

# COVID-19 SCAN DETECTION USING CNN

Vision and Perception course - Project  
presentation



SAPIENZA  
UNIVERSITÀ DI ROMA

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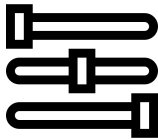
# How the presentation is organized



Section 1 - The Chosen Dataset & Background



Section 2 - The PreTrained CNN Network - ResNet101



Section 3 - Our custom CNN Network

# How the presentation is organized



## **Section 1 - The Chosen Dataset & Background**

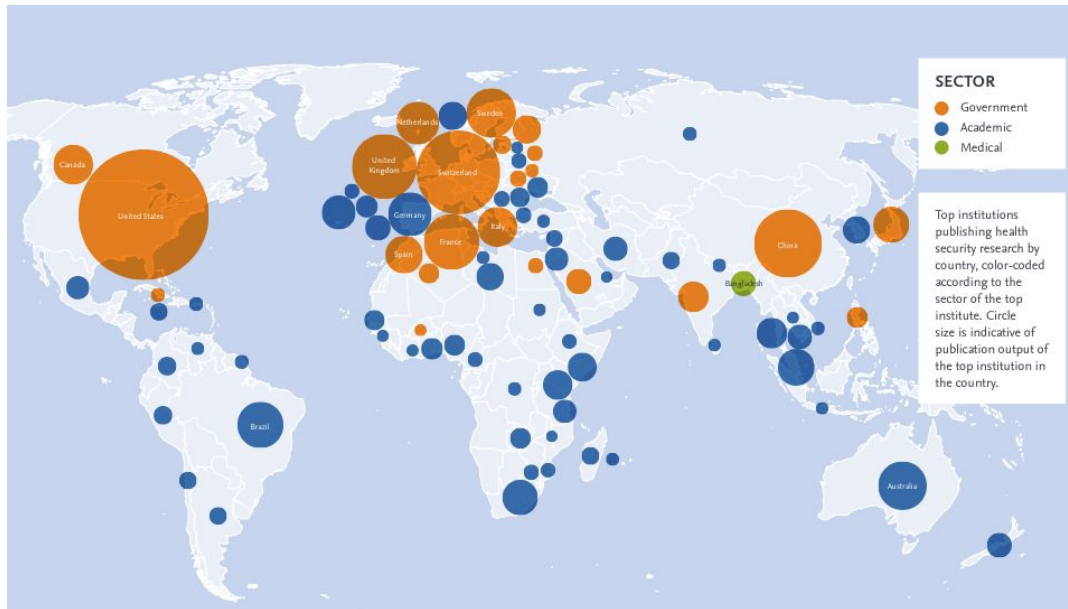


## Section 2 - The PreTrained CNN Network - ResNet101



## Section 3 - Our custom CNN Network

# Section 1 - The Community Effort



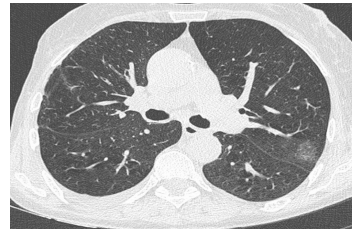
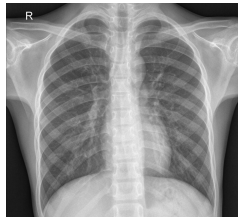
[elsevier.com/connect/infographic-global-research-trends-in-infectious-disease](https://elsevier.com/connect/infographic-global-research-trends-in-infectious-disease)

**researchers** worldwide have been working to advance our understanding of these epidemics and how to respond

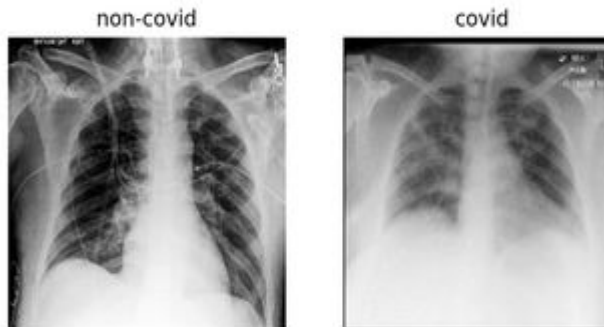
<input type="checkbox"/> 2020	78,857
<input type="checkbox"/> 2019	1,658
<input type="checkbox"/> 2018	1,447
<input type="checkbox"/> 2017	1,475

# Section 1 - The task

A fast **classification** of the radiological imaging such as chest **X-ray** and chest CT-scan can be helpful to **isolate** the infected persons timely and control this epidemic situation



## binary classification

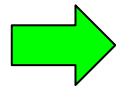
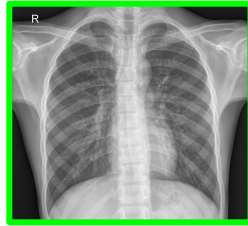


## multiclass classification

Type	Genus or Species
Viral	COVID-19 (SARSr-CoV-2)
	SARS (SARSr-CoV-1)
	Varicella
	Influenza
Bacterial	<i>Streptococcus</i> spp.
	<i>Klebsiella</i> spp.
	<i>Escherichia coli</i>
	<i>Mycoplasma</i> spp.
	<i>Legionella</i> spp.
	Unknown
Fungal	<i>Chlamydomphila</i> spp.
	<i>Pneumocystis</i> spp.
Lipoid	Non applicable

# Section 1 - The task

A fast **classification** of the radiological imaging such as chest **X-ray** and chest CT-scan can be helpful to **isolate** the infected persons timely and control this epidemic situation



**binary** classification



**multiclass** classification

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	<i>Pneumocystis</i> spp.
Lipoid	Non applicable

# Section 1 - The task



Table 4

Comparison of the proposed COVID-19 diagnostic method with other deep learning methods developed using radiology images.

