

Lab 2: pipeline presentation

Winter School 25/01/2022







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Data from Medical Imaging Databank in Valencian Region Medical Image Bank (BIMCV)



BIMCV-COVID19+ (posi)

Images and medical data from COVID-19 patients

BIMCV-COVID19- (neg)

Images and medical data from NO COVID-19 patients







Posi dataset

- 4.854 subjects
- 13.368 sessions
- 23.527 images

Neg dataset

- 5.201 subjects
- 7.565 sessions
- 18.573 images

Note

There are 810 common subjects between the posi and neg datasets. They have different subject ID in each dataset.







Report: opacidades de aspecto intersticioal veolar parcheadas y bilaterales que predominan en ambos lobulos inferiores sospechosas de infeccion por COVID-19 . senos costofrenicos libres .

Labels: COVID 19, alveolar pattern, interstitial pattern, pneumonia Locations: costophrenic angle, lobar, bilateral, lower lobe

| DICOM Fields | Date | Test | Result |
|----------------------------|------------|------|----------|
| Study Date 20200317 | 17.03.2020 | PCR | NEGATIVE |
| Patient's Sex M | 18.03.2020 | PCR | NEGATIVE |
| Patient's Birth Date 1986 | 19.03.2020 | IGG | POSITIVE |
| Modality CR | 19.03.2020 | IGM | POSITIVE |
| Manufacturer GE Healthcare | 20.03.2020 | PCR | POSITIVE |
| ••• | | | |

Image: PNG or NIfTI

DICOM metadata

Medical Report

Labels

Locations of radiological findings

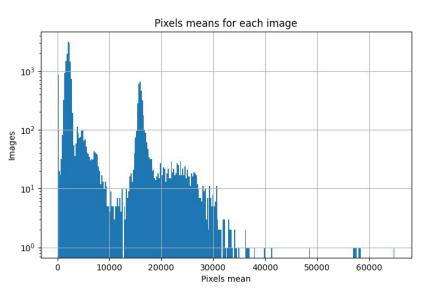
COVID tests (only for posi dataset)

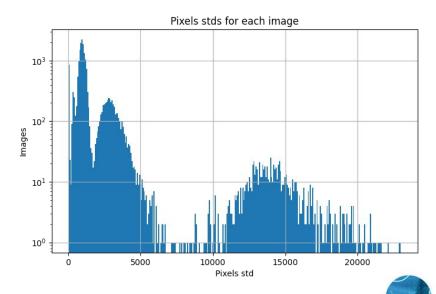
Segmentation (23 images in posi dataset)

[1] From BIMCV COVID-19+ preprint









DEEPHEALTH





| Shapes | Number of images |
|-------------|------------------|
| 2336 x 2836 | 1537 |
| 3480 x 4248 | 1241 |
| 3488 x 4256 | 1068 |
| 3480 x 4240 | 955 |
| 2544 x 3056 | 929 |

- 4082 different shapes
 - Most of them with only one image
- High resolution images
 - Between 2k and 3k pixels of height and width







| Label | Count |
|-------------------|-------|
| normal | 3412 |
| pneumonia | 3214 |
| increased density | 2996 |
| COVID 19 | 2820 |

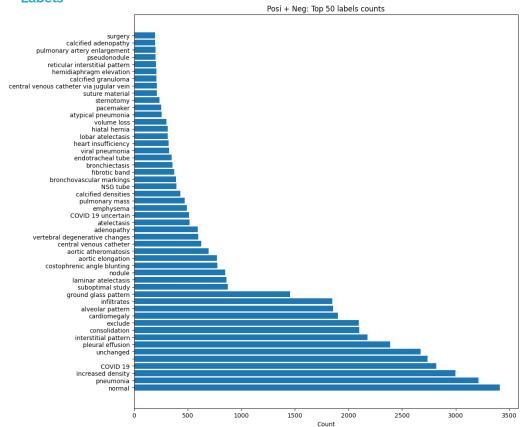
- 190 different labels
- Each sample has a list of labels
- There are samples in the posi dataset without the "COVID 19" label
- There are samples in the neg dataset with the "COVID 19" label



