

JunHyuk Kim 2023 Report-ISIC 2017 improved Unet segmentation task.

This is a brief description of my project. The project is made for segmentation of melanoma. Task finished was ”

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2. Requiriments

2.1. Download Dependencies

Modules that are required to run the code: `import tqdm import torchvision import albumentations import torch import matplotlib import sklearn import glob import PIL import os import pandas import numpy python 3.11.5 window 11`

2.2. Download ISIC 2017 datasets.

The dataset is under <https://challenge.isic-archive.com/data/#2017>. Training data, test data, valid data is required and train ground truth, test ground truth and valid ground truth is also required for the segmenation task. The dataset should be positioned in the same folder as the python codes under data, train data images under train_images folder, and train mask data under train_masks folder. Test data should be in the test_images and test_masks and validation should be in val_images and val_masks folder.

3. Usage

Run the Train.py with python3 this will create the model.pth file. This will create predictions for each epochs and will also create the loss graph.

Run predict.py to get the segmentation and the dice coefficient. This file will save all the segmentation predictions under evaluation_folder.

4. Description

4.1. Brief description of the task

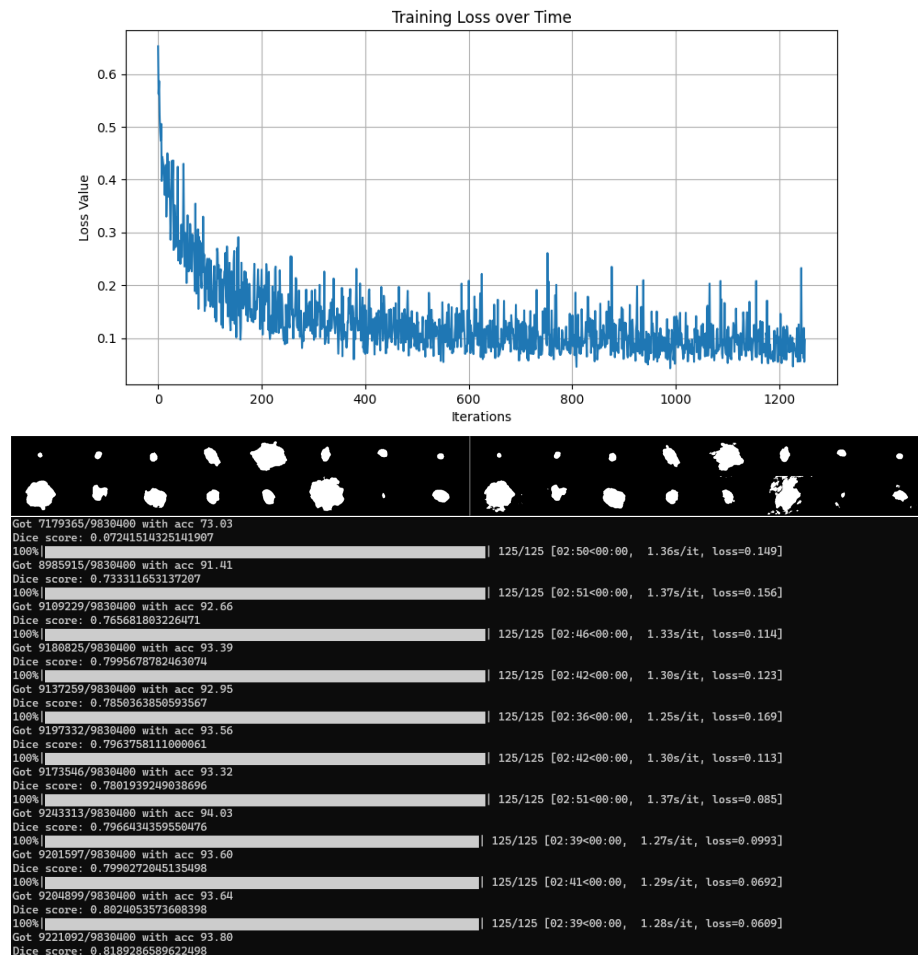
Lesion segmentation is crucial in medical imaging. The code solution that is created solves the problem of the segmentation of lesion by automating it, which can benefit the medical professionals. The ISIC 2017 improved Unet

segmentation task is implemented. The improve Unet is used and the format follows <https://arxiv.org/pdf/1802.10508v1.pdf>. By doing this, the accuracy and speed of segmenation can be increased significantly.

