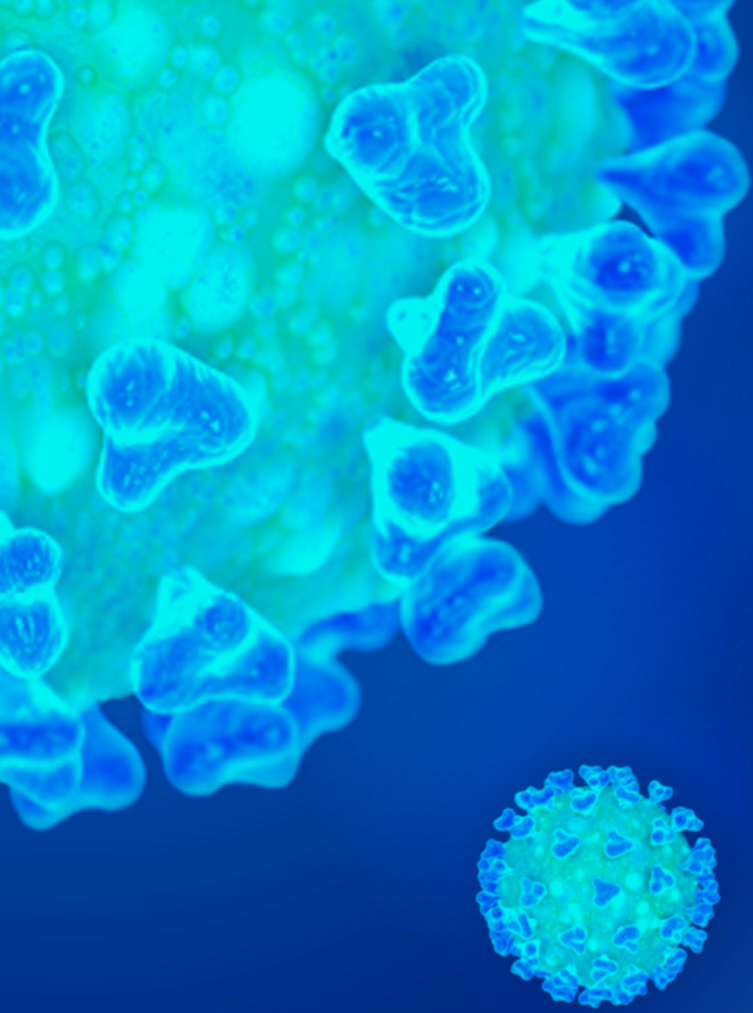


The background features a dark blue gradient with several stylized representations of COVID-19 virus particles. These particles are depicted as spherical structures with a textured, bumpy surface, colored in shades of light blue and white. Some particles are larger and more prominent, while others are smaller and scattered. Additionally, there are larger, more complex structures that resemble lung cells or alveoli, also in light blue and white, with irregular, rounded shapes and internal details. The overall composition is clean and modern, with a focus on the scientific theme of the presentation.