Dataset Description



Labels







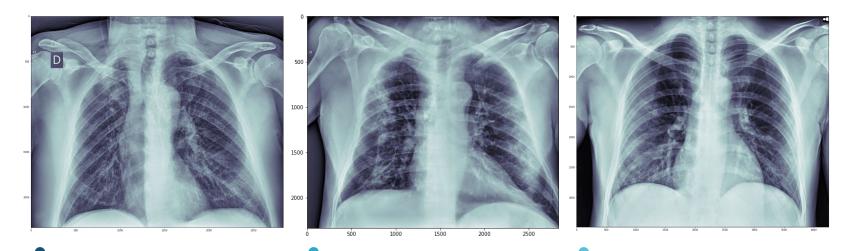


Pneumonia

Dataset Description

Labels examples





COVID 19

Infiltrates

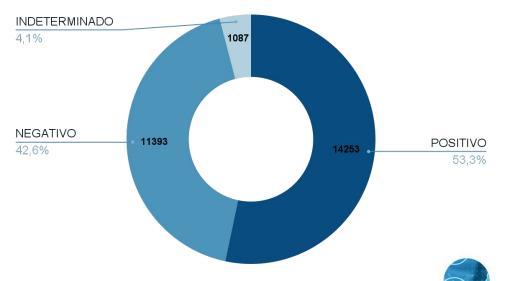




Tests types

| Test | Count |
|------|-------|
| PCR | 16442 |
| IGG | 4273 |
| ACT | 3294 |
| IGM | 2724 |

Tests results









DEEPHEALTH

01

Relabel using COVID tests (not used)

Relabel the samples that don't have the "COVID 19" label and they have a positive COVID test

02

Filter by classification labels

Select the samples that have at least one of the labels that are going to be used for classification, e.g. "COVID 19" vs "normal"

03

Filter by view type

Select the samples that are Anterior-Posterior (AP) or Posterior-Anterior (PA)

04

Filter using manually cleaned data

We manually labeled the AP and PA samples from the posi dataset to remove the ones that are not correct





DEEPHEALTH

01

Invert the "MONOCHROME1" images

The "MONOCHROME1" images have the colors inverted, the background is white and the bones are black

02

Images normalization

Contrast Limited Adaptive Histogram Equalization (CLAHE)

03

Convert to RGB (not used)

Using the jet colormap and dividing the grayscale in ranges to apply the colors

04

Extract lungs ROI

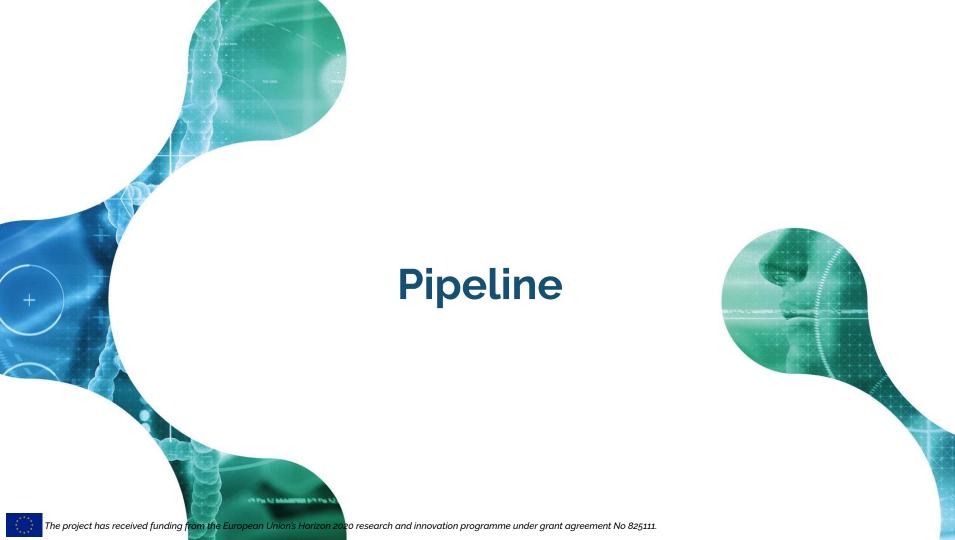
Using manually labeled masks of the lungs. We trained a U-Net to segment the lungs to perform the ROI extraction





