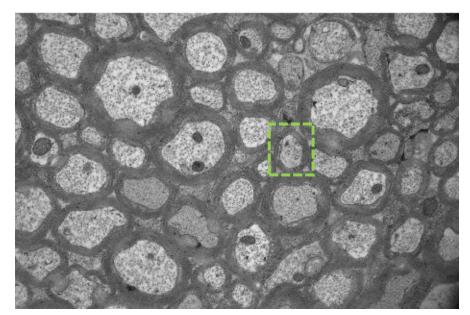
# Project n°8

Segmentation of myelin in electron microscopy images

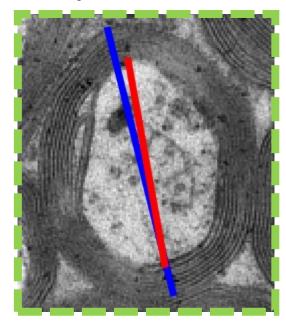
### Context

- Partnership between Lisite (Florence Rossant) and Lariboisière hospital (Anne Joutel) on axon and myelin analysis on Electron Microscopy (EM) mice images.
- Myelin: membran, composed of protid and lipid layers, surrounding axons, aiming at isolating and protecting them, and speeding up the transmission of nerve impulses.
- Clinical stake: extract biological biomarkers such as the g-ratio, that is the ratio between the diameter of the axon and the diameter of its associated myelin sheat

#### → Need in an individual segmentation of all myelin sheats



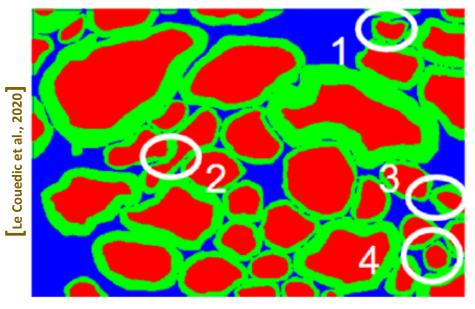
EM image



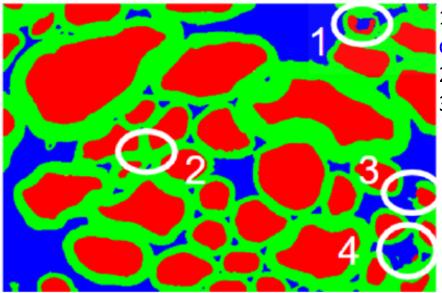
Zoomed EM image

## Preliminary results

- [Le Couedic et al., 2020]: Deep Learning based segmentation of EM images into myelin, axon and background using a U-Net-derived architecture ([Ronneberger et al., 2015])
- Use of 23 mice EM images, associated with their 3-class manual segmentations:
  - ➤ 18 for training
  - > 3 for validation
  - ≥ 2 for test
- Promising results, but room for improvements!



Ground truth image with highlighted errors



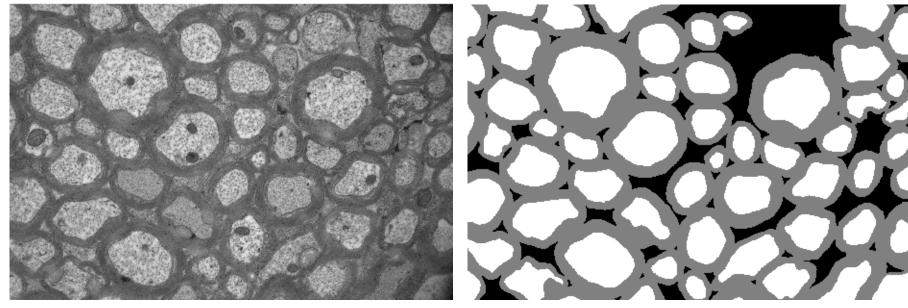
Model segmented image with highlighted errors

- Good segmentation but wrong classification
- 2. & 4. Wrong axon segmentation
- 3. & 4. Wrong myelin segmentation

→ The addition of topological information could improve the segmentation!

# Topological hypotheses to investigate

- The hidden topological constraints:
  - ➤ Manually segmented axons cannot touch background areas
  - > Each manually segmented axon is surrounded by myelin
    - → Supposing we get a pretty accurate binary segmentation of the myelin, each remaining region (connected component) can be further classified into axon or background using Machine Learning (ML)
    - → We hope that focusing on myelin for the segmentation helps 'closing' its contours (error 3)
    - → The ML part can also correct the misclassification of connected components (error 1)



Associated manual segmentation

### Expected outputs

- The goal of the project is to deliver a robust a two-phase algorithm
  - Phase 1: A Deep learning-based binary segmentation of the myelin
  - ➤ Phase 2: A Machine Learning-based classification into background and axon of the remaining regions (connected components of the complement image of the binary segmentation of the myelin)
- Provided data and tools:
  - > EM mice images and their associated manual segmentation into axon, background and myelin
  - > A jupyter notebook of the 3-class segmentation of EM images (Monai library)
- The notebook should be modified to:
  - > Use the binary segmentation of the myelin
  - > Implement topological losses (almost [Hu et al., 2019], if time [Kervadec et al., 2019], or other articles you find interesting)
  - > Include quantitative results during validation and test (Dice's similarity coefficient)
- Expected work:
  - > Read and present the works of [Hu et al., 2019]
  - > Present your work on the project during a final presentation
  - ➤ Give back your modified code

### Some references

#### Articles

- [1] Hu, Xiaoling, et al. "Topology-preserving deep image segmentation." arXiv preprint arXiv:1906.05404 (2019).
- [2] Kervadec, Hoel, et al. "Boundary loss for highly unbalanced segmentation." *International conference on medical imaging with deep learning*. PMLR, 2019.
- [3] Le Couedic, Thomas, et al. "Deep-learning based segmentation of challenging myelin sheaths." 2020 Tenth International Conference on Image Processing Theory, Tools and Applications (IPTA). IEEE, 2020.
- [4] Ronneberger, O., Fischer, P., & Brox, T. (2015, October). U-net: Convolutional networks for biomedical image segmentation. In *International Conference on Medical image computing and computer-assisted intervention* (pp. 234-241). Springer, Cham.

#### Sites web

Pytorch implementation of [1]: https://github.com/HuXiaoling/TopoLoss/blob/master/topoloss\_pytorch.py

Monai tutorials: https://monai.io/start.html#MONAITutorials