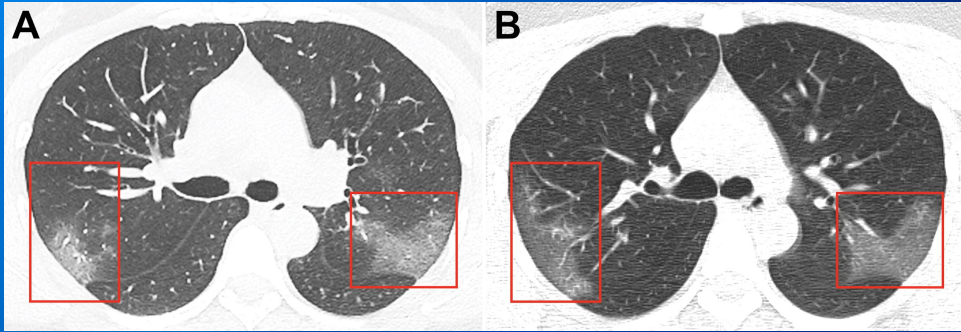
# COVID-19

Classification based on Lung CT Scans

By: Lin Junyuan | DSI-14

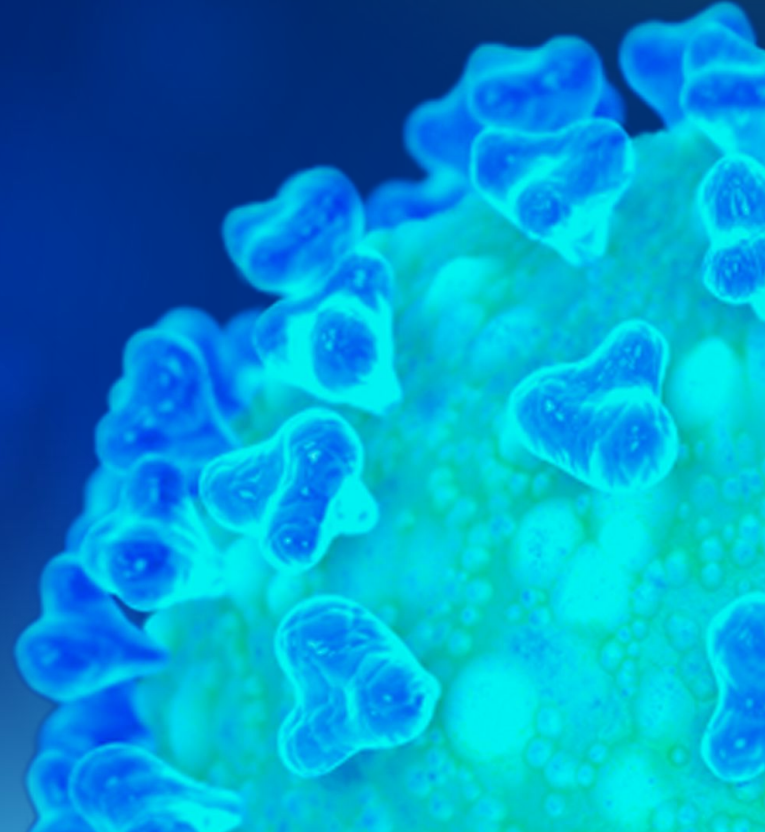
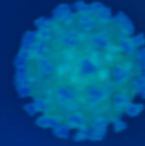


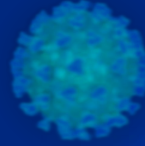
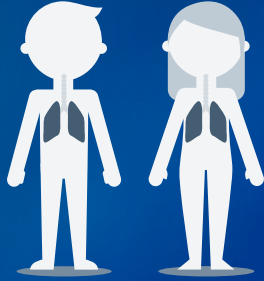
# INTRODUCTION



Credit: CT Imaging of the 2019 Novel Coronavirus (2019-nCoV) Pneumonia  
Junqiang Lei, Junfeng Li, Xun Li, Xiaolong Qi

- Ground Glass Opacities that grows with severity





# Problem Statement

To develop a Convolutional Neural Network-based classification(CNN) model, based on Chest CT Scan data in the axial view, to determine whether Covid infection is present, with a targeted performance of **80%** for both **accuracy** and **sensitivity**.

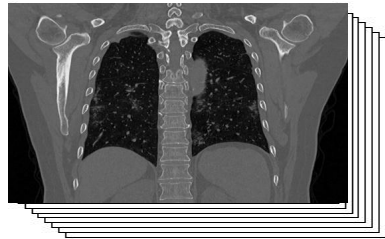


# Data preprocessing

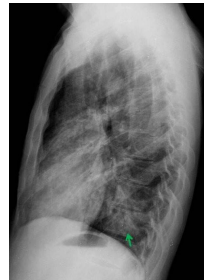
Gathered images from 6 sources, coming in a variety of file formats, views and image resolutions