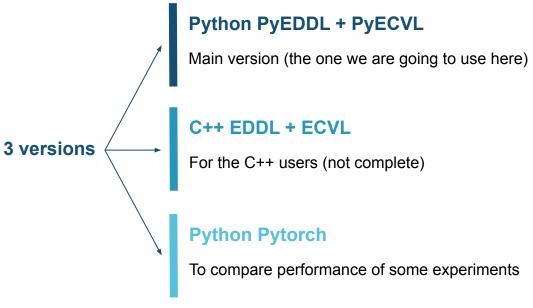


- The original dataset doesn't have the training splits defined
- We created a 80-10-10 (train-val-test) split
- The splits are defined at patient level: to avoid having samples from one patient in different splits















Models

Pretrained VGG and ResNet

Data shape

256x256 and 512x512

Optimizers

Adam and SGD LR: From 1e⁻⁴ to 1e⁻⁷

Regularization

L2 regularization. From 1e⁻² to 1e⁻⁶

Data augmentation

Mirror, Rotation, Brightness and GammaContrast

Losses

2 classes: CE and BCE

4 classes (multilabel): BCE and MSE







| Exp. | Task | Model | Data shape | Optimizer | Learning rate | Regularization | DA |
|------|---|-------------------------|---------------|-----------|------------------|-----------------------|-----|
| 0 | normal vs COVID (With BCE loss) | Pretrained ResNet101 | 256x256 | Adam | 1e ⁻⁵ | L2(1e ⁻⁵) | 1.1 |
| 1 | normal vs COVID vs pneumonia vs infiltrates (With BCE loss) | Pretrained ResNet101 | 512x512 | Adam | 1e ⁻⁵ | L2(1e ⁻⁵) | 1.1 |





| Test loss (BCE) | Test accuracy (%) | Test balanced accuracy (%) |
|-----------------|-------------------|----------------------------|
| 0,404 | 88,57 | 83,57 |





| Metric | normal | COVID | pneumonia | infiltrates |
|-----------|--------|-------|-----------|-------------|
| Accuracy | 0,843 | 0,725 | 0,593 | 0,627 |
| Precision | 0,423 | 0,821 | 0,471 | 0,474 |
| Recall | 0,393 | 0,311 | 0,3 | 0,5 |
| F1-score | 0,407 | 0,451 | 0,366 | 0,486 |







[1] Dataset preprint: https://arxiv.org/abs/2006.01174

[2] BIMCV: https://bimcv.cipf.es/bimcv-projects/bimcv-covid19/

[3] Pipeline: https://github.com/deephealthproject/UC15_pipeline







4 main concepts



YOU CAN INCLUDE IMAGES IN THE CIRCLES











Our phillosophy



MORE QUALITY LESS QUANTITY

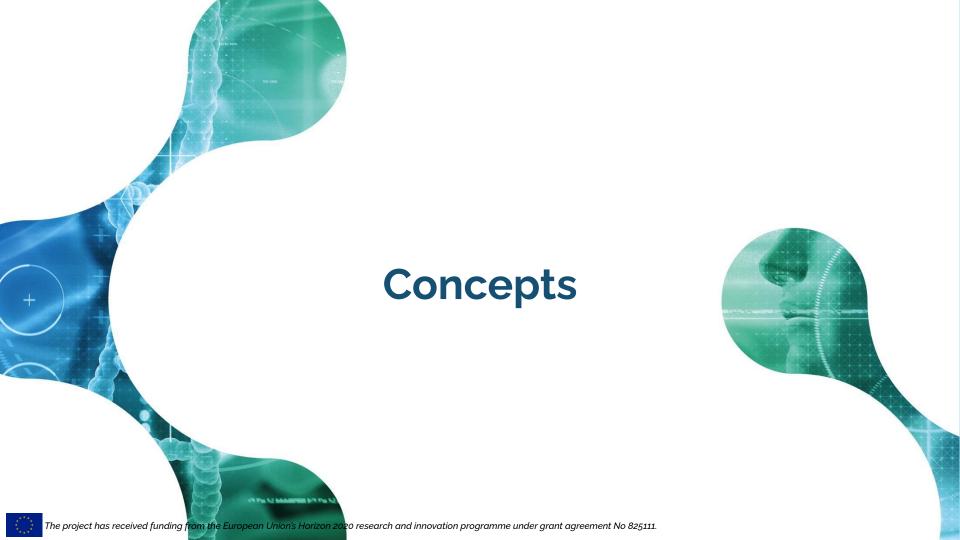
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Double images



