CS440 Introduction to AI

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Class Challenge Report: Image Classification of COVID-19 X-rays

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Task 1 - Binary Classification (Normal vs. COVID-19 X-rays)

Architecture:

For Task 1, we used a CNN model, VGG16, that used pre-trained weights obtained by training on ImageNet. A VGG16 model consists of the 16 layers shown in figure 1.1.2. This was followed by 1 flatten layer (named "flatten"), 1 dense layer (named "dense_feature"), 1 dropout layer (named "dropout"), and lastly, 1 dense layer (named "dense"). The first dense layer, "dense_feature," was used as a feature extraction layer with a RELU activation function, and had 256 nodes. The "dropout" layer had a dropout rate of 0.25 and 256 nodes as well. Then, the second and final dense layer used the sigmoid activation function. This layer had 1 unit for classification since there are only 2 classes of images - normal and COVID-19. The model contained a total of 16 + 3 = 19 layers (Fig. 1.1.1).

Model: "sequential"

Layer (type)	Output	Shape	Param #
vgg16 (Functional)	(None,	7, 7, 512)	14714688
flatten (Flatten)	(None,	25088)	0
dense_feature (Dense)	(None,	256)	6422784
dropout (Dropout)	(None,	256)	0
dense (Dense)	(None,	1)	257

Total params: 21,137,729
Trainable params: 6,423,041
Non-trainable params: 14,714,688

Figure 1.1.1 VGG16 Model Layers

Layer (type)	Output Shape	Param #
 conv2d (Conv2D)	(None, 224, 224, 64)	1792
conv2d_1 (Conv2D)	(None, 224, 224, 64)	36928
max_pooling2d (MaxPooling2D)	(None, 112, 112, 64)	0
conv2d_2 (Conv2D)	(None, 112, 112, 128)	73856
conv2d_3 (Conv2D)	(None, 112, 112, 128)	147584
max_pooling2d_1 (MaxPooling2	(None, 56, 56, 128)	0
conv2d_4 (Conv2D)	(None, 56, 56, 256)	295168
conv2d_5 (Conv2D)	(None, 56, 56, 256)	590080
conv2d_6 (Conv2D)	(None, 56, 56, 256)	590080
max_pooling2d_2 (MaxPooling2	(None, 28, 28, 256)	0
conv2d_7 (Conv2D)	(None, 28, 28, 512)	1180160
conv2d_8 (Conv2D)	(None, 28, 28, 512)	2359808
conv2d_9 (Conv2D)	(None, 28, 28, 512)	2359808
max_pooling2d_3 (MaxPooling2	(None, 14, 14, 512)	0
conv2d_10 (Conv2D)	(None, 14, 14, 512)	2359808
conv2d_11 (Conv2D)	(None, 14, 14, 512)	2359808
conv2d_12 (Conv2D)	(None, 14, 14, 512)	2359808
max_pooling2d_4 (MaxPooling2	(None, 7, 7, 512)	0
flatten (Flatten)	(None, 25088)	0
dense (Dense)	(None, 4096)	102764544
dropout (Dropout)	(None, 4096)	0
dense_1 (Dense)	(None, 4096)	16781312
dropout_1 (Dropout)	(None, 4096)	0
dense_2 (Dense)	(None, 1000)	4097000

Figure 1.1.2 VGG16 Model Layers

The following table consists of hyperparameters we used to fine-tune our VGG16 model:

Hyperparameter	Name
Optimizer	Adam
Loss function	Binary cross-entropy
Parameters	 Total parameters: 21,137,729 Trainable parameters: 6,423,041 Non-trainable parameters: 14,714,688
Regularization	None
Learning rate	0.0001
Batch size	10
Number of Epochs	40

Figure 1.2 VGG16 Model Parameters

Model Accuracy and Loss:

After training was completed, on average, our model produced a training accuracy of 0.95 and a training loss of 0.15 (Fig. 1.3-1.4). Our model also produced a test accuracy of 1.0 and a test loss of approximately 0.09 (Fig. 1.5).