Axial view  
.jpg/.png format



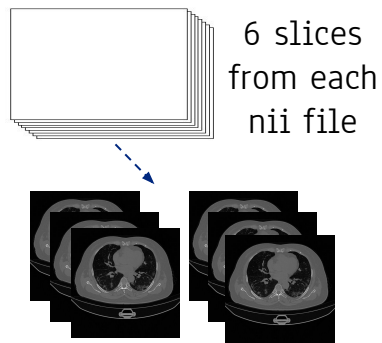
Frontal view  
.nii format



Supine view  
\*This is actually X-ray

# Preprocessing Steps

## Load Image



.jpg/.png

## Process image

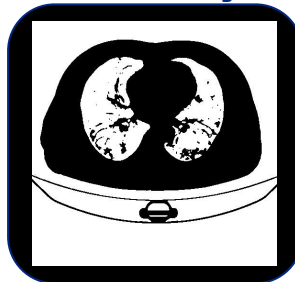
1. Median denoising



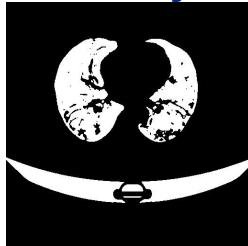
2. Histogram equalization



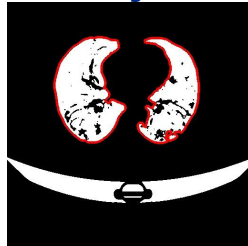
3. Thresholding



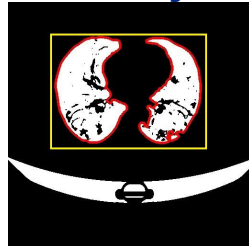
4. Remove background



5. Find Lung contours

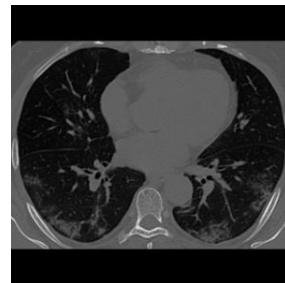


6. Create bounding rectangle

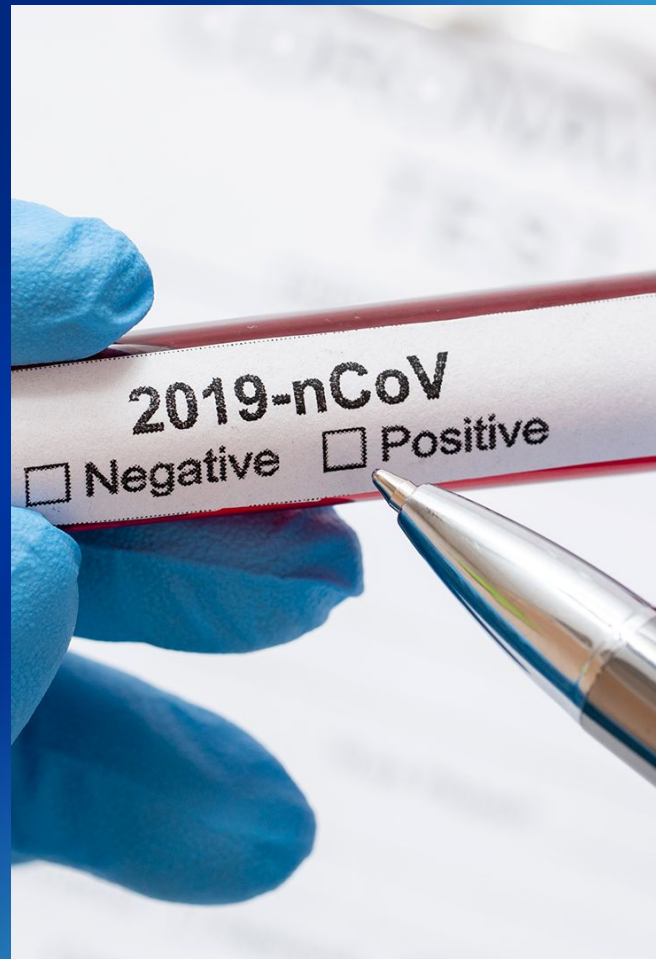
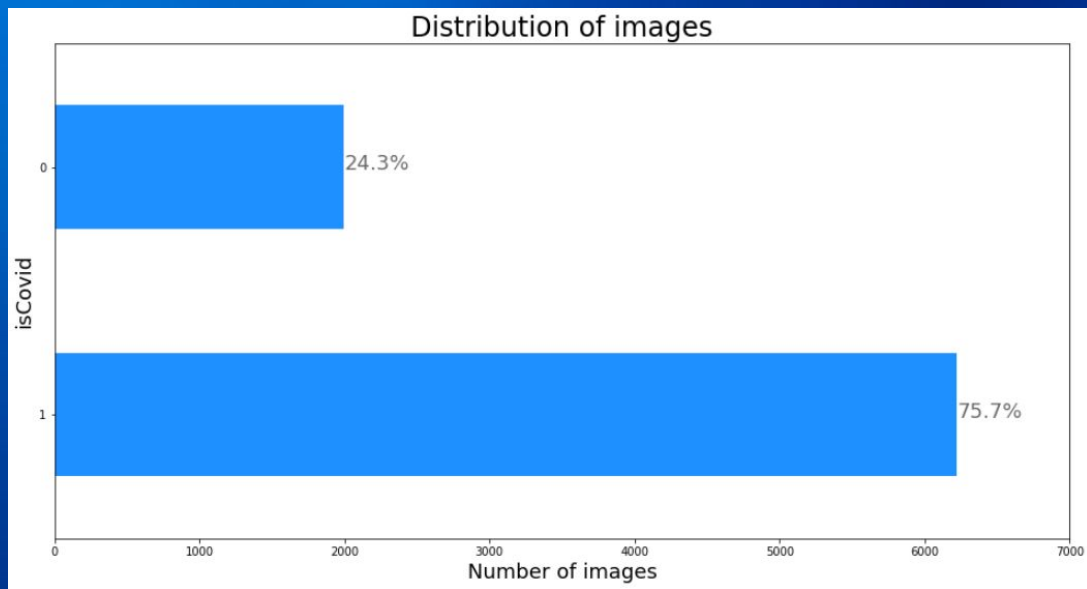


