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INTERMATHS



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Applications of Neural Network Models in Kaggle Competitions

May 22, 2019

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Abstract

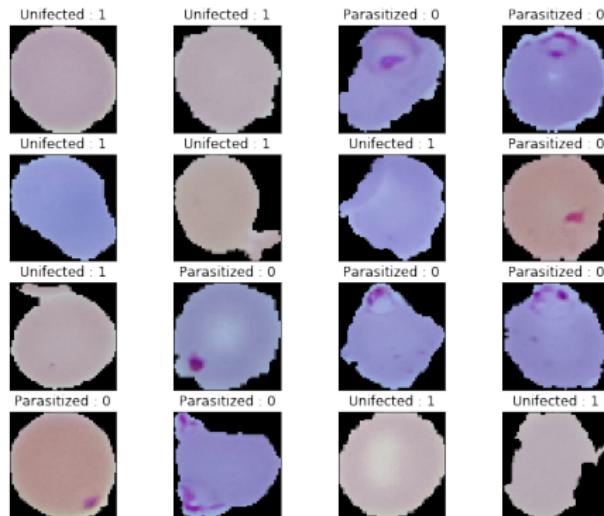
Neural networks are a collection of precise rules which is used in recognizing patterns in data generated from various real life applications such as medicine etc.

Aims

- To understand and implement neural network models on certain medical image datasets via the Kaggle platform
- Classify these images and analyze the performance of the models
- Hyperparameter tuning and comparison of optimizers
- Reduces the workload and misdiagnosis of such diseases by medical practitioners

