

Screening Chest X-rays for Covid-19 with Deep Learning

Eric Robert Gill – June 2021

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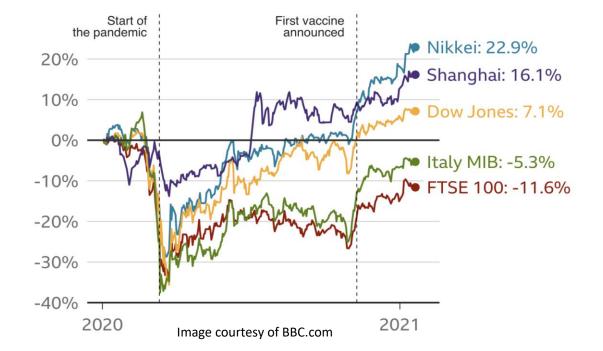
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Context – The Coronavirus

• 3,467,722 deaths since 31st December 2019¹.

uoc.edu

- Predicted economic losses of up to \$8.8 trillion worldwide².
- Market instability despite vaccination announcements.



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2. Baig et al., "Deaths, panic, lockdowns and US equity markets: The case of COVID-19 pandemic," Finance Research Letters, 2021

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Context - Prevention

- Strategies generally involve detection and isolation.
- Testing with PCR and antigens tests.
- Can be expensive¹, require resources.
- Deep learning and X-rays show potential to help^{2, 3, 4}.

1. Neilan et al. "Clinical Impact, Costs, and Cost-effectiveness of Expanded Severe Acute Respiratory Syndrome Coronavirus 2 Testing in Massachusetts," *Clinical Infectious Diseases* 2020

- 2. M. E. H. Chowdhury et al., "Can AI Help in Screening Viral and COVID-19 Pneumonia?," IEEE Access 2020
- 3. S. Minaee et al., "Deep-COVID: Predicting COVID-19 from chest X-ray images using deep transfer learning," *Medical Image Analysis* 2020
- 4. P. Kedia et al., "CoVNet-19: A Deep Learning model for the detection and analysis of COVID-19 patients," *Applied Soft Computing* 2020



Objectives

- Use deep learning to classify images of lungs into one of three classes: Covid-19, Normal or Viral Pneumonia.
- Investigate Convolutional Neural Networks and techniques to increase performance and compare results with baseline results found in literature.
- Investigate feasibility of applying deep learning to this kind of medical screening.
- Compare results obtained with results from a pretrained model using transfer-learning to fine-tune its parameters.



Medical Context

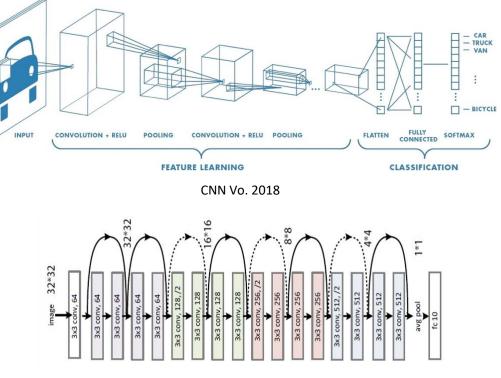
- Medical imaging heavily used in diagnosis.
- Chest x-rays (CXR) considerably cheaper than alternatives¹.
- Pneumonia detected in CXR with deep learning (DL)^{2, 3}.
- Several papers succeed in detecting Covid-19 in CXR with DL^{4, 5, 6}.

- 1. Wielputz et.al., Deutsches Arzteblatt International, 2014.
- 2. N. M. Elshennawy et.al. Deep-Pneumonia Framework Using Deep Learning Models Based on Chest X-Ray Images," *Diagnostics 2020*
- 3. T. Rahman et al., "Transfer learning with deep Convolutional Neural Network (CNN) for pneumonia detection using chest X-ray," *Applied Sciences (Switzerland)*, 2020
- 4. M. E. H. Chowdhury et al., "Can AI Help in Screening Viral and COVID-19 Pneumonia?," *IEEE Access* 2020
- 5. S. Minaee et al., "Deep-COVID: Predicting COVID-19 from chest X-ray images using deep transfer learning," *Medical Image Analysis* 2020
- 6. P. Kedia et al., "CoVNet-19: A Deep Learning model for the detection and analysis of COVID-19 Eric Robert Gill patients," *Applied Soft Computing* 2020



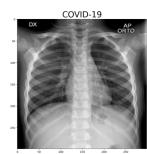
Techniques employed

- Convolutional Neural Network (CNN).
- Data Augmentation.
- Batch Normalization.
- L2 Regularization.
- Dropout.
- Transfer learning.