## Output image

- Crop original image using bounded region
- Zero pad and resize to (224, 280)



# Data distribution



# Data preparation

## STEP 01 - Train-Test-Split

- There are multiple images from each patient.
- Will need to split on patientID to prevent data leakage

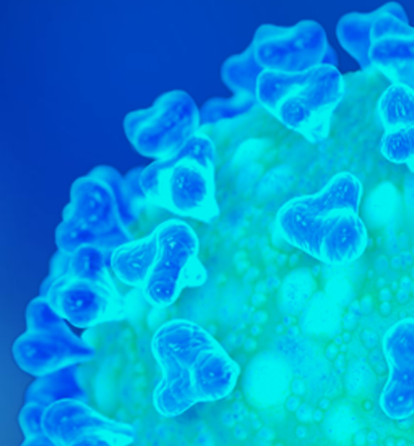
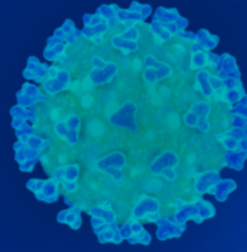
## STEP 02 - Data augmentation

- Keras / ImageDataGenerator
- Randomization Parameters
  - Horizontal shift +/- 5%
  - Vertical shift +/- 5%
  - Horizontal flip
  - fill\_mode='nearest',
  - Brightness variation +/- 20%





# MODELING



# Initialize model structure

Conv2d (filters = 32, kernel size = 3, 'relu')
Conv2d (filters = 64, kernel size = 3, 'relu')
Max Pooling (size = (2,2))
Dropout (ratio=0.25)
Conv2d (filters = 32, kernel size = 3, 'relu')
Max Pooling (size = (2,2))
Global Average Pooling
Dense (filters=256, 'relu')
Dense (filters=128, 'relu')
Dropout (ratio=0.5)
Dense (filters=64, 'relu')
Dense (filters=1, 'sigmoid')

Model: "model1\_1"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 222, 278, 32)	320
conv2d_1 (Conv2D)	(None, 220, 276, 64)	18496
max_pooling2d (MaxPooling2D)	(None, 110, 138, 64)	0
dropout (Dropout)	(None, 110, 138, 64)	0
conv2d_2 (Conv2D)	(None, 108, 136, 32)	18464
max_pooling2d_1 (MaxPooling2D)	(None, 54, 68, 32)	0
global_average_pooling2d (GlobalAveragePooling2D)	(None, 32)	0
dense (Dense)	(None, 256)	8448
dense_1 (Dense)	(None, 128)	32896
dropout_1 (Dropout)	(None, 128)	0
dense_2 (Dense)	(None, 64)	8256
dense_3 (Dense)	(None, 1)	65

