# COVID-19 CLASSIFICATION USING CHEST X-RAY IMAGES

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#### **PROJECT OBJECTIVE**

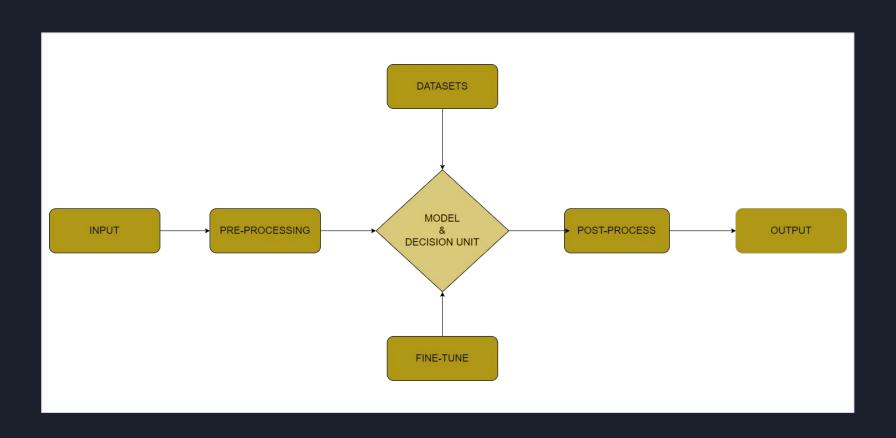
 The aim of the project is to develop a novel deep neural network based model for highly accurate detection of COVID-19 infection from the chest X- Ray images of the patients.

 The aim of our project is to make use of modern AI techniques to detect the COVID-19 patients using X-Ray images in an automated manner, particularly in settings where radiologists are not available, and help make the proposed testing technology scalable

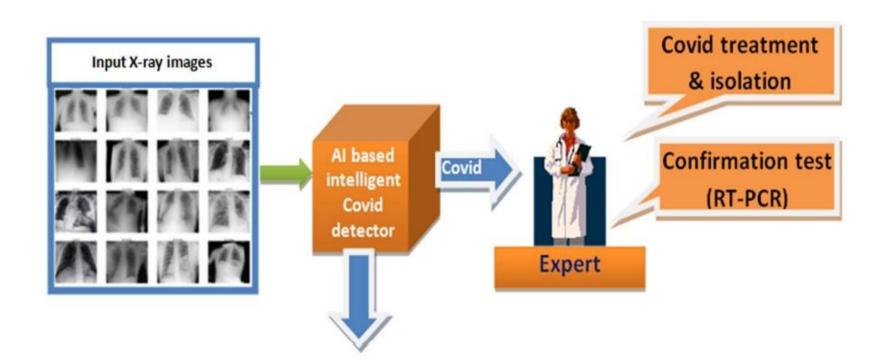
#### **FEATURES OF PROJECT**

- Early diagnosis is very essential for both early intervention to the patient and prevent the risk of transmission of the disease. In this Project we have presented a neural network architecture which will reduce overfitting.
- It can also be used in situation where the possibilities are insufficient whether in terms RT-PCR test or doctor. We have used a fuzzy neural network instead of taking a fully connected layer for the feature information.
- We all know that Convolutional neural network is an excellent feature extractor so integrating it with the fuzzy neural network will increase its accuracy in tests.