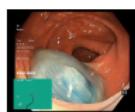
Kaggle Datasets

Figure: Cells from the Pre-processed Malaria Dataset

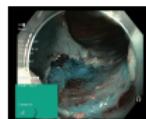


Kaggle Datasets

Figure: Endoscope images of the 8 classes from the Kvasir dataset



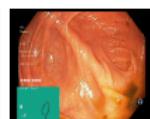
(a)
dyedlifted-
polyps



(b) dyed-
resection-
margins



(c) normal-
esophagus



(d) normal-
cecum



(e) normal-
pylorus



(f) normal-
z-line



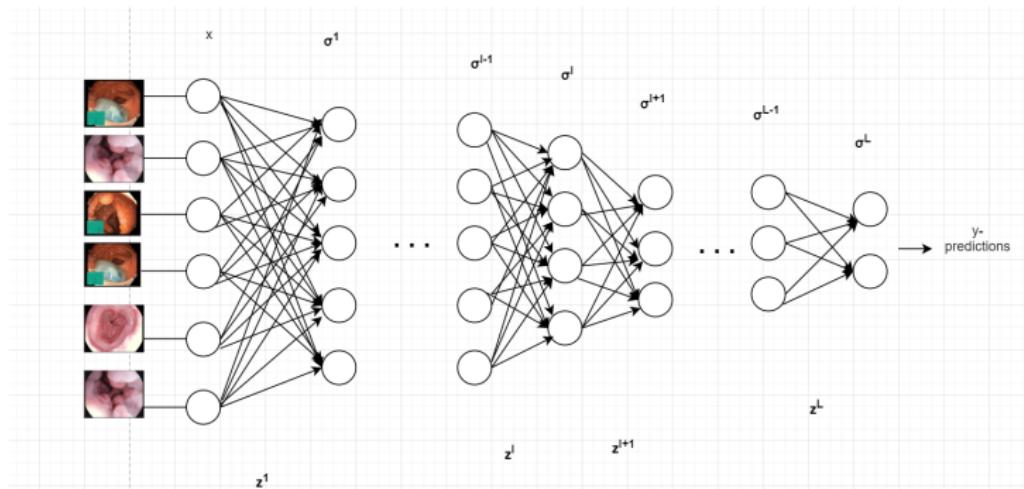
(g) polyps



(h) ulcerative-
colitis

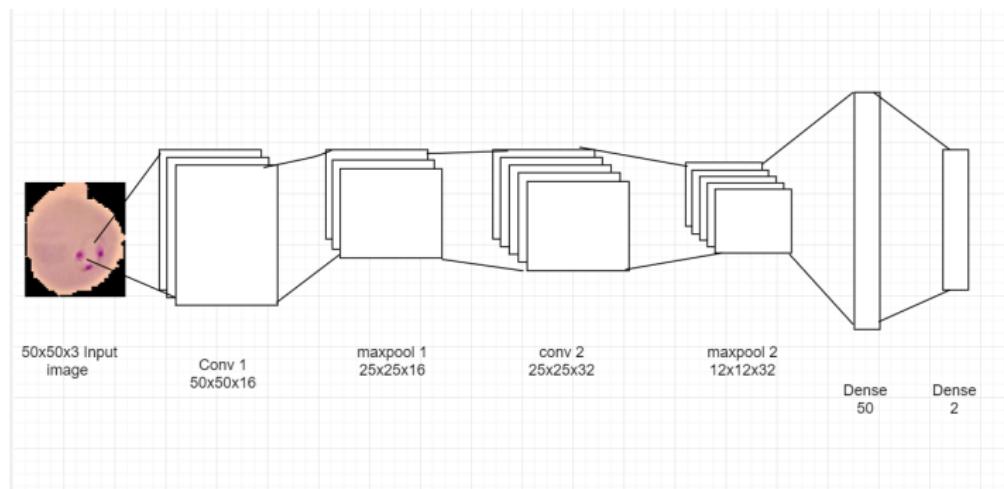
Topology Of Neural Network Models

The Fully Connected Neural Network (FCN) is usually comprised of three (3) basic layers namely: the input layer, the hidden layers and the output layers.



Topology Of Neural Network Models

A standard architecture of the Convolutional Neural Network (CNN) is comprised of the convolutional layers with a nonlinear activation function, the pooling layers and the fully connected layers.



Topology Of Neural Network Models

- Parameters with activation:

$$\sigma_i^{[l]} = \sigma^l(z_i^{[l]}) = \sigma^l\left(\sum_i (w_i^{[l]})^T \cdot x_i^{[l-1]} + b_i^{[l]}\right),$$

- where the activation can be:

$$ReLU, \quad \sigma(z_i) = \max(0, z_i),$$

$$Softmax, \quad \sigma_i(z_i) = \frac{\exp^{-z_i}}{\sum_j \exp^{-z_j}}.$$

- Loss function:

$$\mathfrak{L} = -\frac{1}{m} \sum_{i=1}^m y_i \log(\hat{y}_i) + (1 - y_i) \log(1 - \hat{y}_i).$$

Topology Of Neural Network Models

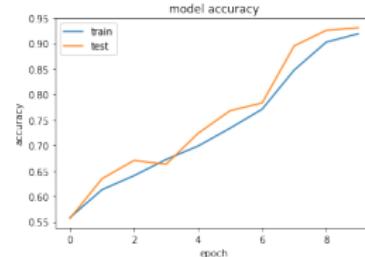
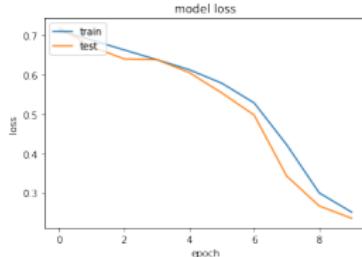
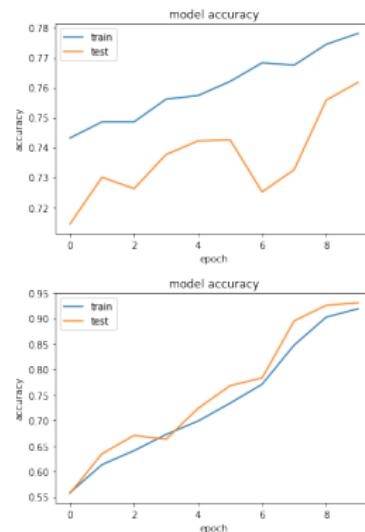
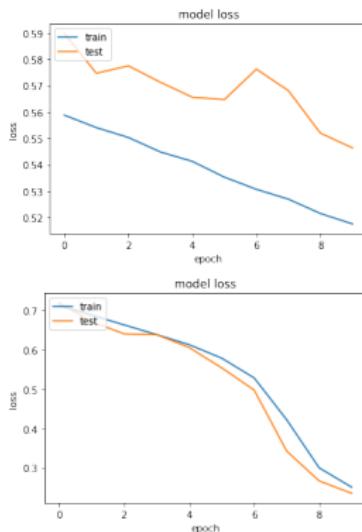
- Parameter update via backpropagation:

$$z_i = z_i - \eta \Delta_i z_i.$$

- Regularization:

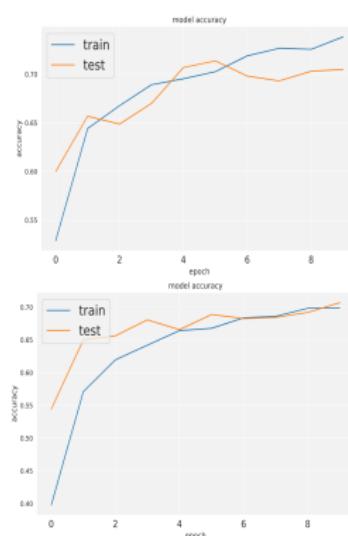
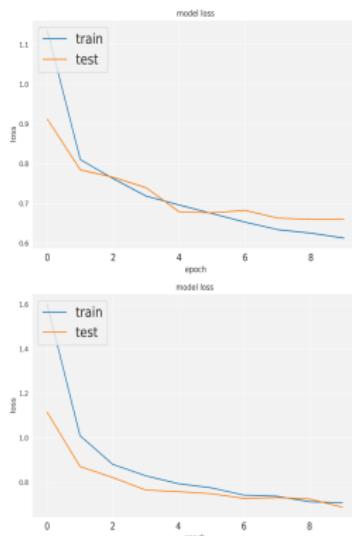
$$\mathcal{L}_{L1} = \mathcal{L} + \lambda \sum_{i=0}^m |z_i|, \quad \lambda, z_i \text{ are the parameters.}$$

Results: Analysis of Models for Malaria Samples



The FCN had an accuracy of 77.81% with loss 0.5176 on the train set and accuracy of 76.18% with loss 0.5464 on the test set. The CNN had an accuracy of 91.94% with loss 0.2511 on the train set and accuracy of 93.11% with loss 0.2356 on the test set.

Results: Analysis of Models for Kvasir Samples



The FCN had an accuracy of 73.84% with loss 0.6130 on the train set and accuracy of 70.50% with loss 0.6602 on the test set. The CNN had an accuracy of 71.08% with loss 0.6942 on the train set and accuracy of 70.06% with loss 0.6975 on the test set.

Results: Analysis of Models

Table: Confusion Matrix and Classification Report for Malaria

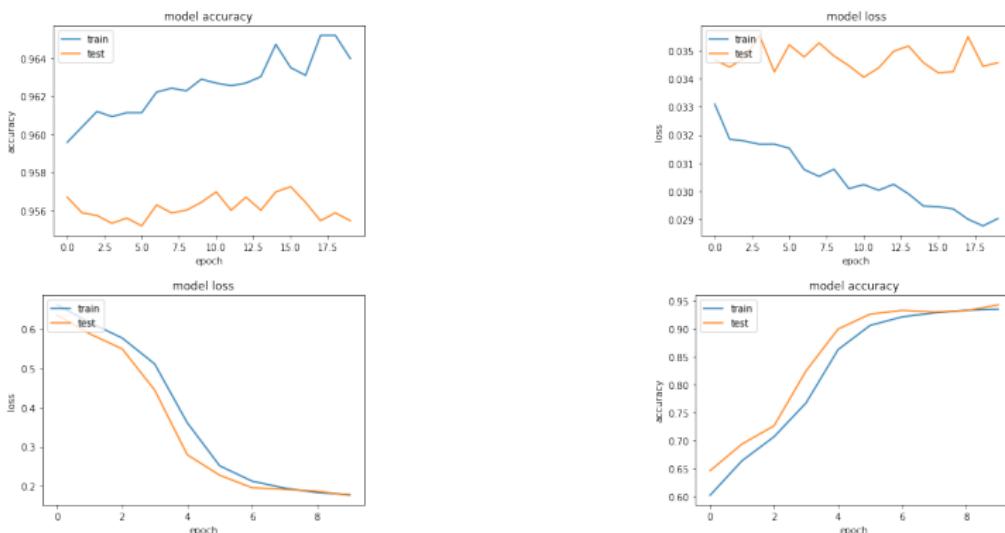
| Malaria Classes | 0. Parasitized | 1. Uninfected | recall | total |
|-----------------|----------------|---------------|--------|-------|
| 0. Parasitized | 2470 | 264 | 0.90 | 2734 |
| 1. Uninfected | 116 | 2662 | 0.96 | 2778 |
| Precision | 0.96 | 0.91 | - | - |
| f1-score | 0.93 | 0.93 | - | - |
| total | 2586 | 2926 | - | 5512 |