Study	Type of Images	Number of Cases	Method Used	Accuracy (%)
Ioannis et al. [43]	Chest X-ray	224 COVID-19(+) 700 Pneumonia 504 Healthy	VGG-19	93.48
Wang and Wong [42]	Chest X-ray	53 COVID-19(+) 5526 COVID-19 (-) 8066 Healthy	COVID-Net	92.4
Sethy and Behra [45]	Chest X-ray	25 COVID-19(+) 25 COVID-19 (-)	ResNet50+ SVM	95.38
Hemdan et al. [41]	Chest X-ray	25 COVID-19(+) 25 Normal	COVIDX-Net	90.0
Narin et al. [44]	Chest X-ray	50 COVID-19(+) 50 COVID-19 (-)	Deep CNN ResNet-50	98
Ying et al. [46]	Chest CT	777 COVID-19(+) 708 Healthy	DRE-Net	86
Wang et al. [47]	Chest CT	195 COVID-19(+) 258 COVID-19(-)	M-Inception	82.9
Zheng et al. [48]	Chest CT	313 COVID-19(+) 229 COVID-19(-)	UNet+3D Deep Network	90.8
Xu et al. [49]	Chest CT	219 COVID-19(+) 224 Viral pneumonia 175 Healthy	ResNet + Location Attention	86.7
<b>Proposed Study</b>	Chest X-ray	125 COVID-19(+) 500 No-Findings	DarkCovidNet	98.08
		125 COVID-19(+) 500 Pneumonia 500 No-Findings		87.02



# Section 1 - The Chosen Dataset (1)

## COVID-19 Image Data Collection

Joseph Paul Cohen<sup>1,2</sup> Paul Morrison<sup>3</sup> Lan Dao<sup>4</sup>

### Abstract

This paper describes the initial COVID-19 open image data collection. It was created by assembling medical images from websites and publications and currently contains 123 frontal view X-rays.

### 1. Motivation

In the context of a COVID-19 pandemic, it is crucial to streamline diagnosis. Data is the first step to developing any diagnostic tool or treatment. While there exist large public datasets of more typical chest X-rays (Wang et al., 2017; Bustos et al., 2019; Irvin et al., 2019; Johnson et al., 2019; Demner-Fushman et al., 2016), there is no collection of COVID-19 chest X-rays or CT scans designed to be used for computational analysis.

In this paper, we describe the public database of pneumonia cases with chest X-ray or CT images, specifically COVID-19 cases as well as MERS, SARS, and ARDS. Data will be collected from public sources in order not to infringe patient confidentiality. Example images shown in Figure 1.

Our team believes that this database can dramatically improve identification of COVID-19. Notably, this would provide essential data to train and test a Deep Learning-based system, likely using some form of transfer learning. These tools could be developed to identify COVID-19 characteristics as compared to other types of pneumonia or in order to predict survival.

Currently, all images and data are released under the following URL: <https://github.com/leee8023/covid-chestxray-dataset>. As stated above, images collected have already been made public.

### 2. Expected outcome

This dataset can be used to study the progress of COVID-19 and how its radiological findings vary from other types of

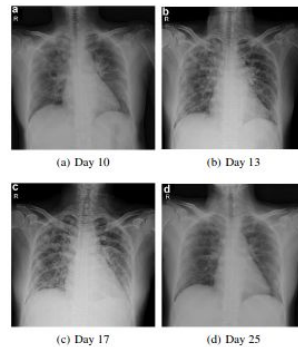


Figure 1. Example images from the same patient (#19) extracted from Cheng et al. (2020). This 55 year old female survived a COVID-19 infection.

pneumonia. Similarly to the outcome of the Chest Xray14 (Wang et al., 2017) dataset which enabled significant advances in medical imaging, tools can be developed to predict not only the type of pneumonia, but also its outcome. Eventually, our model could take inspiration from work by Rajpurkar et al. (2017), which could predict pneumonia, as well as Cohen et al. (2019), which deployed such models.

Tools could be built to triage cases in the absence of physical tests, particularly in the context of polymerase chain reaction (PCR) tests shortage (Satyanarayana, 2020; Kelly Geraldine Malone, 2020). These tools could predict patient outcomes such as survival, allowing a physician to plan ahead for specific patients and facilitate management. In extreme situations, where physicians could be faced with the extraordinary decision to choose which patient should be allocated healthcare resources (Yascha Mounk, 2020), such a tool could potentially serve as a measuring device.

We collected scan images found attached to this research paper

Different resolutions from (255, 249) up to (4280, 3520)

- **COVID-Xray-5k dataset**
  - a. with 184 samples containing clinical findings of COVID-19 from 216 patients

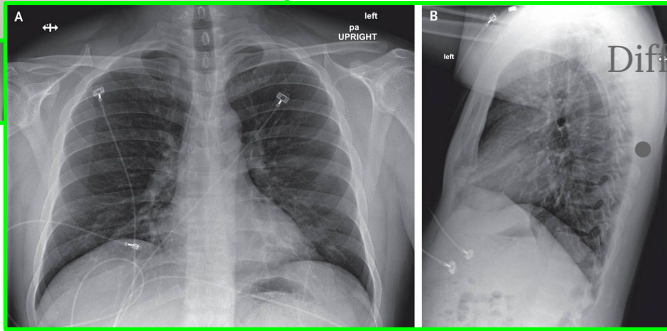
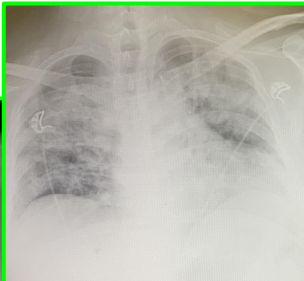
Table 1. Number of images per category in COVID-Xray-5k dataset.

Split	COVID-19	Non-COVID
Training Set	84 (420 after augmentation)	2000
Test Set	100	3000

<sup>1</sup>Mila, Quebec Artificial Intelligence Institute <sup>2</sup>University of Montreal <sup>3</sup>Department of Mathematics and Computer Science, Fontbonne University <sup>4</sup>Faculty of Medicine, University of Montreal. Correspondence to: Joseph Paul Cohen <joseph@josephpcohen.com>.