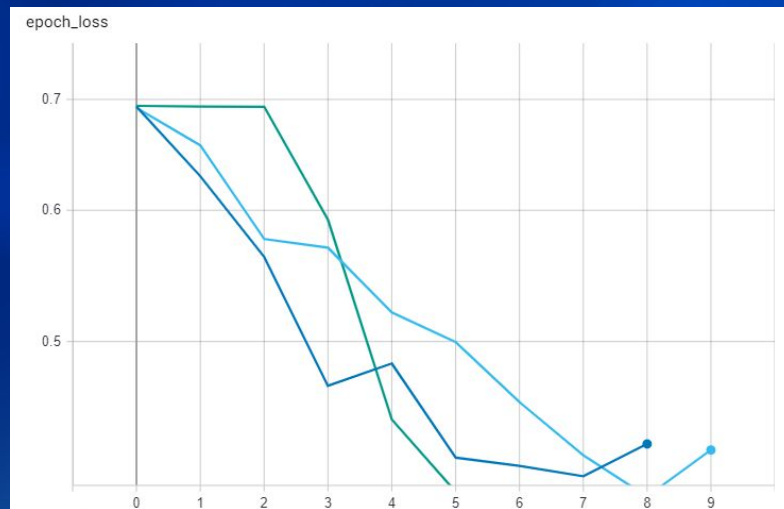
Total params: 86,945

Trainable params: 86,945

Non-trainable params: 0

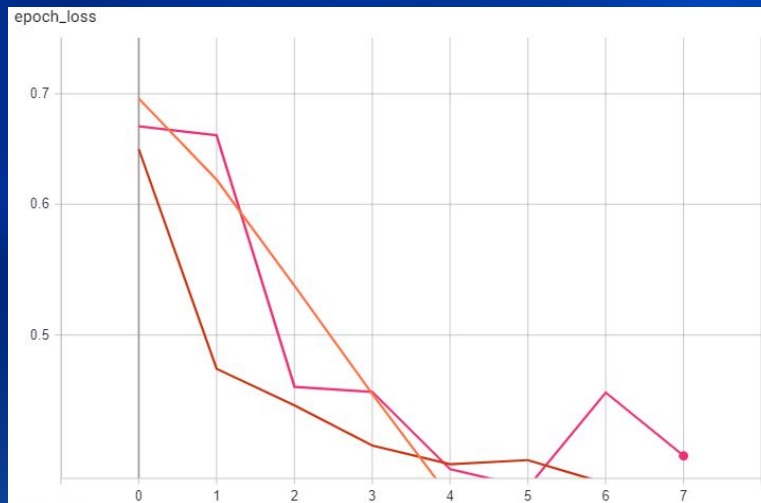
# Hyperparameter tuning [Number of filters]

Initial model | model with half number of filters | model with twice number of filters