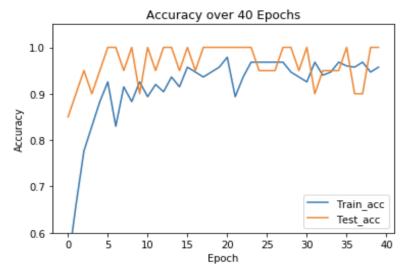


Figure 1.3 VGG16 Model Accuracy

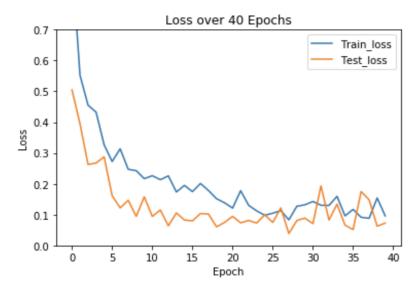


Figure 1.4 VGG15 Model Loss

Figure 1.5 VGG16 Model Testing Accuracy & Loss

T-SNE Visualization:

We visualized features of the training data by extracting the "dense feature" layer for the t-SNE plot. By observing the t-SNE graph, we concluded that the model performed relatively well, as seen by the 2 distinct clusters representing COVID-19 (blue) and normal (orange) data points (Fig. 1.6).

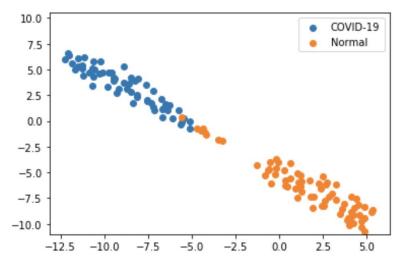


Figure 1.6 VGG16 t-SNE Visual

Task 2 - Multi Classification (Normal, COVID-19, Pneumonia-Bacterial, and Pneumonia-Viral)

Architecture:

For Task 2, we compared a VGG16 and ResNet50, both of which used pre-trained weights obtained by training on ImageNet. The VGG16 model was followed by 1 flatten layer (named "flatten"), 1 dense layer (named "dense_feature"), 1 dropout layer (named "dropout_1"), and lastly, 1 dense layer (named "dense"). The first dense layer, "dense_feature," was used as a feature extraction layer, with a RELU activation function, and had 256 nodes. The "dropout_1" layer had a dropout rate of 0.5. The model outputted to the second and last "dense" layer which used the softmax activation function. This layer had 4 units for classification since there were 4 classes of images - normal, COVID-19, Pneumonia-Bacterial, and Pneumonia-Viral. The model contained a total of 16 + 4 = 20 layers (Fig 1.1.2, 2.1.1).

Model: "sequential"

Layer (type)	Output	Shape	Param #
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vgg16 (Functional)	(None,	7, 7, 512)	14714688
flatten (Flatten)	(None,	25088)	0
dense_feature (Dense)	(None,	256)	6422784
dropout_1 (Dropout)	(None,	256)	0
dense (Dense)	(None,	4)	1028
			=======
Total params: 21,138,500			
Trainable params: 6,423,812			
Non-trainable params: 14,714	1 600		
Mon-cramable params: 14,/14	,000		

Figure 2.1.1 VGG16 Model Layers

A ResNet50 model consists of 50 layers. The ResNet50 model was followed by 1 2D convolutional layer (named "conv2d_8"), 1 flatten layer (named "flatten"), 1 dense layer (named "dense_feature"), 1 dropout layer (named "dropout"), and lastly, 1 dense layer (named "dense"). The "conv2d_8" layer convoluted the output of the ResNet layer to 1024 nodes with a kernel size of (1,1) and padding. The first dense layer, "dense_feature", was used as a feature extraction layer, with 256 nodes and a RELU activation function. The "dropout" layer had a dropout rate of 0.25. The model outputted to the second and last "dense" layer, which used the softmax activation function. This layer had 4 units for classification since there were 4 classes of images - normal, COVID-19, Pneumonia-Bacterial, and Pneumonia-Viral. The model contained a total of 50 + 5 = 55 layers (Fig 2.1.2).

