

**Given a A FULLY AUTOMATED DEEP LEARNING-BASED NETWORK FOR DETECTING COVID-19 FROM A NEW AND LARGE LUNG CT SCAN DATASET paper**

**understand the paper and run the code**

**What is 1\*1 convolution in FPN?**

**How three features are extracted from the top layer of FPN?**

**Run the code**

**Calculate the dimension differences occurring while using FPN.**

- 1\*1 convolution is for changing the dimension of the depth.
  - (32,32,512) 1\*1 conv with 1 filter is (32,32,1)
  - If we use 512 filters then our dimensions are (32,32,512).
- There are five features
  - conv(1\*1) of C5
  - conv(1\*1) of C4
  - conv(1\*1) of C3
  - conv(3\*3, stride=2) of C5
  - activation on P6 and conv(3\*3, stride=2) of P6
- RESNET50V2:
  - C3.shape (None, 32, 32, 512)
  - C4.shape (None, 16, 16, 1024)
  - C5.shape (None, 16, 16, 2048)
- FPN:
  - NET 5 features
  - Feature 1
    - P5 = conv(1\*1) of C5 ->reduced to 256 featuremaps
    - P5.shape(conv1\*1) (None, 16, 16, 256)
    - P5 Upsampled.shape  
KerasTensor(type\_spec=TensorSpec(shape=(None, None, None, 256), dtype=tf.float32, name=None),  
name='P5\_upsampled/resize/ResizeNearestNeighbor:0',  
description="created by layer 'P5\_upsampled'")
    - P5.shape(conv 3\*3) (None, 16, 16, 256)
  - Feature 2
    - P4=conv(1\*1) of C4 ->reduced to 256 feature maps
    - P4.shape(conv1\*1) (None, 16, 16, 256)
    - P4.shape(concatenate P5\_upsampled,P4) (None, 16, 16, 512)
    - P4\_upsampled.shape (None, None, None, 512)
    - P4.shape(conv 3\*3) (None, 14, 14, 256)
  - Feature 3
    - P3=conv(1\*1) of C3 ->reduced to 256 feature maps
    - P3.shape(conv1\*1) (None, 32, 32, 256)
    - P3.shape(concatenate P4\_upsampled,P3) (None, 32, 32, 768)
    - P3.shape(conv 3\*3) (None, 30, 30, 256)

- Feature 4
  - **P6=conv(3\*3,stride=2) of C5**
  - **P6.shape(conv 3\*3,stride=2) (None, 8, 8, 256)**
- Feature 5
  - **P7=activation on P6 and conv(3\*3,stride=2) of P6**
  - **P7.shape(P6-relu,conv 3\*3,stride=2) (None, 4, 4, 256)**