 'Chest X Ray Image after Rotating')



```
# Color jittering

data_transform = transforms.Compose([
    transforms.Resize((320, 320)),
    ### Your code here, use API like transforms.XXX() ###
    transforms.ColorJitter(brightness=0.1, contrast=0, saturation=1, hue=0.1),
    ### End your code ###
    transforms.ToTensor()
])
dataset = XRayDataset('nih/images-small', data_transform)

sns.set_style("white")
generated_image = dataset.__getitem__(0)
inverted_image = transform_invert(copy.deepcopy(generated_image), data_transform)
plt.imshow(inverted_image, cmap='gray')
plt.colorbar()
plt.title(['Chest X Ray Image after Color Jittering'])
```

✓ 0.4s

Text(0.5, 1.0, 'Chest X Ray Image after Color Jittering')

