## Labwork on **image classification**

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Solutions will be given on this online notebook.

## Chest CT-Scan images Dataset

- (a) Download the data here.
- (b) Install and load the 'OpenImageR' package.
- (c) Load and plot any image with the 'readImage' and 'imageShow' functions.
- (d) Compute the dimensions of the image, convert it to a black and white image with the 'rgb\_2gray' function and recompute its dimensions.
- (e) Compute the  $\boldsymbol{HOG}$  descriptors of the image with the 'HOG' function.
- (f) Write an R script that scraps and loads all the downloaded images, computes their HOG descriptors and their labels (adenocarcinoma, large.cell.carcinoma, squamous.cell.carcinom or normal) and saves them into a descriptors matrix and a labels vector.
- (g) In the *scraping* part of script, add a test that avoids duplicates.
- (h) Evaluate the predictive power of the processed dataset with tree and forest (and other) classifiers.
- (i) Try to increase the accuracy of your best models.

https://www.kaggle.com/mohamedhanyyy/chest-ctscan-images