4.2. Reproducibility After 10 epochs, the model relably provided dice score around 80 percent.

4.3. provide example inputs, outputs and plots of your algorithm

For every epoch, the validation set is used to test the segmenation. ##### 4.3.1.
The output when train.py is ran



4.3.2. The output expected after running predict.py

```
Got 36461547/39321600 with acc 92.73  
Dice score: 0.8383143544197083
```



4.4. Describe any specific pre-processing you have used with references if any. Justify your training,

4.4.1. pre-processing I used transformation to increase the number of samples the model is trained on. For transformation the albumentations module was used. This makes the transformation much easier. Transformation used are, rotate, horizontalflip, verticalflip. Also the values were normalized. I also resized the images into 256x256 sizes to increase the performance of training and also because my model works on that sizes.

4.4.2. Training The training is done in batch size of 16 with 10 epochs. The optimizer used is ADAM and the loss function used is Dice loss function. The scheduler was used to reduce the learning rate at the end of the function. The training accuracy and saved predictions are from validation imagesets. GradScaler is used to make the training faster.

4.5 Commentation Each class and functions/methods are commented. Comment includes a brief description of the function of the class, method or the function and followed by the parameters it receives and the return values it returns.

5. Algorithm/ImprovedUnet

This program uses improved Unet to segment the different types of skin conditions such as melanoma. The improved Unet structure explained ### 5.1. Context layer and Localization Layer. Context Layer consists of 2d convolutional layer with kernel size 3, InstanceNorm2d, LeakyReLU and dropout layer with p value of 3. After that, another 2d convolutional layer with kernel size 3 is applied followed by instanceNorm2d and LeakyReLU. Context layer normally receives the same output as input channel, which means going through context layer doesn't really change the dimension of the tensor.

Localization Layer consists of 2d convolutional layer with kernel size 3, InstanceNorm2d, LeakyReLU. After that, another 2d convolutional layer with kernel size 1 is applied followed by instanceNorm2d and LeakyReLU. Localization layer changes the dimension of the tensor given by halving it. This is because it receives the concatenation of skip_connection and the previous output from the previous layer.

5.2. Layer 1 to Layer 5.

Firstly, 2d convolutional layer is applied to the input tensor X, and the output is saved. First convolutional layer changes the dimension of the tensor to 3 to 64, but all the other pathways doubles the dimension of the tensor. The output is fed into the context layer, and the output of the context layer and the first convolutional layer is added element wise using the “torch.add()” function. The output is saved in the name of skip_connection_n (where n is the layer which the context pathway is in).

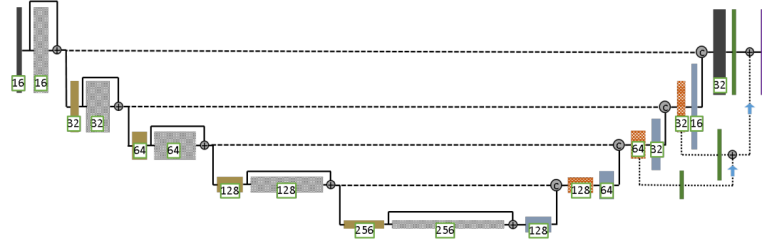
Then this continues 4 more times except for the 5th time the skip connection is not saved.

5.3. Layer 5 to Layer 1.

In the layer 5 after applying the context pathway for 5 times, localization pathway is applied. Localization pathway consists of first upsampling the tensor (which doubles the size of the image) and a convolutional layer with kernel size of 1, which halves the amount of features the tensor has. After this, the skip layer which corresponds to the layer which the localization layer is positioned is concatenated to the previous output. Then the output is fed into the localization layer, which will halve the amount of features the tensor has. This is because the number of features is doubled after concatenation. This continues until the output reaches the layer 1, where instead of localization a normal convolutional layer with kernel size 3 is applied. This doesn't change the amount of the feature map.

During the localization pathway, in Layer 3 localization and Layer 4 localization, and after the final convolutional layer, segmentation layer which is just a convolutional layer with output channel of 1 is applied and the output is saved under the name of segmentation 1, segmentation 2 and segmentation 3. Segmentation 1 is upsampled and added element wise to segmentation 2, and the combination of those are upsampled and added to segmentation 3 element wise.

Finally the combination of all are ran through Sigmoid function and the model re-



turns the output.