# Section 1 - The Chosen Dataset (2)



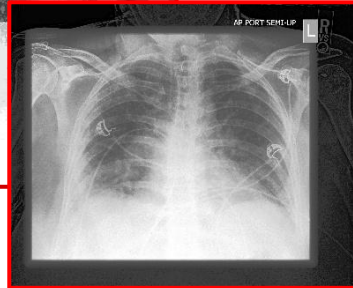
Different resolutions from (255, 249) up to (4280, 3520)

- **COVID-Xray-5k dataset**

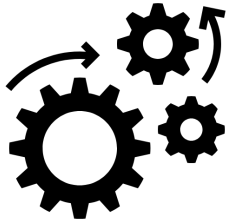
- a. with 84 CT images containing clinical findings of COVID-19 from 216 patients

**Table 1. Number of images per category in COVID-Xray-5k dataset.**

Split	COVID-19	Non-COVID
Training Set	84 (420 after augmentation)	2000
Test Set	100	3000



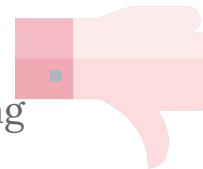
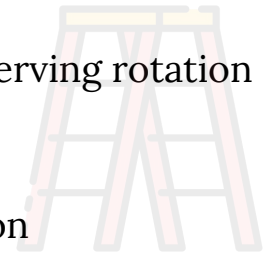
# Section 1 - The Chosen Dataset (2)



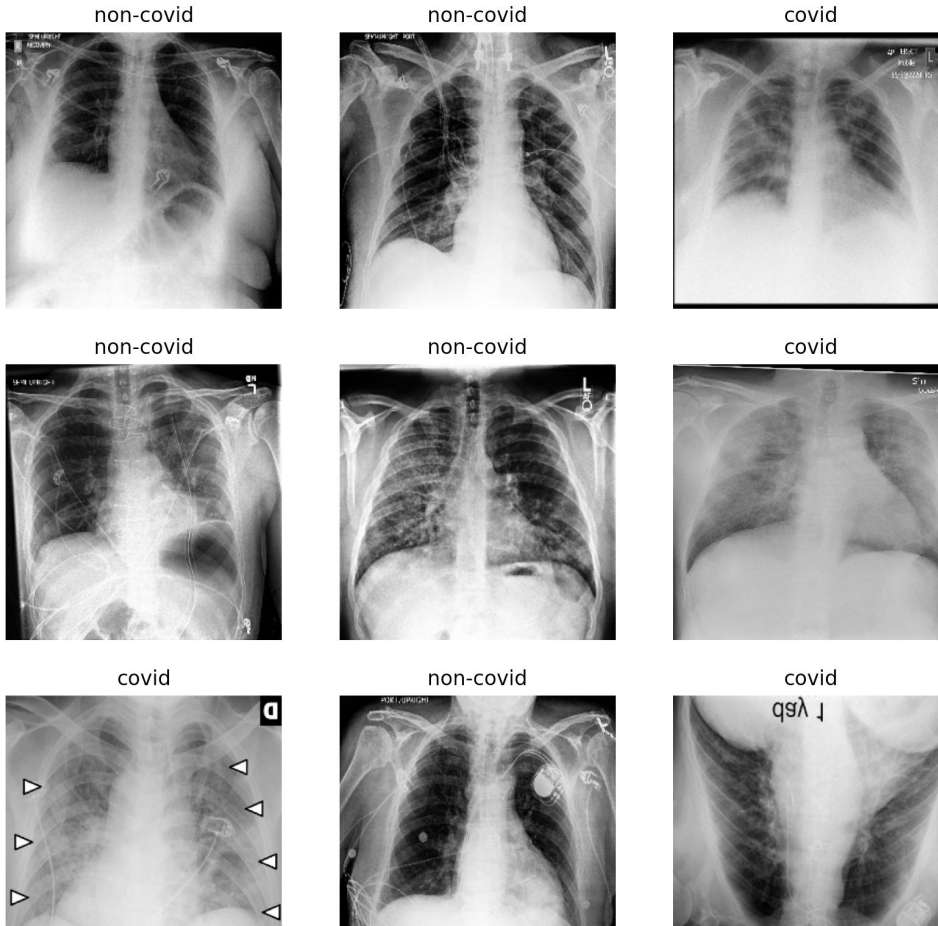
We obtained **more than 1000** images from the initial dataset after the data augmentation phase



1. **rotation** = (few degrees) size preserving rotation
  2. **flip\_left\_right** = mirroring
  3. **centered crop**
  4. **contrast jitter** = contrast alteration
- A. translation
  - B. gaussian noise
  - C. vertical mirroring
  - D. skew



# Section 1 - The Chosen Dataset ()



pros:

- community effort
- sharing is caring

cons:

- few images
- overfitting
- need doctors supervision

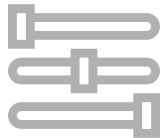
# How the presentation is organized



Section 1 - The Chosen Dataset & Background



**Section 2 - The PreTrained CNN Network - ResNet101**



Section 3 - Our custom CNN Network

# Section 2 - The PreTrained CNN Network - ResNet101 (1)

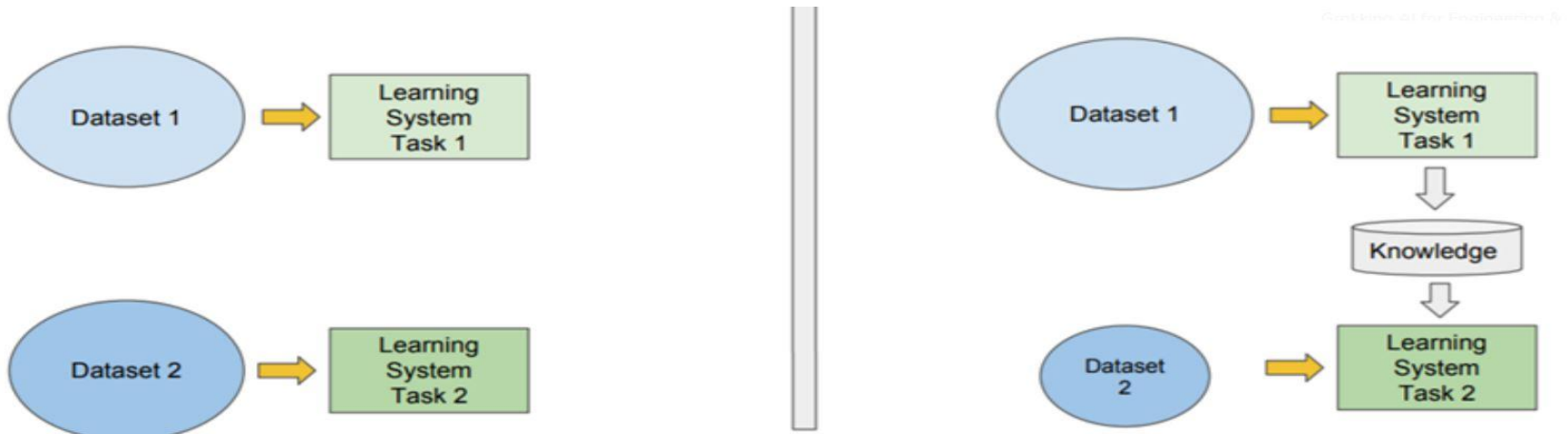
## Traditional Machine Learning

- Isolated
- Single Task Learning
- Knowledge not accumulated

vs

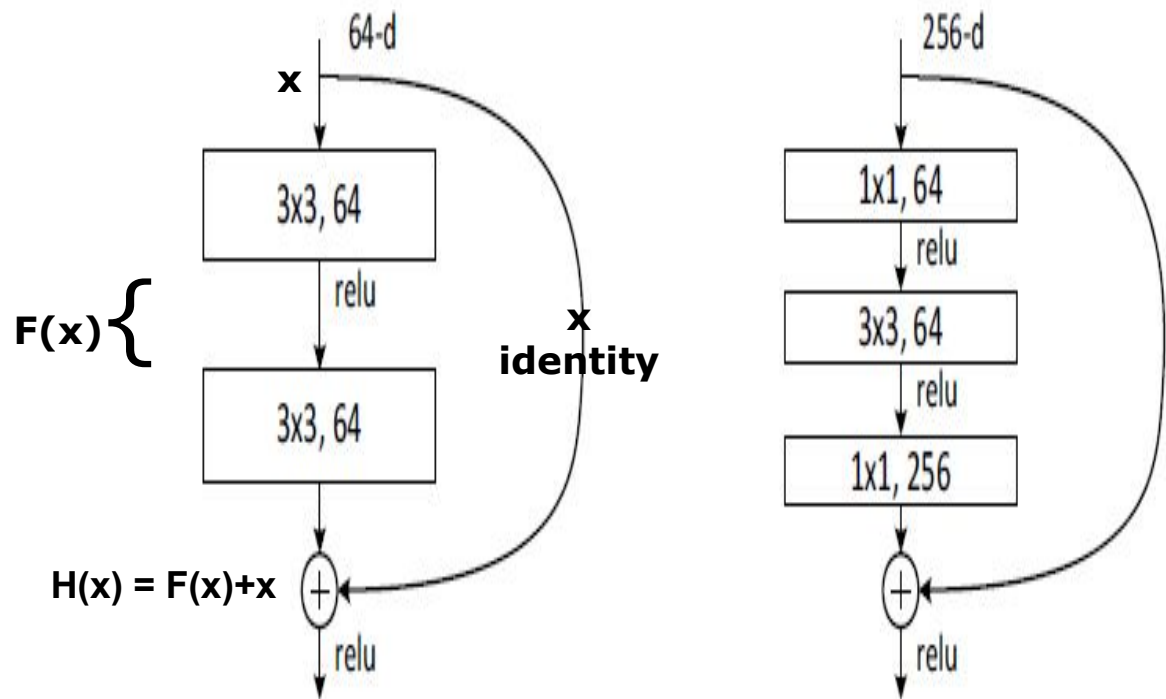
## Transfer Learning

- Learning relies on previously learned tasks
- Learning process faster
- More accurate
- Less train data



# Section 2 - The PreTrained CNN Network - ResNet101 (1)

- “Depth Revolution”
- Vanishing/exploding gradient
- **Residual Network:**
  1. Identity Mapping
  2. Projections
- **Bottleneck design**



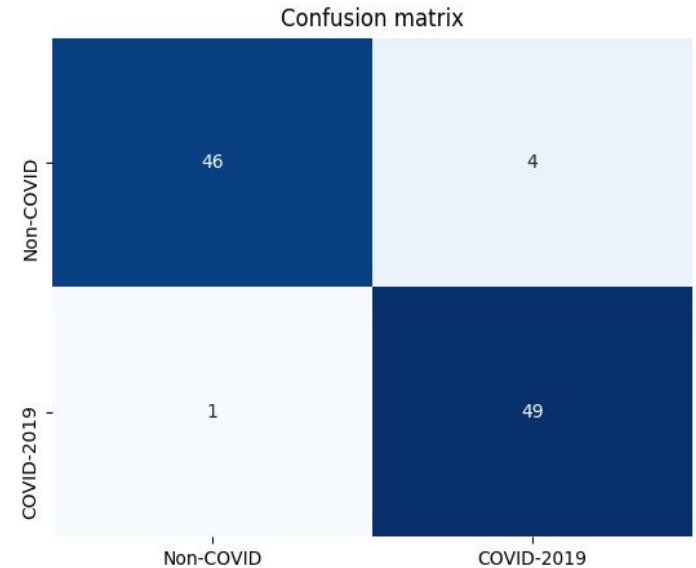
## Pagina 15



# Section 2 - The PreTrained CNN Network - ResNet101 (3)



	MIRRO RING	MIRRORING ROTATION	MIRRORING ROTATION CONTRAST JITTER	MIRRORING ROTATION CONTRAST JITTER CROP
<b>SENSITI VITY (TP/P)</b>	0.85	0.905	0.95	0.98
<b>SPECIFI CITY (TN/N)</b>	0.82	0.86	0.89	0.92
<b>ACCURA CY (TP+TN)/ (TOT)</b>	0.835	0.88	0.92	0.95



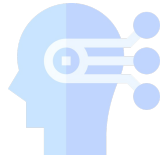
↓

sensitivity : 0.98  
specificity : 0.92  
accuracy : 0.95  
f1\_score : 0.9514

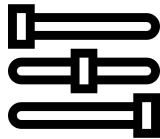
# How the presentation is organized



Section 1 - The Chosen Dataset & Background



Section 2 - The PreTrained CNN Network - ResNet101



**Section 3 - Our custom CNN Network**

# Section 3 - Our custom CNN Network (1)

## Major issues due to the small dataset:

- Overfitting
- Vanishing gradient
- Oscillations in both training and testing

## Solutions adopted:

- Batch normalization
- L2 weight decay
- Dropout

```
model = tf.keras.Sequential([
    data_augmentation,
    layers.experimental.preprocessing.Rescaling(1./255)

    layers.Conv2D(16, 5, padding='valid', activation='relu'),
    layers.MaxPooling2D( (2,2) ), #stride default
    layers.BatchNormalization(momentum=0.8),

    layers.Conv2D(32, 3, padding='same', activation='relu'),
    layers.MaxPooling2D( (2,2) ), #stride default
    layers.BatchNormalization(momentum=0.8),

    layers.Conv2D(64, 3, padding='same', activation='relu'),
    layers.MaxPooling2D( (2,2) ), #stride default
    layers.BatchNormalization(momentum=0.8),

    layers.Conv2D(128, 3, padding='same', activation='relu'),
    layers.MaxPooling2D( (2,2) ), #stride default
    layers.BatchNormalization(momentum=0.8),

    layers.Dropout(0.5),
    layers.Flatten(),

    layers.Dense(256, activation='relu') #First FC layers
    layers.Dense(2)                    #second FC. 2 is our #classes
])
```

