

# Alternative Methods to Detect COVID-19

A Digitial Signal and Image Management Project

Davoli Sofia - 813479

Filosa Alberto - 815589





#### **PIPELINE**



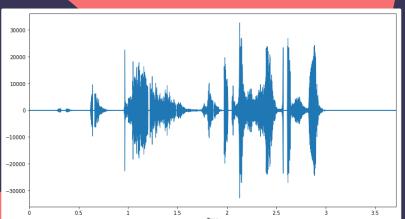


# 01 AUDIO

**Audio Recognition** 

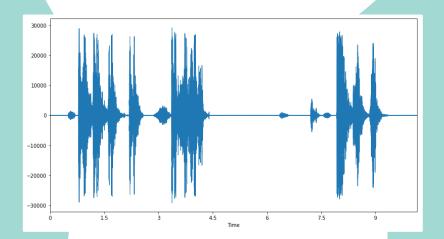






## **Not COVID**





#### **AUDIO AUGMENTATION**

Only COVID-19 audio



#### **ADD NOISE**

Simply add some random value into audio



#### **CHANGE SPEED**

Change speed of audio (randomly)

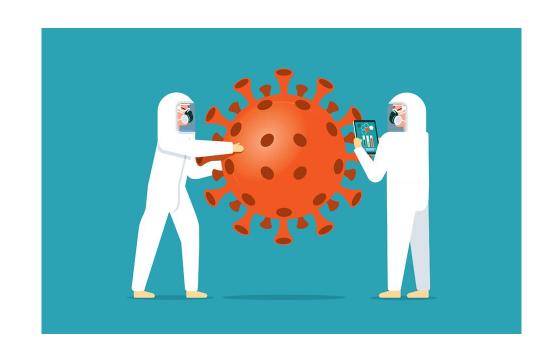


#### **PITCH SHIFTING**

Adjust the intonation of the voice (randomly)

#### **FEATURE EXTRACTION**

- Chroma STFT 1;
- Chroma STFT 6;
- Chroma STFT 12;
- Spectral Centroid;
- Spectral Bandwidth;
- Rolloff;
- Zero Crossing Rate;
- MFCC from 1 to 20.





#### **MODELS**



#### **SVM**

C: 10

Gamma: 0.0001



#### **LOGISTIC REGRESSION**

Solver: LibLinear Balanced Weights



#### **NEURAL NETWORK**

6 Layers

Activation Function: ReLu



#### **CNN**

2 Models



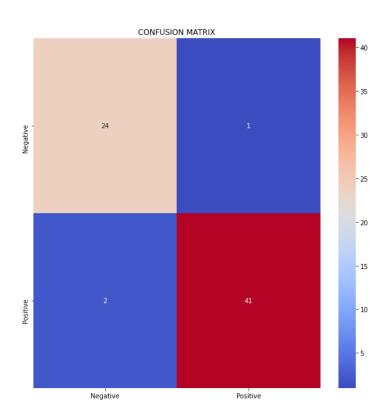
#### **BEST MODEL - LOGISTIC**

#### Feature Importance:

- $Coeff_{MFCC_3} = -0.57758$
- $Coeff_{MFCC_7} = -0.32325$
- $Coeff_{MFCC_8} = 0.36815$
- $Coeff_{MFCC_{10}} = 0.40355$
- $Coeff_{MFCC_{12}} = 0.48419$
- $Coeff_{MFCC_{14}} = 0.45921$



		Precision	Recall	F1-Score
	Negative	0.92	0.96	0.94
	Positive	0.98	0.95	0.96

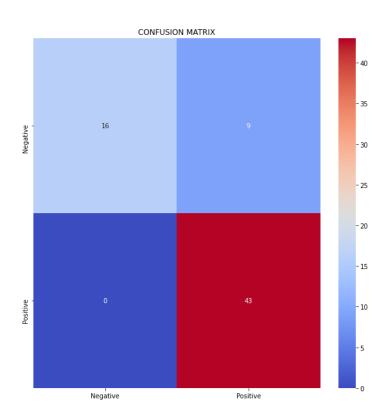


#### **BEST MODEL (NN)**

- Sequential()
- Input(27, 1)
- Convolutional 1D (512, 10)
- Convolutional 1D (256, 10)
- MaxPooling()
- Dense(10)
- Flatten()
- Dense(2) with Softmax



	Precision	Recall	F1-Score
Negative	0.96	0.96	0.96
Positive	0.98	0.97	0.98



#### LIVE TUTORIAL

You can record your cough to see if you are Positive to COVID-19.