Layer (type)	Output Shape	Param #
resnet50 (Functional)	(None, 7, 7, 2048)	23587712
conv2d_9 (Conv2D)	(None, 7, 7, 1024)	2098176
flatten (Flatten)	(None, 50176)	0
dense_feature (Dense)	(None, 128)	6422656
dropout (Dropout)	(None, 128)	0
dense (Dense)	(None, 4)	516
Motel marger 22 100 060		

Total params: 32,109,060 Trainable params: 32,055,940 Non-trainable params: 53,120

Figure 2.1.2 ResNet50 Model Layers

The following table consists of hyperparameters we used to fine-tune our VGG16 model:

Hyperparameter	Name
Optimizer	Adam
Loss function	Categorical cross-entropy
Parameters	 Total parameters: 21,138,500 Trainable parameters: 6,423,812 Non-trainable parameters: 14,714,688
Regularization	None
Learning rate	0.0001
Batch size	10
Number of Epochs	100

Figure 2.2.1 VGG16 Model Parameters

The following table consists of hyperparameters we used to fine-tune our ResNet50 model:

Hyperparameter	Name
Optimizer	Adam
Loss function	Categorical cross-entropy
Parameters	 Total parameters: 32,109,060 Trainable parameters: 32,055,940 Non-trainable parameters: 53,120
Regularization	None
Learning rate	0.00001
Batch size	10
Number of Epochs	100

Figure 2.2.2 ResNet50 Model Parameters

Model Accuracy and Loss:

After training was completed, the VGG16 model produced a test accuracy of approximately 0.64 and a test loss of approximately 0.76 (Fig. 2.3.1-2.3.3).

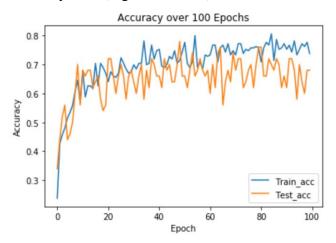


Figure 2.3.1 VGG16 Model Accuracy

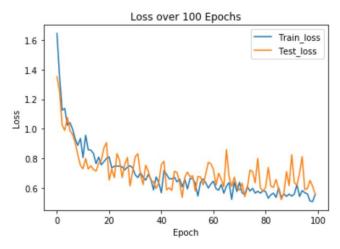


Figure 2.3.2 VGG16 Model Loss

Test loss: 0.7577974796295166
Test accuracy: 0.6388888955116272

Figure 2.3.3 VGG16 Test Accuracy and Test Loss

After training was completed, the ResNet50 model produced a test accuracy of approximately 0.69 and a test loss of approximately 0.86 (Fig. 2.4.1-2.4.3).

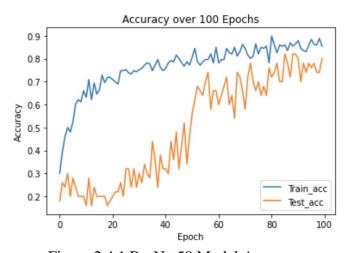


Figure 2.4.1 ResNet50 Model Accuracy

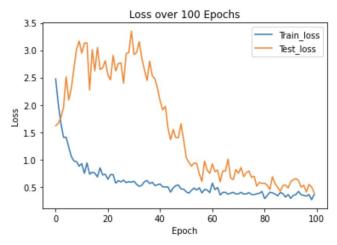


Figure 2.4.2 ResNet50 Model Loss

Figure 2.4.3 ResNet50 Model Test Accuracy & Loss

T-SNE Visualization:

For the VGG16 model, we visualized features of the training data by extracting the "dense feature" layer for the t-SNE plot. By observing the t-SNE graph, we can conclude that the model performed relatively decently, with 2 distinct clusters for COVID-19 and normal data points, but undistinguishable clusters for pneumonia-bacteria and pneumonia-viral (Fig. 2.5.1).