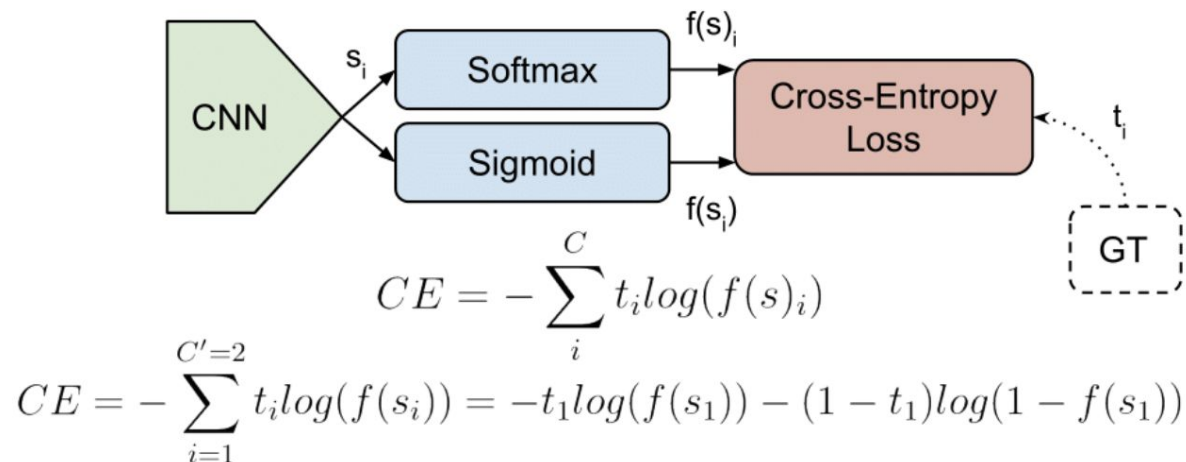
# Section 3 - Our custom CNN Network (2)

**Loss function** experiments:

1. BinaryCrossEntropy
2. CategoricalCrossentropy (Sparse)

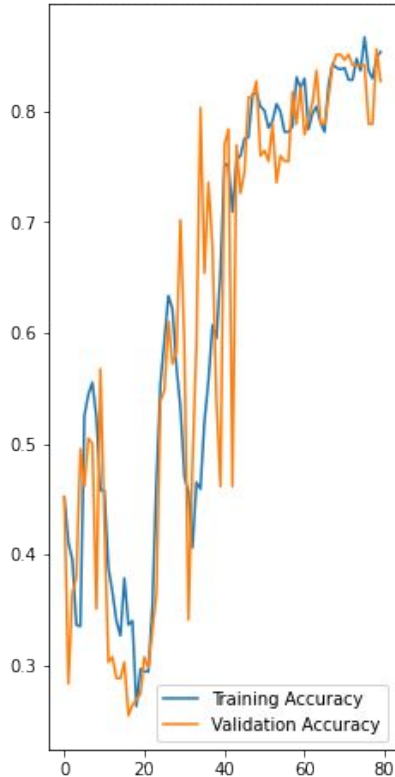
The first iterates over two classes (0,1) so it works well with a **sigmoid** before computation.

The second is meant for two or more classes, so take advantage of a **softmax** activation.