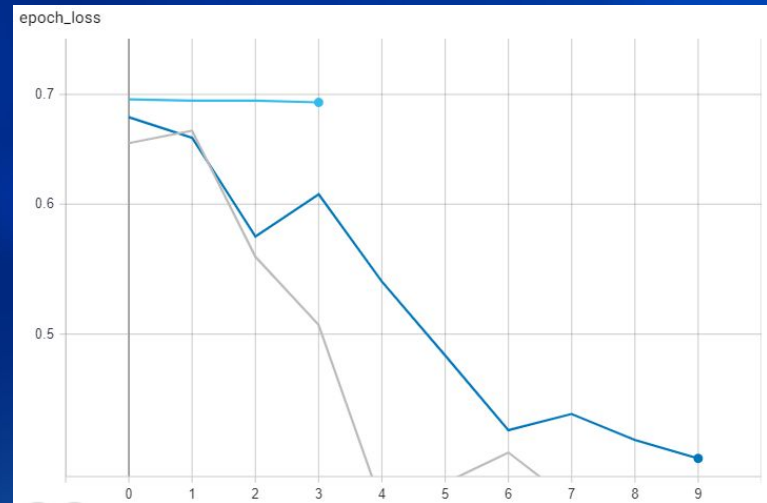
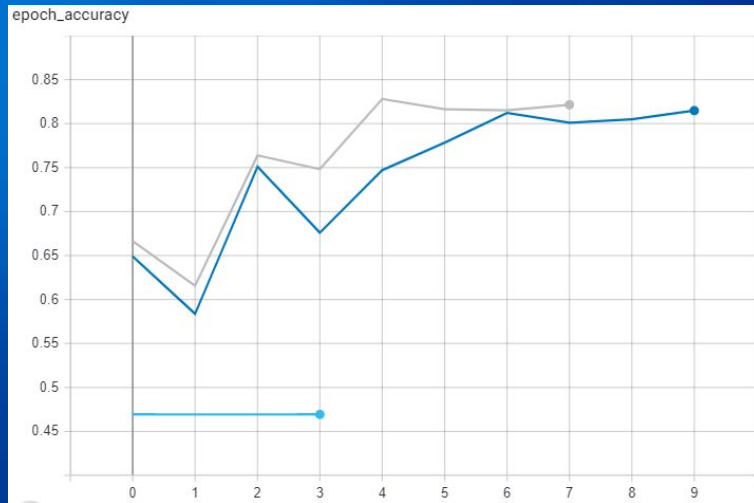
# Hyperparameter tuning [Size of kernels]

Initial model (size:3) | model with kernel size: 5 | model with kernel size: 7



# Hyperparameter tuning [Number of layers]

Initial model | model with 1 less conv layer and 1 less dense layer |  
model with 1 more conv layer and 1 more dense layer





# Model evaluation

	model_name	true_negative	false_positive	false_negative	true_positive	accuracy	sensitivity	specificity
0	model1_1	1027	231	337	955	0.78	0.74	0.82
1	model1_2	866	392	74	1218	0.82	0.94	0.69
2	model1_3	947	311	105	1187	0.84	0.92	0.75
3	model2_1	896	395	63	1208	0.82	0.95	0.69
4	model2_2	853	438	32	1239	0.82	0.97	0.66
8	model2_3	917	404	50	1206	0.82	0.96	0.69
5	model3_1	920	321	135	1178	0.82	0.90	0.74
6	model3_2	941	300	173	1140	0.81	0.87	0.76
7	model3_3	1241	0	1313	0	0.49	0.00	1.00

# Selected model structure

Conv2d (filters = 64, kernel size = 5, 'relu')
Conv2d (filters = 128, kernel size = 5, 'relu')
Max Pooling (size = (2,2))
Dropout (ratio=0.25)
Conv2d (filters = 64, kernel size = 5, 'relu')
Max Pooling (size = (2,2))
Global Average Pooling
Dense (filters=512, 'relu')
Dense (filters=256, 'relu')
Dropout (ratio=0.5)
Dense (filters=128, 'relu')
Dense (filters=1, 'sigmoid')

Model: "model\_selected"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 220, 276, 64)	1664
conv2d_1 (Conv2D)	(None, 216, 272, 128)	204928
max_pooling2d (MaxPooling2D)	(None, 108, 136, 128)	0
dropout (Dropout)	(None, 108, 136, 128)	0
conv2d_2 (Conv2D)	(None, 104, 132, 64)	204864
max_pooling2d_1 (MaxPooling2D)	(None, 52, 66, 64)	0
global_average_pooling2d (GlobalAveragePooling2D)	(None, 64)	0
dense (Dense)	(None, 512)	33280
dense_1 (Dense)	(None, 256)	131328
dropout_1 (Dropout)	(None, 256)	0
dense_2 (Dense)	(None, 128)	32896
dense_3 (Dense)	(None, 1)	129

