

## **README NORMAL LIGHT**

**First:** Introduce the three scripts in the folder where all the images taken with normal light are placed.

**Whats in the folder:** There are three scripts:

- `Area_fibrosis.m` : Script of a function that measures the fibrotic area based on a color filtering of pink-purple.
- `Area_tissue.m` : Script in function form that measures total tissue area based on a color filtering that delete white spaces and fill gaps corresponding to tissue lost in the processing.
- `Area_measure_normal_light.m` : Script that reads all the images in .tif format of the folder and measures the fibrosis area (running `Area_fibrosis.m`), the tissue area (running `Area_tissue.m`) and the total area of the image (based on each one dimensions). It saves the results in a .csv file on the same folder. It's recommended to run ">> *Results*" on the Command Window to visualize it in MATLAB.

**Implementation:** Set the MATLAB directory to the folder where the images and scripts are. RUN ONLY '`Area_measure_normal_light.m`' :

Open '`Area_measure_normal_light.m`' on MATLAB and press Run. The results will be automatically saved on a .csv file named '`Area_measure_normal_light.csv`'

**Changes:** You can adapt these scripts to your studies making all changes needed like changing the name of the csv file or even setting new filtering functions by Color Threshold.