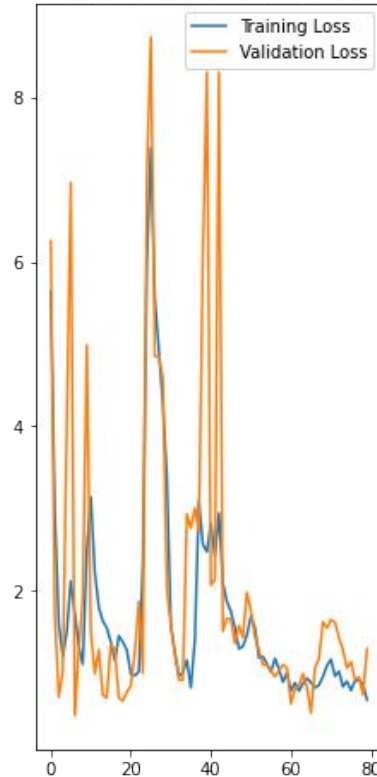


# Section 3 - Our custom CNN Network (3)

Training and Validation Accuracy

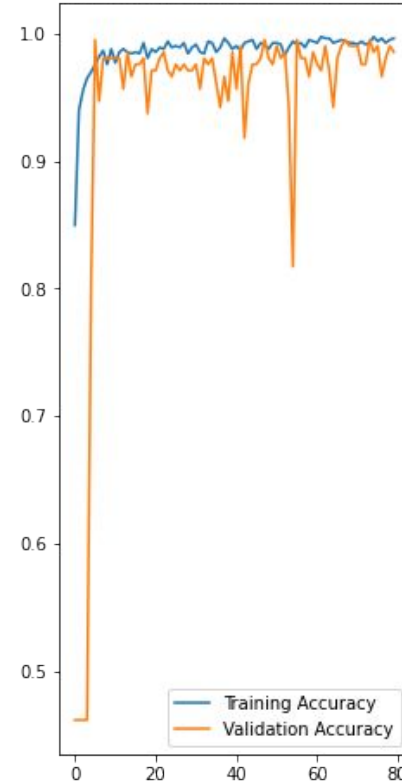


Training and Validation Loss

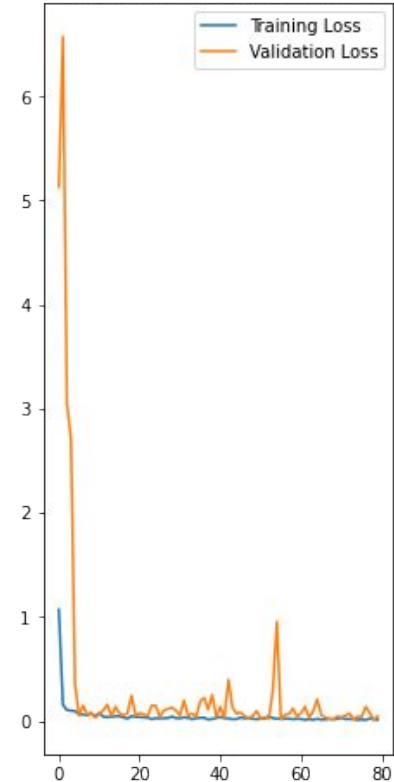


**Binary Cross Entropy** (best results after a very long hyperparams tuning)

Training and Validation Accuracy



Training and Validation Loss



**Sparse Cat. Cross Entropy** (one of the first obtained results: only 2 epochs to achieve an optimal val\_acc)

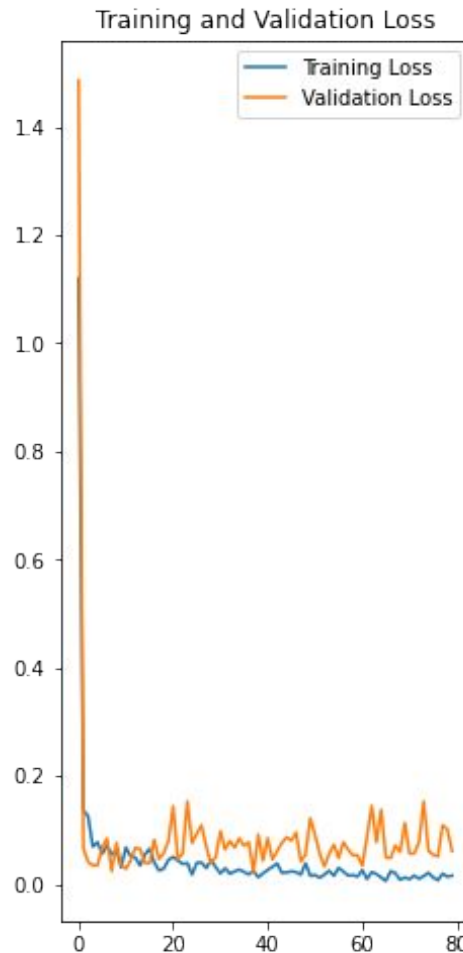
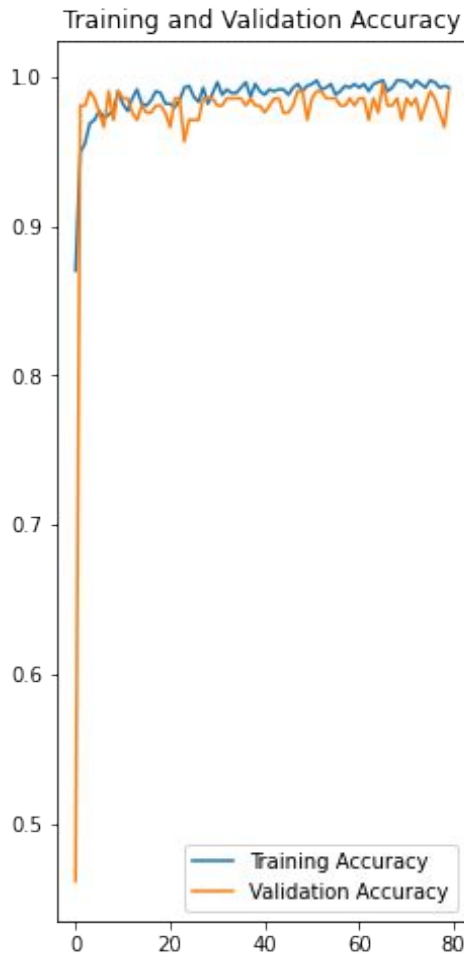
# Section 3 - Our custom CNN Network (4)

**Data augmentation** experiments:

	MIRRORING	MIRRORING ROTATION	MIRRORING ROTATION CONTRAST JITTER	MIRRORING ROTATION CONTRAST JITTER CROP
<b>SENSITIVITY (TP/P)</b>	0.71	0.75	0.78	0.86
<b>SPECIFICITY (TN/N)</b>	0.93	0.95	0.96	1.0
<b>ACCURACY (TP+TN)/(TOT)</b>	0.82	0.85	0.87	0.93

Note: some other techniques have been tried, but not relevant in terms of network accuracy

# Section 3 - Our custom CNN Network (5)



## Best Model

Conv2D: (16) (32) (64) (128)  
Dense\_1: (256)  
Dense\_2: (2)

(Not so deep, but very efficient)

**Batch size:** 64

**Epochs:** 80

**Dropout:** 0.5

**Batch norm. momentum:** 0.8

**Loss:** sparse categorical cross entropy

**Optimizer:** Adam with 0.9 LR decay rate

**Starting LR:**  $1e^{-2}$

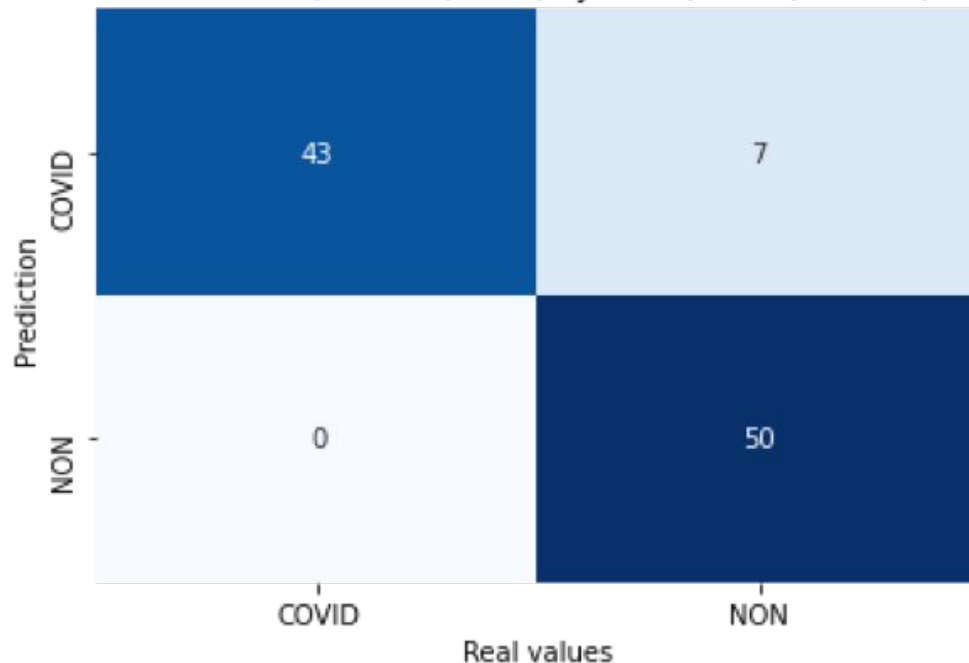
**Activation function:** relu



# Section 3 - Our custom CNN Network (6)

To have a practical response of it, a brief **inference** part is realized, with very good results