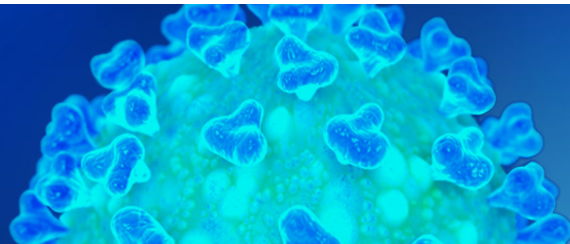
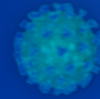
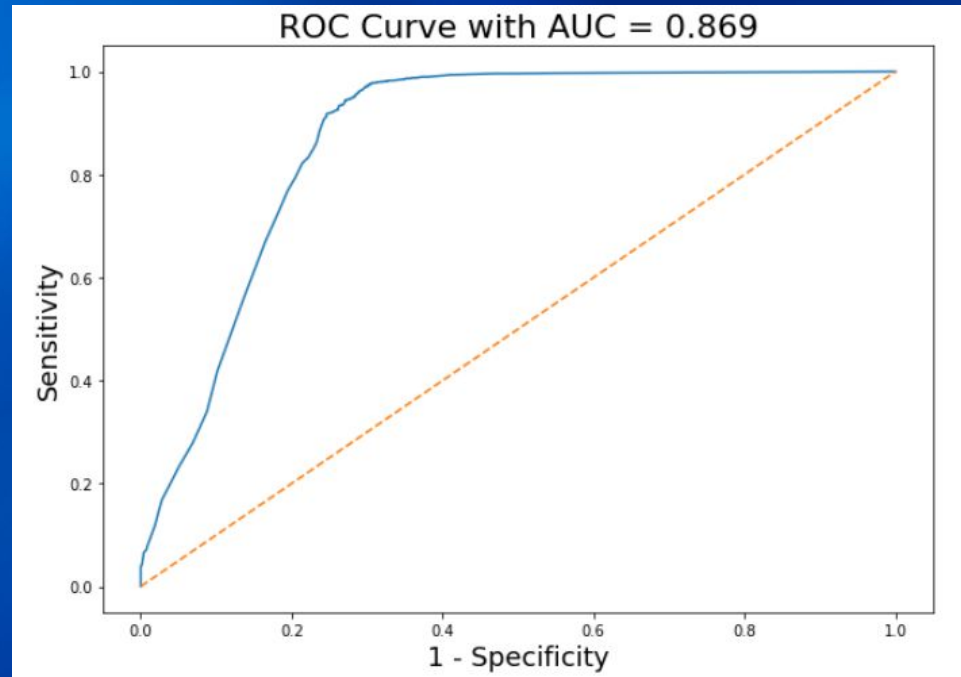
Total params: 609,089

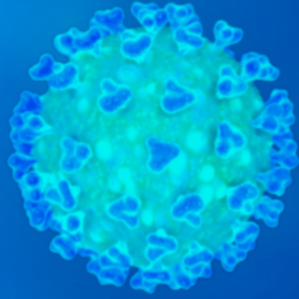
Trainable params: 609,089

Non-trainable params: 0

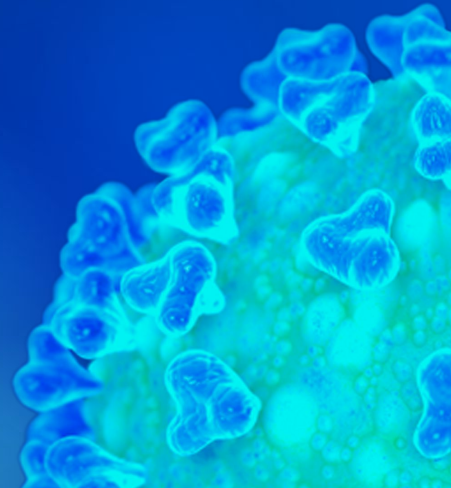
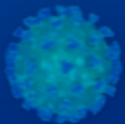
## Selected model performance

	model_name	true_negative	false_positive	false_negative	true_positive	accuracy	sensitivity	specificity
0	model1_1	1027	231	337	955	0.78	0.74	0.82
1	model1_2	866	392	74	1218	0.82	0.94	0.69
2	model1_3	947	311	105	1187	0.84	0.92	0.75
3	model2_1	896	395	63	1208	0.82	0.95	0.69
4	model2_2	853	438	32	1239	0.82	0.97	0.66
8	model2_3	917	404	50	1206	0.82	0.96	0.69
5	model3_1	920	321	135	1178	0.82	0.90	0.74
6	model3_2	941	300	173	1140	0.81	0.87	0.76
7	model3_3	1241	0	1313	0	0.49	0.00	1.00
9	model_selected	947	374	61	1195	0.83	0.95	0.72