Results: Analysis of Models

Table: Confusion Matrix and Classification Report for Kvasir dataset

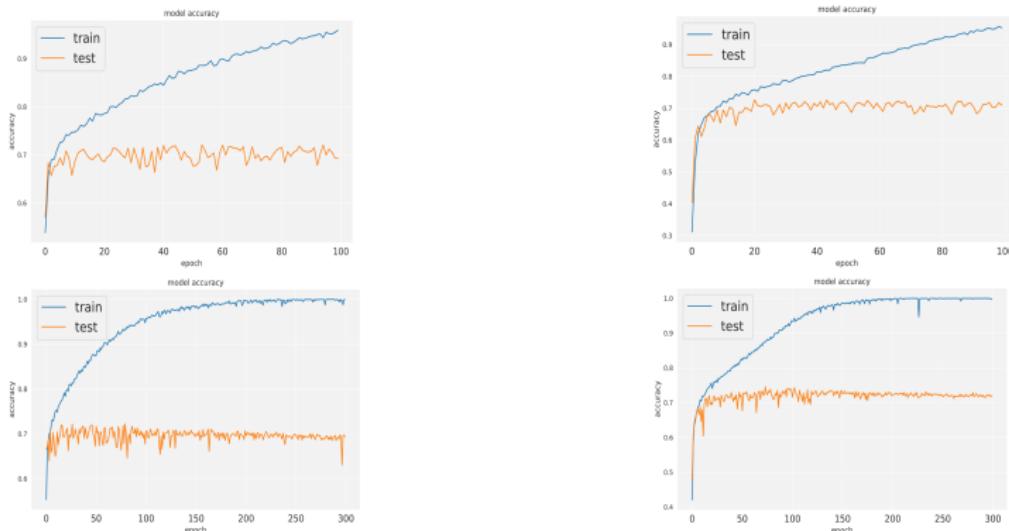
| Endoscopic Classes | 0. | 1. | 2. | 3. | 4. | 5. | 6. | 7. | recall | total |
|---------------------------|------|------|------|------|------|------|------|------|--------|-------|
| 0. Dyed-lifted polyps | 105 | 80 | 0 | 1 | 0 | 0 | 13 | 0 | 0.53 | 199 |
| 1. Dyed-resection margins | 53 | 130 | 0 | 0 | 0 | 0 | 5 | 1 | 0.69 | 189 |
| 2. Esophagitis | 0 | 1 | 108 | 0 | 6 | 82 | 0 | 0 | 0.55 | 197 |
| 3. Normal Cecum | 0 | 0 | 0 | 180 | 0 | 0 | 17 | 7 | 0.88 | 204 |
| 4. Normal Pylorus | 0 | 0 | 1 | 0 | 195 | 5 | 2 | 0 | 0.96 | 203 |
| 5. Normal Z-line | 0 | 0 | 25 | 0 | 22 | 174 | 0 | 1 | 0.78 | 222 |
| 6. Polyps | 2 | 0 | 0 | 29 | 6 | 0 | 125 | 22 | 0.68 | 184 |
| 7. Ulcerative colitis | 1 | 1 | 0 | 30 | 3 | 0 | 53 | 114 | 0.56 | 202 |
| Precision | 0.65 | 0.61 | 0.81 | 0.75 | 0.84 | 0.67 | 0.58 | 0.79 | - | - |
| f1-score | 0.58 | 0.65 | 0.65 | 0.81 | 0.90 | 0.72 | 0.63 | 0.66 | - | - |
| total | 161 | 212 | 134 | 240 | 232 | 261 | 215 | 145 | - | 1600 |