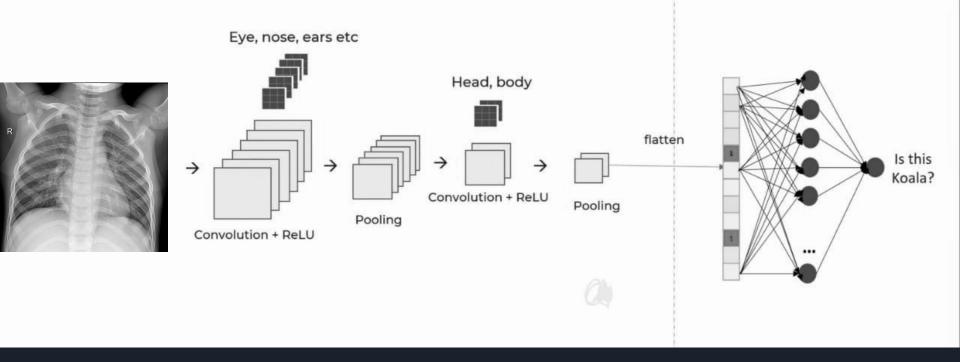
## **BLOCK DIAGRAM**

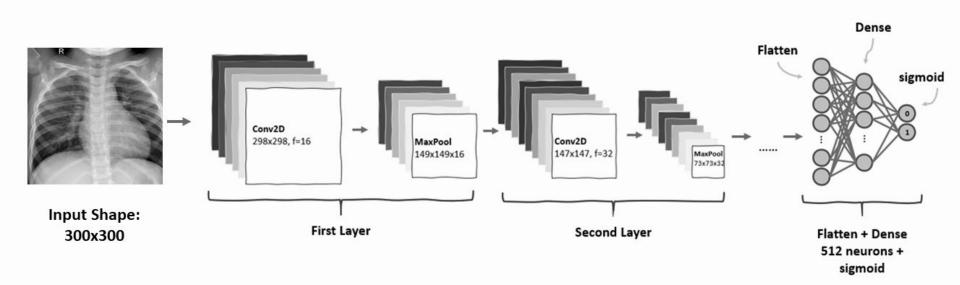


#### **Proposed Systems Architecture**

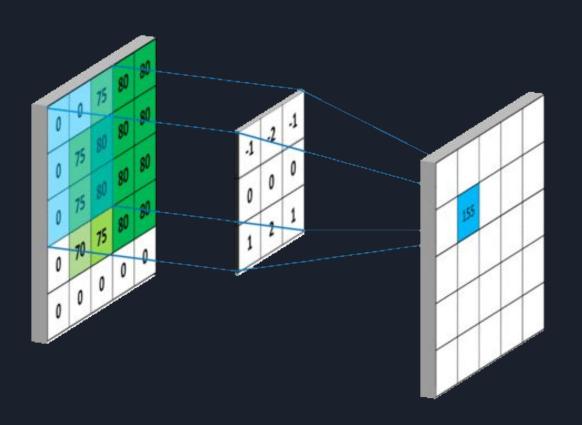


## **CLASSIFICATION USING CNN**

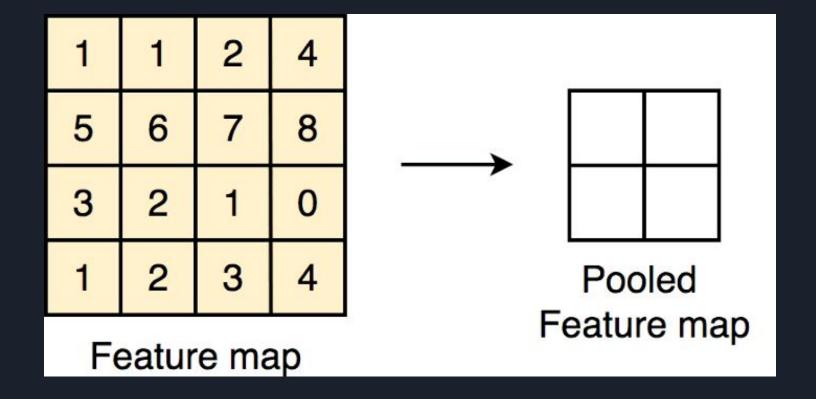




# **Convolutional Operation**

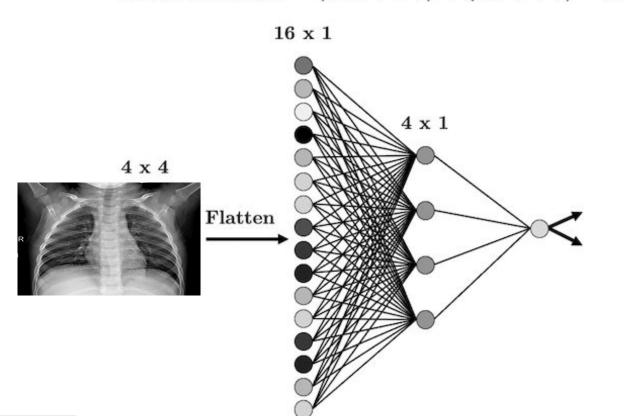


# MAX POOLING



## **CLASSIFICATION**

Total Parameters = 
$$(16 \times 4 + 4) + (4 \times 1 + 1) = 73$$



#### **CLASSIFICATION USING FUZZY LOGIC**

- Fuzzy logic is a method that was introduced to handle a range of values between TRUE and FALSE. This model of reasoning has helped us to reach closer to human reasoning. Most of the models which work on crisp data set only give output in one of the two forms: False (Value 0) or True (Value 1), Negative (Value 0) or Positive (Value 1) but what if there are other possibilities like partially True or partially False.
- A model which can produce the following possible decisions between FALSE and TRUE: Surely True, might be True, Not Sure about True or False, Might Be False, surely False. All these decisions can be achieved with the help of Fuzzy logic.
  - ➤ A. Fuzzy Based Image Processing
  - B. Fuzzy Based Classification on X-ray Images