# Conclusion and Recommendations





# Conclusions



Conv2d (filters = 64, kernel size = 5, 'relu')
Conv2d (filters = 128, kernel size = 5, 'relu')
Max Pooling (size = (2,2))
Dropout (ratio=0.25)
Conv2d (filters = 64, kernel size = 5, 'relu')
Max Pooling (size = (2,2))
Global Average Pooling
Dense (filters=512, 'relu')
Dense (filters=256, 'relu')
Dropout (ratio=0.5)
Dense (filters=128, 'relu')
Dense (filters=1, 'sigmoid')

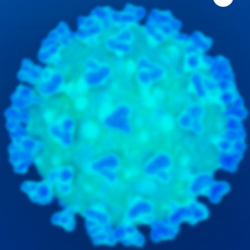
- The selected model was able to perform to our targeted values of minimum 80% for both accuracy and sensitivity.
- It has attained an ROC-AUC score of 0.86.

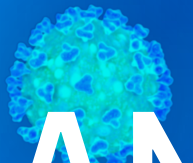
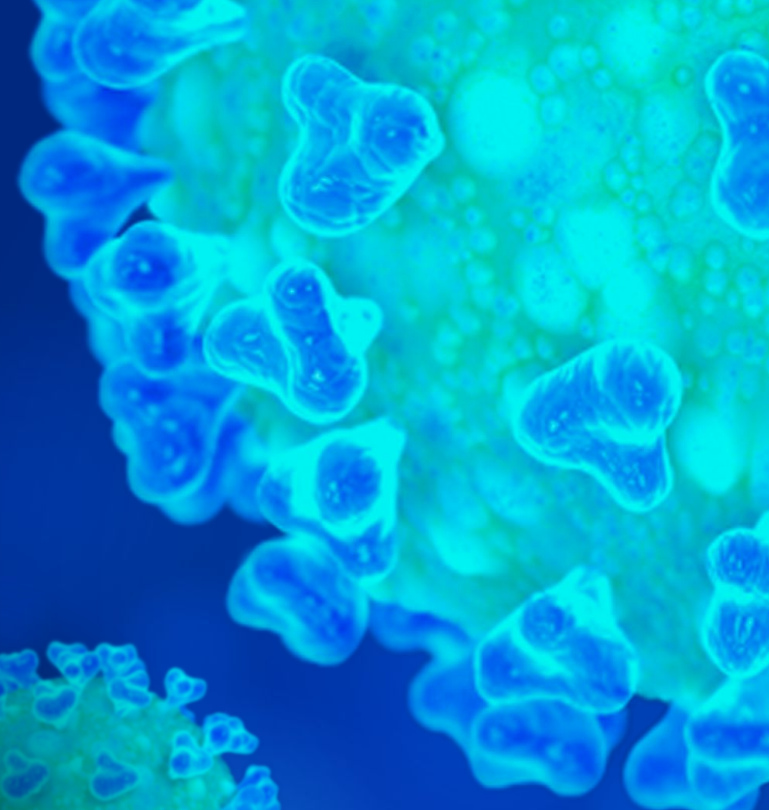


# Recommendations



- To continue to build up the database  
(More quality data -> Better performance)
- To do Multi-Class classification to separate between Covid infection and other types of viral/bacterial infection
- To try other methodologies, i.e. Transfer Learning, 3D CT Scan CNNs





# THANKS!



<https://www.linkedin.com/in/junyuanlin/>

DID YOU KNOW?

CNN was inspired by  
research done on cat brains.

“Receptive Fields of Single Neurons in the Cat’s Striate Cortex”  
conducted by Hubel and Wiesel

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