PERFORMANCE OF : /content/drive/My Drive/vision/dataset/model.h5



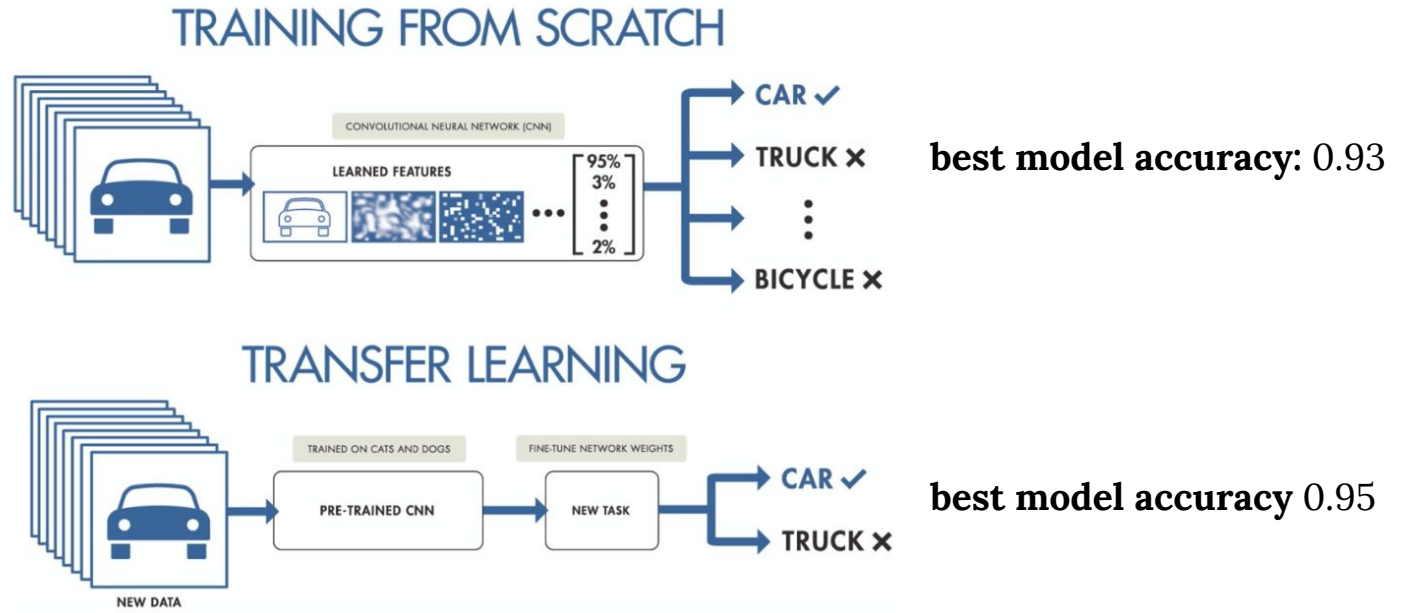
**Confusion Matrix**



sensitivity : 0.86  
specificity : 1.0  
accuracy : 0.93  
f1\_score : 0.9247

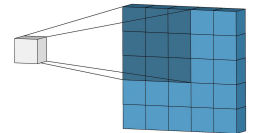
The blind testset is the same we used for the first part with the transfer learning, to make also a further comparison