Results: Hyperparameter Tuning And Optimizer Analysis



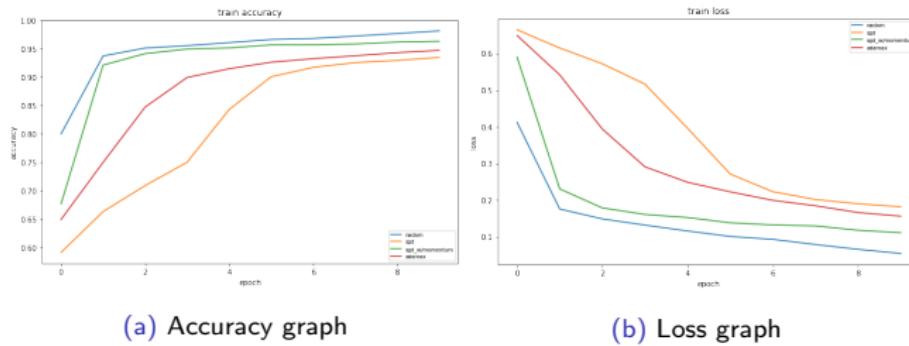
Model an accuracy of 96.43% with loss 0.0290 on the train set and accuracy of 95.58% with loss 0.0351 on the test set. Model with regularization had an accuracy of 93.48% with loss 0.2149 on the train set and accuracy of 93.65% with loss 0.2135 on the test set.

Results: Hyperparameter Tuning And Optimizer Analysis



Model at 100 epoch an accuracy of 95.28% for the train with accuracy of 69.85% for the test and with regularization had an accuracy of 93.40% for the train with accuracy of 71.00% for the test. Model at 300 epoch had an accuracy of 99.98% for the train with accuracy of 69.90% for the test and with regularization had an accuracy of 99.92% for the train with accuracy of 72.69% for the test.

Results: Hyperparameter Tuning And Optimizer Analysis



(a) Accuracy graph

(b) Loss graph

Figure: Further Optimizer Comparison Loss and Accuracy for the malaria Dataset Model on the Test samples

| train | MAcc | Mloss |
|--------|--------|--------|
| Nadam | 98.10% | 0.0545 |
| SGD | 93.44% | 0.1820 |
| Adamax | 94.69% | 0.1563 |
| SGDmom | 96.29% | 0.1112 |

Results: Hyperparameter Tuning And Optimizer Analysis

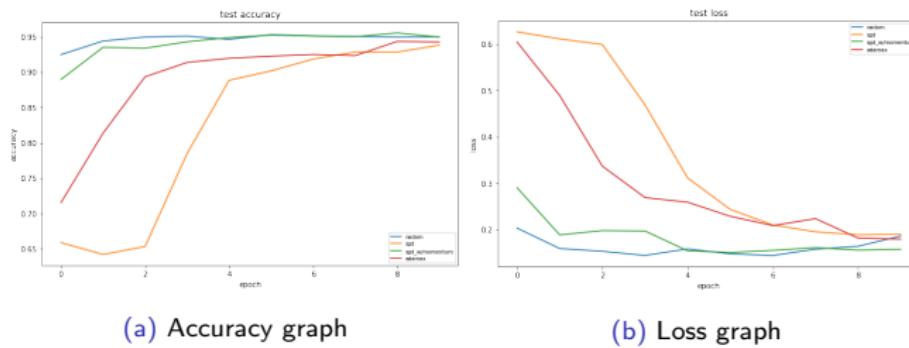
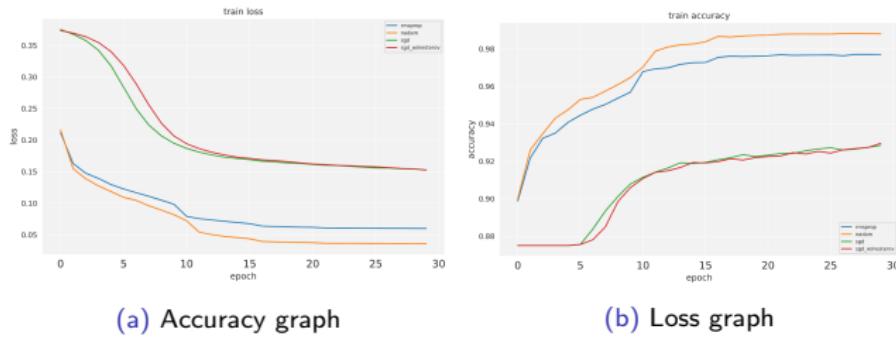