**BUT** remember, the Accuracy is 97% and it is not a swab!

```
[ ] signal, rate = librosa.load("audio.wav")
 ipd.Audio(signal, rate = rate)
 /usr/local/lib/python3.6/dist-packages/librosa/core/audio.py:162: UserWarning: PySoundFile failed. Trying audioread instead.
  warnings.warn("PySoundFile failed. Trying audioread instead.")
   ▶ 0:00 / 0:02 ———
def kwow if you are posivite(audio):
   signal, rate = librosa.load(audio)
   #-- Create DataFrame
   new audio = pd.DataFrame()
   new audio["filename"] = 0
   new_audio.loc[0] = audio
   #-- Add Features
   new audio["chroma stft 1"] = np.mean(librosa.feature.chroma stft(y = signal, sr = rate)[0])
   new audio["chroma stft 6"] = np.mean(librosa.feature.chroma stft(y = signal, sr = rate)[5])
   new audio["chroma stft 12"] = np.mean(librosa.feature.chroma stft(y = signal, sr = rate)[11])
   new audio["spectral centroid"] = np.mean(librosa.feature.spectral centroid(y = signal, sr = rate))
  new audio["spectral bandwidth"] = np.mean(librosa.feature.spectral bandwidth(y = signal, sr = rate))
   new audio["rolloff"] = np.mean(librosa.feature.spectral rolloff(v = signal, sr = rate))
   new audio["zcr"] = np.mean(librosa.feature.zero crossing rate(y = signal))
   for i in range(0, 20):
     new audio["mfcc " + str(i)] = np.mean(librosa.feature.mfcc(y = signal, sr = rate)[i])
  y pred = model.predict(new audio[new audio.columns[1:]], verbose=0)
  y pred = np.argmax(y pred, axis = 1)
  if y pred == 1:
     print("Oh Man! You Are Positive to COVID-19! Stay Safe!")
     print("Lucky You! You are Negative to COVID-19. Wear your mask and Stay Safe!")
   return y pred
```

[ ] kwow if you are posivite("audio.wav")



02

# **IMAGE**

X-Ray Chest Recognition



# **COVID** Histograms

### **Not COVID**









Shear angle in counter-clockwise direction in degrees

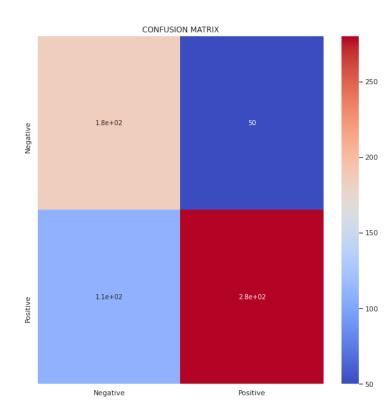
#### **MODEL:** ResNet - 50

#### Fine Tuning:

- GlobalAveragePooling2D()
- Dense() with 128 Neurons
- Dropout(0.2)
- Dense(1) with Sigmoid Activation



	Precision	Recall	F1-Score
Negative	0.63	0.79	0.70
Positive	0.85	0.72	0.78



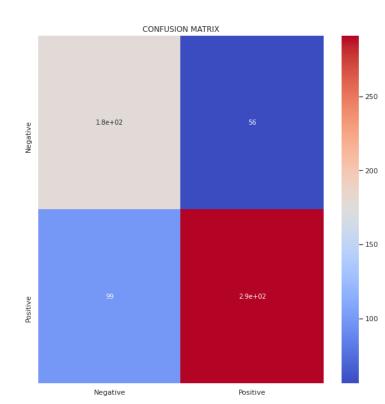
#### **MODEL:** EfficientNet – B7

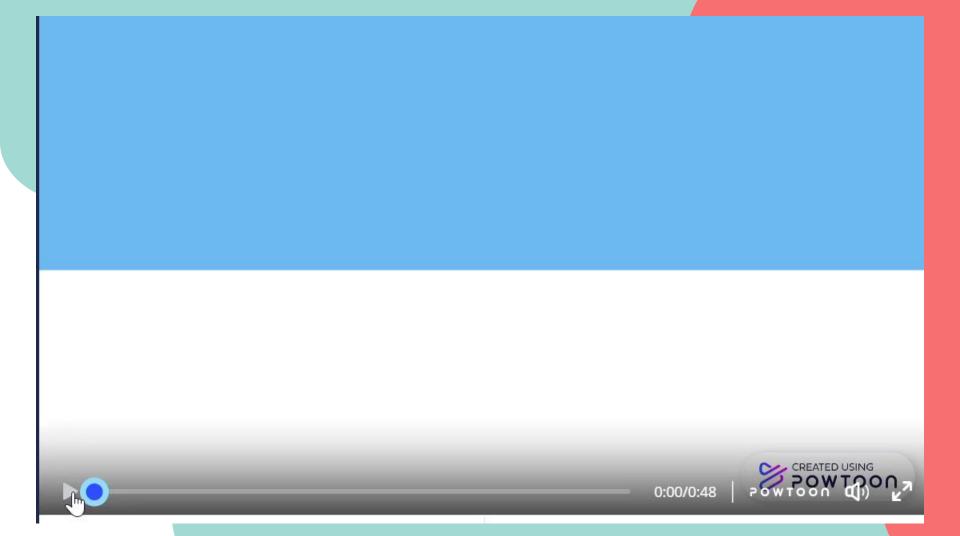
#### Fine Tuning:

- GlobalAveragePooling2D()
- Dense() with 128 Neurons
- Dropout(0.2)
- Dense(1) with Sigmoid Activation



	Precision	Recall	F1-Score
Negative	0.64	0.76	0.70
Positive	0.84	0.75	0.79





# **THANKS**

Do you have any question?