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Gallery image feature 02

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Image with info



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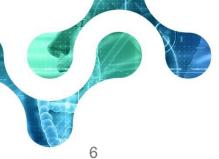
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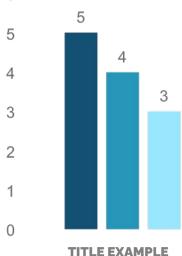




Bar charts



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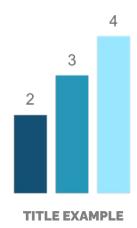
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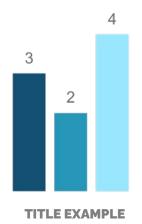
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Sharp line chart



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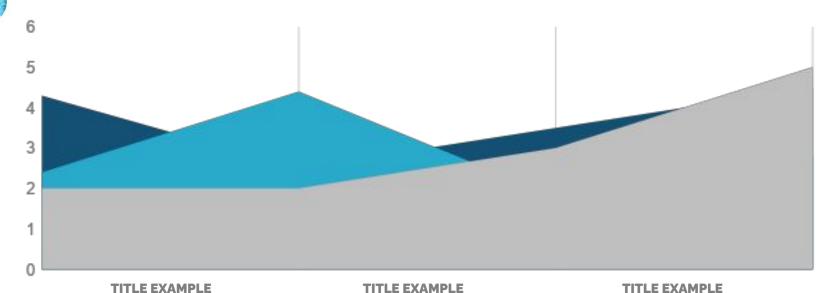




Area chart



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Curve line chart

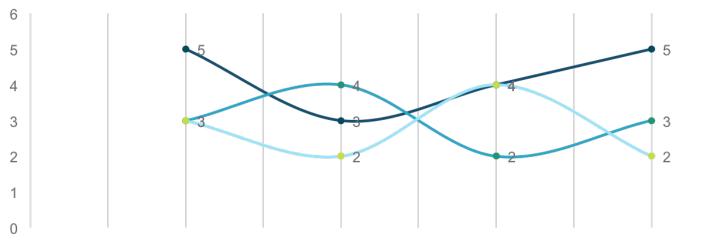


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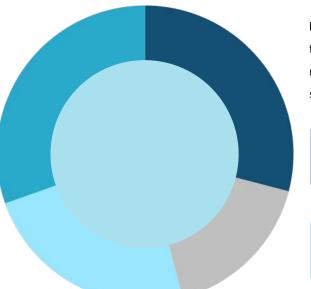
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Pie chart

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Timeline one

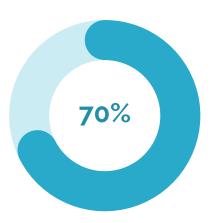


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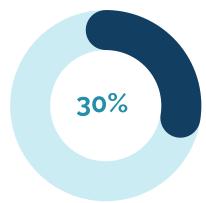
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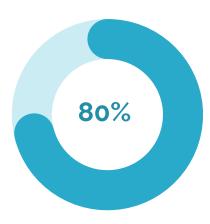
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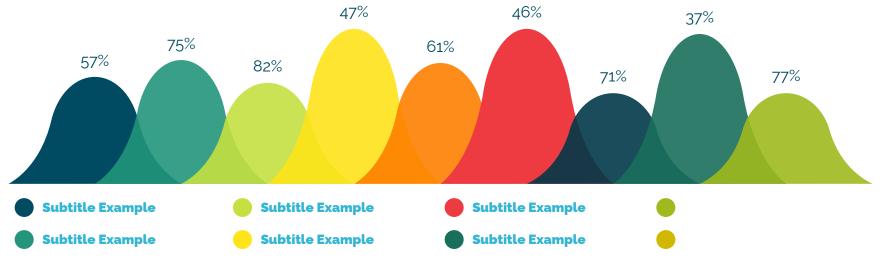


























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