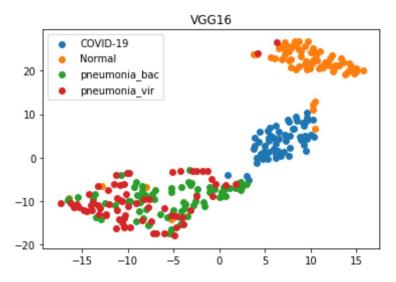


Figure 2.5.1 VGG16 t-SNE Visual

For the ResNet50 model, we visualized features of the training data by extracting the "dense feature" layer for the t-SNE plot. By observing the t-SNE graph, we can conclude that the model performed relatively decently, with 2 distinct clusters for COVID-19 and normal data points, but undistinguishable clusters for pneumonia-bacteria and pneumonia-viral (Fig. 2.5.2).

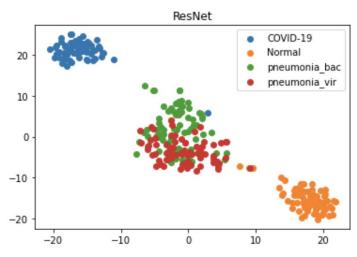


Figure 2.5.2 ResNet50 t-SNE Visual

Comparison of Performance:

For Task 2, we used the following 2 models: VGG16 and ResNet50. Comparing the two models' hyperparameters, the ResNet50 model had 32,109,060 total parameters - 32,055,940 (trainable) and 53,120 (non-trainable), whilst the VGG16 model had 21,138,500 total parameters - 6,423,812 (trainable) and 14,714,688 (non-trainable). As for the rest of the parameters, we kept them the same for both models as seen in figure 2.6.1.

Hyperparameter	ResNet50	VGG16
Optimizer	Adam	Adam
Loss function	Categorical cross-entropy	Categorical cross-entropy
Parameters	 Total parameters: 32,109,060 Trainable parameters: 32,055,940 Non-trainable parameters: 53,120 	 Total parameters: 21,138,500 Trainable parameters: 6,423,812 Non-trainable parameters: 14,714,688
Regularization	None	None
Learning rate	0.0001	0.0001

Batch size	10	10
Number of Epochs	100	100

Figure 2.6.1

Model Analysis:

For a successful model, a higher test accuracy and a lower test loss are desirable. Furthermore, the t-SNE visualization produced by the model should correctly classify the 4 classes - COVID-19, normal, pneumonia-bacteria and pneumonia-viral. This is seen through the clear separation of data points into 4 distinct clusters.

In terms of test accuracy, the ResNet50 model performed better than VGG16 with test accuracies of 0.69 and 0.64 respectively. As for the test loss, the ResNet50 model also performed better than VGG16 with test losses of 0.86 and 0.89 respectively. In terms of the t-SNE visualization, the ResNet50 model appears to have a clearer distinction between the COVID-19 and normal X-rays data points. However, both models failed to successfully separate the 2 pneumonia clusters (seen by the mix of red and green data points). Although the test and training accuracies and losses for the VGG16 model were more consistent with each other, the ResNet50 model had the best performance overall.

Upon reflection, introducing overfitting-prevention techniques such as regularization and early-stopping into training could have improved the overall performance of both models. Despite adding dropout layers in both models, the test accuracies were relatively low and the test losses were relatively high. Therefore, replacing or even adding different regularization techniques might have reduced the test losses and in turn, increased test accuracies. Overall, with the given time and limited resources, we believe our model modestly showcased our ability to perform binary and multi-class classification.¹

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¹ We really tried. Thank you for reading our report, and have a wonderful winter break :)