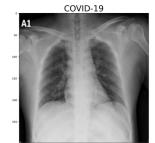


ResNet18, programmersought.com

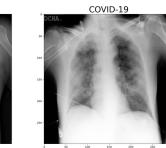
Data



Normal



Normal

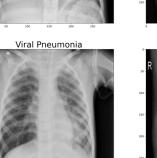


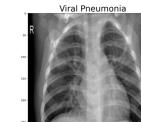
Normal



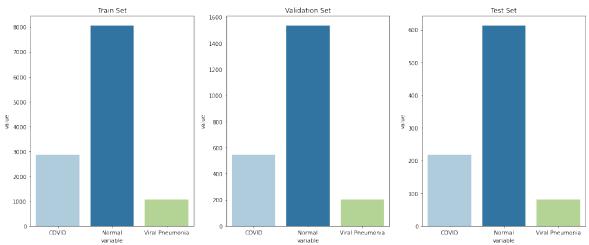
Viral Pneumonia







| Class | Number of images | Percentage of Total |
|-----------------|------------------|---------------------|
| COVID-19 | 3,616 | 23.86% |
| Normal | 10,192 | 67.26% |
| Viral Pneumonia | 1,345 | 8.88% |



8000 -7000 -6000 -5000 -

Experiments - 1

Experiment 1

| Layer type | Output | Kernel/Filter | Stride | Padding |
|-----------------|--------------|---------------|--------|---------|
| Convolutional | (295,295,32) | 5 | 1 | 0 |
| ReLU | (295,295,32) | N/A | N/A | N/A |
| Max Pooling | (147,147,32) | 2x2 | 1 | 0 |
| Convolutional | (138,138,64) | 5 | 1 | 0 |
| ReLU | (138,138,64) | N/A | N/A | N/A |
| Max Pooling | (69,69,64) | 2x2 | 1 | 0 |
| Convolutional | (50,50,128) | 5 | 1 | 0 |
| ReLU | (50,50,128) | N/A | N/A | N/A |
| Max Pooling | (25,25,128) | 2x2 | 1 | 0 |
| Flatten | (80,000, 1) | N/A | N/A | N/A |
| Fully Connected | (128, 1) | N/A | N/A | N/A |
| Output | (3, 1) | N/A | N/A | N/A |

Experiment 2

- Data Augmentation
 - Random Grayscale: 0.05p.
 - Random Vertical Flip: 0.08p.
 - Random Rotation: \pm 10 degrees.

Experiments - 2

Experiment 3

| Layer type | Output | Kernel/Filter | Stride | Padding |
|---------------------|--------------|---------------|--------|---------|
| Convolutional | (295,295,32) | 5 | 1 | 0 |
| BatchNorm (0.04) | (295,295,32) | N/A | N/A | N/A |
| ReLU | (295,295,32) | N/A | N/A | N/A |
| Max Pooling | (147,147,32) | 2x2 | 1 | 0 |
| Convolutional | (138,138,64) | 5 | 1 | 0 |
| BatchNorm (0.06) | (138,138,64) | N/A | N/A | N/A |
| ReLU | (138,138,64) | N/A | N/A | N/A |
| Max Pooling | (69,69,64) | 2x2 | 1 | 0 |
| Convolutional | (50,50,128) | 5 | 1 | 0 |
| BatchNorm (0.06) | (50,50,128) | N/A | N/A | N/A |
| ReLU | (50,50,128) | N/A | N/A | N/A |
| Max Pooling | (25,25,128) | 2x2 | 1 | 0 |
| Flatten | (80,000, 1) | N/A | N/A | N/A |
| Fully Connected | (128, 1) | N/A | N/A | N/A |
| Output | (3, 1) | N/A | N/A | N/A |

Experiment 4

- L2 Regularization
 - Weight decay set to 1⁻⁶.
- Data Augmentation
 - Random Grayscale: 0.05p.
 - Random Vertical Flip: 0.07p.
 - Random Rotation: \pm 8 degrees.