Figure: Further Optimizer Comparison Loss and Accuracy for the Malaria Dataset Model on the Test samples

| test | MAcc | Mloss |
|--------|--------|--------|
| Nadam | 94.99% | 0.1861 |
| SGD | 93.80% | 0.1900 |
| Adamax | 92.70% | 0.1794 |
| SGDmom | 94.96% | 0.1577 |

Results: Hyperparameter Tuning And Optimization Algorithms Analysis



(a) Accuracy graph

(b) Loss graph

Figure: Further Optimizer Comparison Loss and Accuracy for the Kvasir Dataset Model on the Test samples

| train | KAcc | Kloss |
|---------|--------|--------|
| Nadam | 98.80% | 0.0358 |
| SGD | 92.95% | 0.1527 |
| RMSprop | 97.70% | 0.0599 |
| SGDnest | 92.95% | 0.1526 |

Results: Hyperparameter Tuning And Optimization Algorithms Analysis

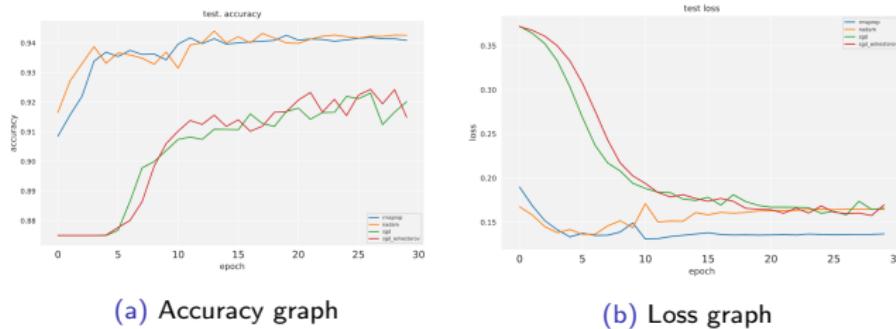


Figure: Further Optimizer Comparison Loss and Accuracy for the Kvasir Dataset Model on the Test samples

| test | KAcc | Kloss |
|---------|--------|--------|
| Nadam | 94.26% | 0.1648 |
| SGD | 92.02% | 0.1658 |
| RMSprop | 94.09% | 0.1368 |
| SGDnest | 91.49% | 0.1696 |

Conclusion

- Although the FCN is a good classifier, they are not good feature extractors but the CNN best quality is identifying and extracting the features from the images during training as seen in the malaria samples.
- For datasets with less samples, the choice of the hyperparameter or optimizer can help improve the performance of the network.
- A deeper CNN model with a combination of the best observable hyperparameters showed significant improvement with an accuracy of 94.99% on the malaria dataset and accuracy of 94.26% on the kvasir dataset for the CNN model.

Conclusion

- This suggest that in training more complex and deeper networks the adaptive methods still have a lot of improvements to be made in order to improve its generalization on the test samples.
- Also, further future work can be carried out by analyzing how higher order optimization algorithms can affect the performance of deep learning models on the medical imaging datasets.

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