## Fuzzy Based Classification on X-ray Images

#### **Feature Extraction:**

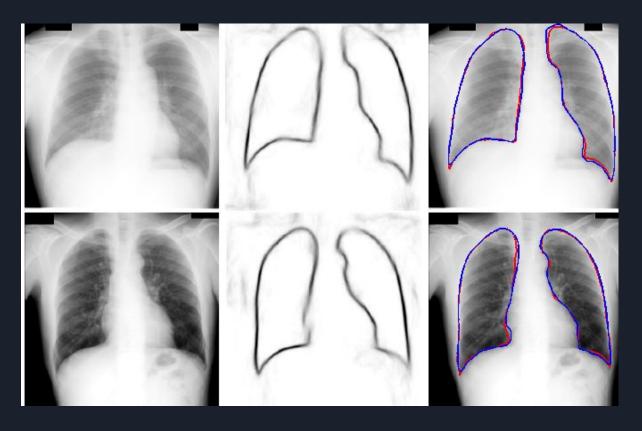
- In pattern classification, we use features like color, shape, spatial location, and texture. It is not suitable to consider color and texture features for x-ray image classification. The reason is that x-ray images are gray-scale images, and their texture characteristics are remarkably similar. It is recommended to use combinations of texture and shape features together.
- Histogram Adjustment
- Noise Removal
- Edge Detection



(a) Input Image



(b) Histogram Image



Edge Detection

## **System Implementation and Algorithm**

We have taken datasets that are publicly available. As we are doing binary classification, we have taken positive covid 19 X-ray images from the GitHub repository which is being updated by a group of doctors on a regular basis. As we wanted a balanced dataset therefore, we have taken the same number of normal X-ray images from Kaggle as we have taken from GitHub repository.

Link: <u>GitHub - ieee8023/covid-chestxray-dataset: We are building an open</u> <u>database of COVID-19 cases with chest X-ray or CT images.</u>

	import of import	os athlik shuti rando math numpy matpl keras eras.l eras.m eras.f ensorf	o impo il om / as r lotlik s layers models prepro Flow.	np po.pyp s imp ocess keras	olot oort oort sing s.pre	* * import image processing imp	port i	image as imager									
		FILE_PATH = Path('chestxray/metadata.csv').absolute() MAGE_PATH = Path('chestxray/images/').absolute()															
In [5]:	df = pd.read_csv(FILE_PATH)																
In [6]:	df.head	l()															
Out[6]:	patie	ntid	offset	sex	age	f	inding	RT_PCR_positive	survival	intubated	intubation_present	went_ic	u	date	location	folder	
	0	2	0.0	M	65.0	Pneumonia/Viral/C	OVID- 19	Υ	Y	N	N	1	٧	January 22, 2020	Cho Ray Hospital, Ho Chi Minh City, Vietnam	images	20
	1	2	3.0	M	65.0	Pneumonia/Viral/C	OVID- 19	Y	Y	N	N	1	٧	January 25, 2020	Cho Ray Hospital, Ho Chi Minh City, Vietnam	images	20
	2	2	5.0	M	65.0	Pneumonia/Viral/C	OVID-	Υ	Υ	N	N	1	۷	January 27, 2020	Cho Ray Hospital, Ho Chi Minh City,	images	20
														2020	Vietnam		

Cho Ray

```
In [3]: model = Sequential()
        model.add(Conv2D(32, kernel_size=(3, 3), activation='relu', input_shape=(224, 224, 3)))
        model.add(Conv2D(64, kernel_size=(3, 3), activation='relu'))
        model.add(MaxPooling2D(pool size=(2, 2)))
        model.add(Dropout(0.25))
        model.add(Conv2D(64, kernel size=(3, 3), activation='relu'))
        model.add(MaxPooling2D(pool size=(2, 2)))
        model.add(Dropout(0.25))
        model.add(Conv2D(128, kernel size=(3, 3), activation='relu'))
        model.add(MaxPooling2D(pool size=(2, 2)))
        model.add(Dropout(0.25))
        model.add(Flatten())
        model.add(Dense(64, activation='relu'))
        model.add(Dropout(0.5))
        model.add(Dense(1, activation='sigmoid'))
        model.compile(loss=keras.losses.binary_crossentropy, optimizer='adam', metrics=['accuracy'])
        model.summary()
        Model: "sequential"
                                     Output Shape
         Layer (type)
                                                                Param #
         conv2d (Conv2D)
                                     (None, 222, 222, 32)
                                                                896
```

```
conv2d 1 (Conv2D)
                           (None, 220, 220, 64)
                                                     18496
max pooling2d (MaxPooling2D (None, 110, 110, 64)
dropout (Dropout)
                          (None, 110, 110, 64)
                                                      0
conv2d 2 (Conv2D)
                           (None, 108, 108, 64)
                                                      36928
max pooling2d 1 (MaxPooling (None, 54, 54, 64)
                                                      0
2D)
dropout_1 (Dropout)
                          (None, 54, 54, 64)
                                                      0
conv2d 3 (Conv2D)
                           (None, 52, 52, 128)
                                                      73856
max pooling2d 2 (MaxPooling (None, 26, 26, 128)
                                                      0
2D)
```

# **RESULTS**





# **Conclusion**



Early diagnosis is very essential for both early intervention to the patient and prevent the risk of transmission of the disease. In this Project we have presented a neural network architecture which will reduce overfitting. It can also be used in situation where the possibilities are insufficient whether in terms RT-PCR test or doctor.



# Thank You !!!

ANY QUESTIONS?