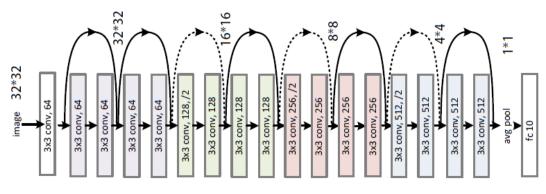


Experiment 5

| Layer type | Output | Kernel/Filter | Stride | Padding |
|---------------------|--------------|---------------|--------|---------|
| Convolutional | (295,295,32) | 5 | 1 | 0 |
| BatchNorm (0.04) | (295,295,32) | N/A | N/A | N/A |
| ReLU | (295,295,32) | N/A | N/A | N/A |
| Max Pooling | (147,147,32) | 2x2 | 1 | 0 |
| Convolutional | (138,138,64) | 5 | 1 | 0 |
| BatchNorm (0.06) | (138,138,64) | N/A | N/A | N/A |
| Dropout (0.08) | (138,138,64) | N/A | N/A | N/A |
| ReLU | (138,138,64) | N/A | N/A | N/A |
| Max Pooling | (69,69,64) | 2x2 | 1 | 0 |
| Convolutional | (50,50,128) | 5 | 1 | 0 |
| BatchNorm (0.06) | (50,50,128) | N/A | N/A | N/A |
| Dropout (0.10) | (50,50,128) | N/A | N/A | N/A |
| ReLU | (50,50,128) | N/A | N/A | N/A |
| Max Pooling | (25,25,128) | 2x2 | 1 | 0 |
| Flatten | (80,000, 1) | N/A | N/A | N/A |
| Fully Connected | (128, 1) | N/A | N/A | N/A |
| Output | (3, 1) | N/A | N/A | N/A |

Experiment 6

ResNet18



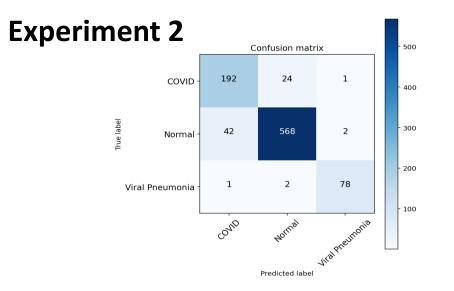
Courtesy of programmersought.com



Results - 1

Experiment 1 Confusion matrix 500 2 168 47 COVID 400 True label 0 29 300 Normal 200 7 73 1 Viral Pneumonia 100 COVID Normal Vital Preum Predicted label

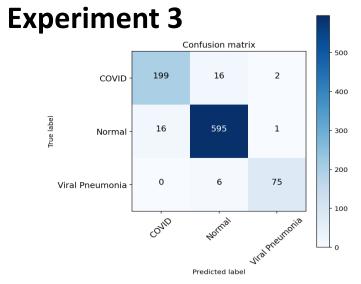
| | Precision | Recall | F1-score | Support |
|--------------|-----------|--------|----------|---------|
| COVID | 84.85% | 77.42% | 80.96% | 217 |
| Normal | 91.52% | 95.26% | 93.35% | 612 |
| Pneumonia | 97.33% | 90.12% | 93.59% | 81 |
| Accuracy | | | 90.55% | 910 |
| Macro avg | 91.23% | 87.60% | 89.30% | 910 |
| Weighted avg | 90.45% | 90.55% | 90.42% | 910 |



| | Precision | Recall | F1-score | Support |
|--------------|-----------|--------|----------|---------|
| COVID | 81.70% | 88.48% | 84.96% | 217 |
| Normal | 95.62% | 92.81% | 94.20% | 612 |
| Pneumonia | 96.30% | 96.30% | 96.30% | 81 |
| Accuracy | | | 92.09% | 910 |
| Macro avg | 91.21% | 92.53% | 91.82% | 910 |
| Weighted avg | 92.36% | 92.09% | 92.18% | 910 |



Results - 2



| | Precision | Recall | F1-score | Support |
|--------------|-----------|--------|----------|---------|
| COVID | 92.56% | 91.71% | 92.13% | 217 |
| Normal | 96.43% | 97.22% | 96.83% | 612 |
| Pneumonia | 96.15% | 92.59% | 94.34% | 81 |
| Accuracy | | | 95.49% | 910 |
| Macro avg | 95.05% | 93.84% | 94.43% | 910 |
| Weighted avg | 95.49% | 95.49% | 95.49% | 910 |