# Final considerations

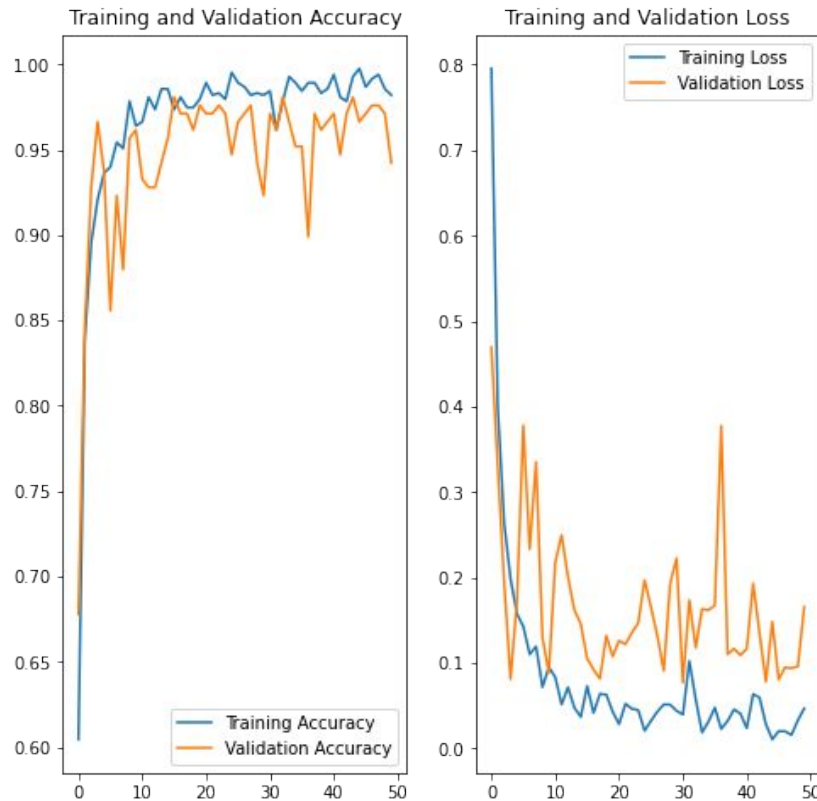


## Future Works:

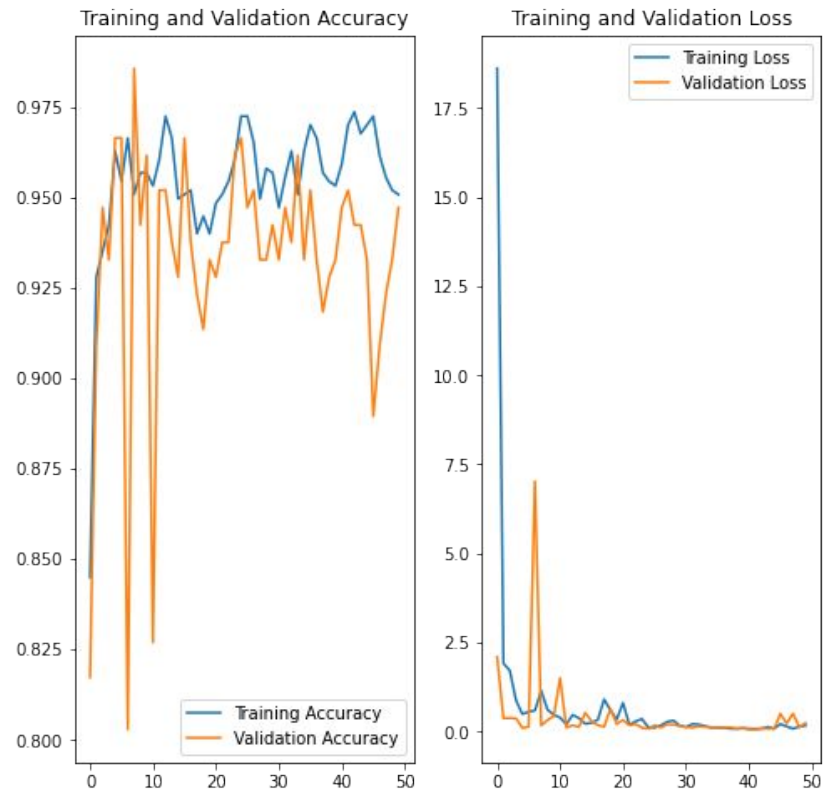
- 1) extend the dataset
- 2) multiclass classification (covid, pneumonia, pneumothorax ...)



# Appendix: additional exp. fail plots (1)

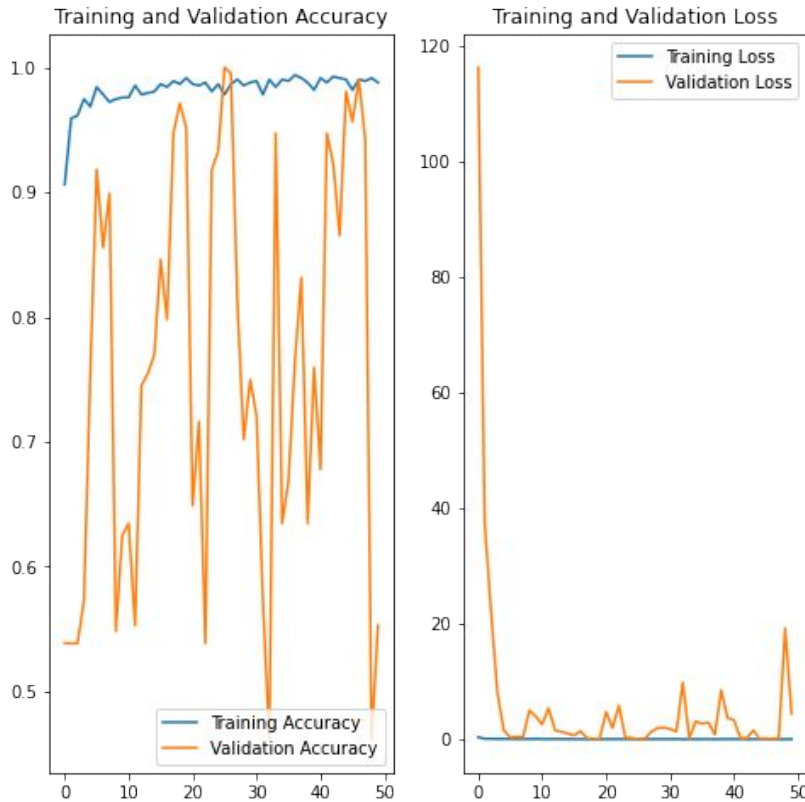


Conv2D: 16 32 64 128 256 512  
No batch normalization, no LR decay  
Very long training time  
Good accuracy but unstable: bad inference

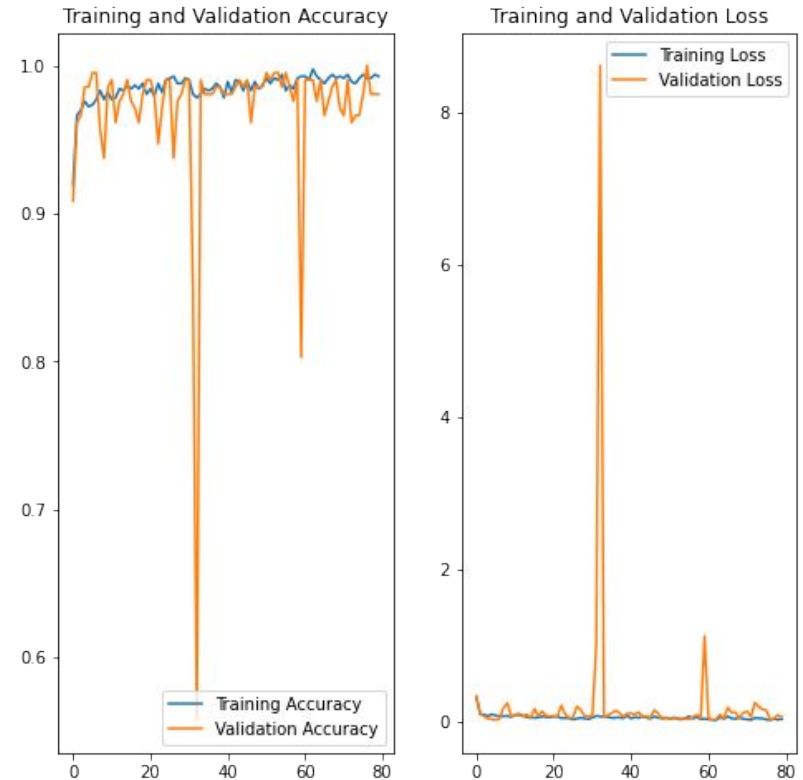


Conv2D: 16 32 64 128 256  
First introduction of the batch norm  
Very unstable training  
Val\_accuracy oscillating, bad inference

# Appendix: additional exp. fail plots (2)



Deep network (32 32 64 64 ... 512)  
No data augmentation  
Batch size small (8), big b.n. momentum (0.99)  
LR: 0.1 without decay (very unstable net)



Standard network (shown before)  
Batch norm before max pooling  
No LR decay: instability  
Inference not so bad

THANKS FOR  
YOUR  
ATTENTION