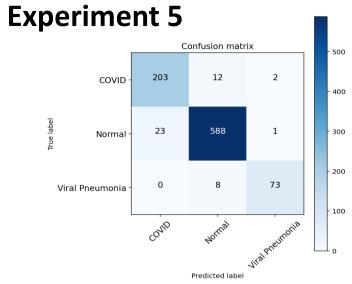
Experiment 4 Confusion matrix - 500 18 2 197 COVID 400 ue label 5 13 594 - 300 Normal - 200 0 4 77 Viral Pneumonia - 100 COND Jormal Viral Prieur

| | Precision | Recall | F1-score | Support |
|--------------|-----------|--------|----------|---------|
| COVID | 93.81% | 90.78% | 92.27% | 217 |
| Normal | 96.43% | 97.06% | 96.74% | 612 |
| Pneumonia | 91.67% | 95.06% | 93.33% | 81 |
| Accuracy | | | 95.38% | 910 |
| Macro avg | 93.97% | 94.30% | 94.12% | 910 |
| Weighted avg | 95.38% | 95.38% | 95.37% | 910 |

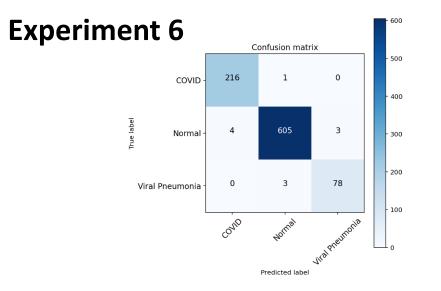
Predicted label



Results - 3



| | Precision | Recall | F1-score | Support |
|--------------|-----------|--------|----------|---------|
| COVID | 89.82% | 93.55% | 91.65% | 217 |
| Normal | 96.71% | 96.08% | 96.39% | 612 |
| Pneumonia | 96.05% | 90.12% | 92.99% | 81 |
| Accuracy | | | 94.95% | 910 |
| Macro avg | 94.20% | 93.25% | 93.68% | 910 |
| Weighted avg | 95.01% | 94.95% | 94.96% | 910 |



| | Precision | Recall | F1-score | Support |
|--------------|-----------|--------|----------|---------|
| COVID | 98.18% | 99.54% | 98.86% | 217 |
| Normal | 99.34% | 98.86% | 99.10% | 612 |
| Pneumonia | 96.30% | 96.30% | 96.30% | 81 |
| Accuracy | | | 98.79% | 910 |
| Macro avg | 97.94% | 98.23% | 98.08% | 910 |
| Weighted avg | 98.80% | 98.79% | 98.79% | 910 |

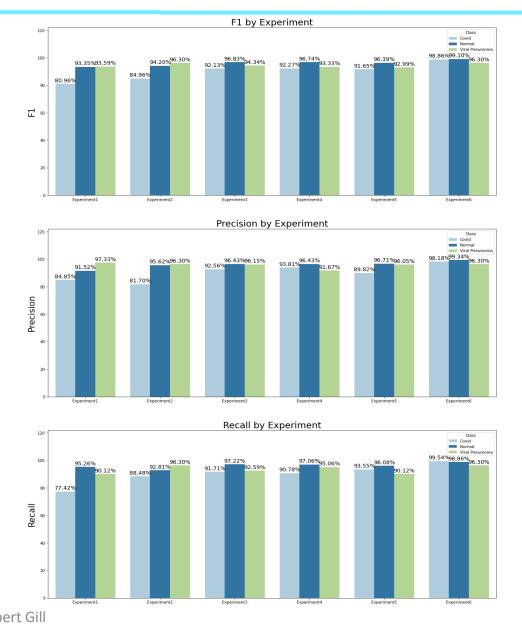
Discussion

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- Data augmentation improved recall on Covid class^{1, 2}.
- Batch Normalization improved three metrics on Covid class.
- L2 regularization produced higher precisions in Normal and Covid classes than preceding experiments.
- Dropout produced highest recall in Covid.
- Transfer Learning shows high potential¹.
- 1. S. Minaee et al., "Deep-COVID: Predicting COVID-19 from chest X-ray images using deep transfer learning," *Medical Image Analysis* 2020
- 2. P. Kedia et al., "CoVNet-19: A Deep Learning model for the detection and analysis of COVID-19 Eric Robert Gill patients," *Applied Soft Computing* 2020





Conclusions

- Compelling results suport hypothesis.
- High recall achieved for Covid and Viral Pneumonia.
- X-rays cheap, massively available.
- Further studies with more data are feasible.
- Transfer learning appears key.



Thank you for your time