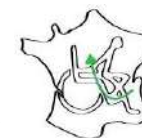
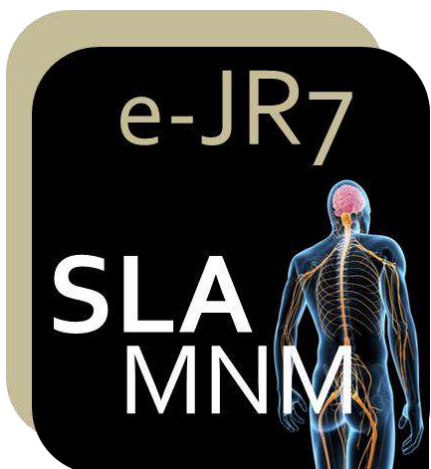


Propagation and toxicity of pathological determinants in Amyotrophic Lateral Sclerosis

7^{èmes} Journées de la Recherche sur la SLA et les Maladies du Neurone Moteur
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INTRODUCTION

- Pathological determinants are present in ALS motor neurons and also in glial cells

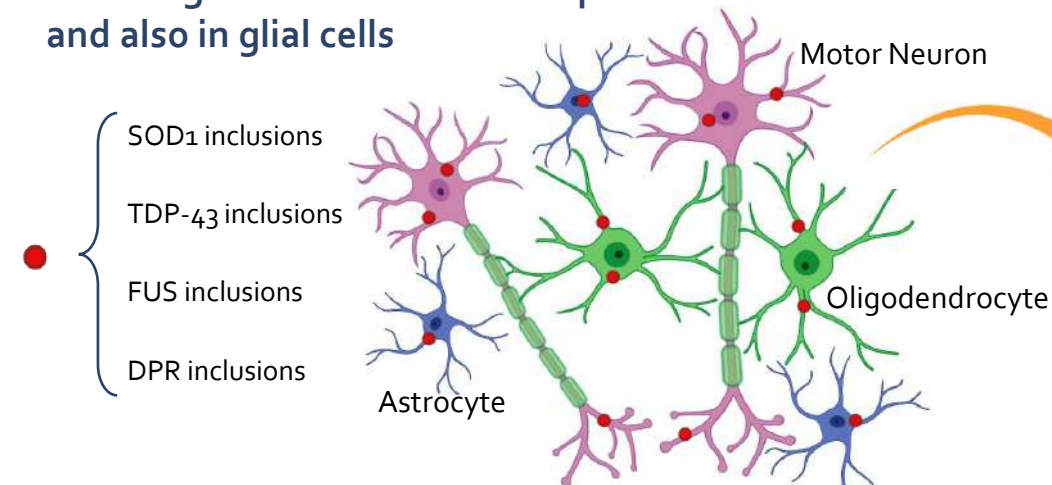


Figure 1. Scheme of motor neurons and glial cells illustrating accumulations of pathological determinants.

- The USP19-dependent MAPS pathway

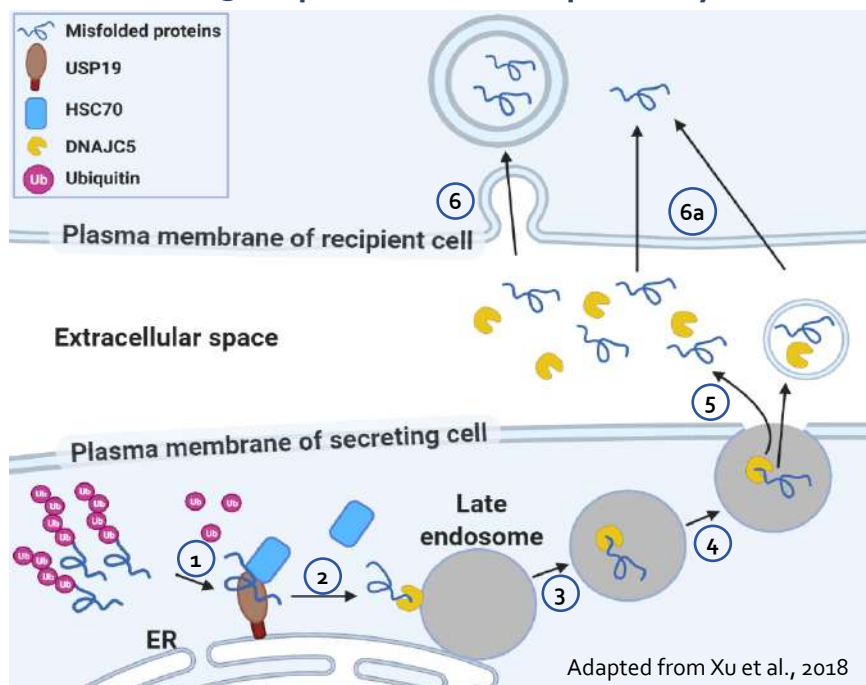


Figure 3. The MAPS pathway

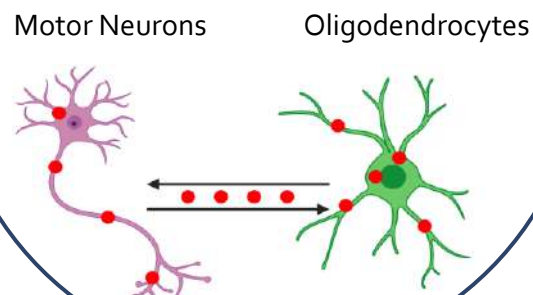
1. A misfolded protein (MP) can be recruited by the deubiquitinase USP19 localized on the ER.

2,3. The MP can then be transferred to the lumen of a late endosome (LE) via a translocation of the DNAJC5-MP complex
4,5. LE moves towards the cell membrane, fusion occurs and the MP can be released in the extracellular space.

6,6a. The MP may enter a recipient cell either by endocytosis or translocation

Adapted from Xu et al., 2018

HYPOTHESIS :
Could the pathological determinants of ALS spread from cell to cell and contribute to the progression of the disease ?



- ALS onset is anatomically localized and spread to the whole body

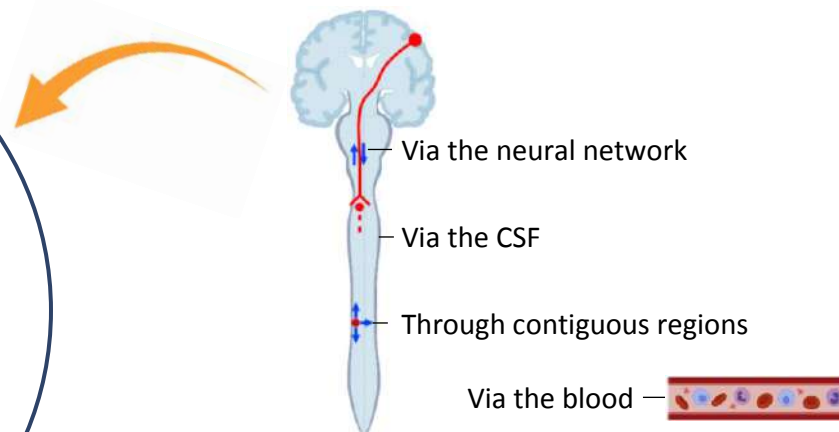


Figure 2. Scheme of the CNS and the different hypothesis of ALS pathology propagation

- USP19: Isoforms and roles

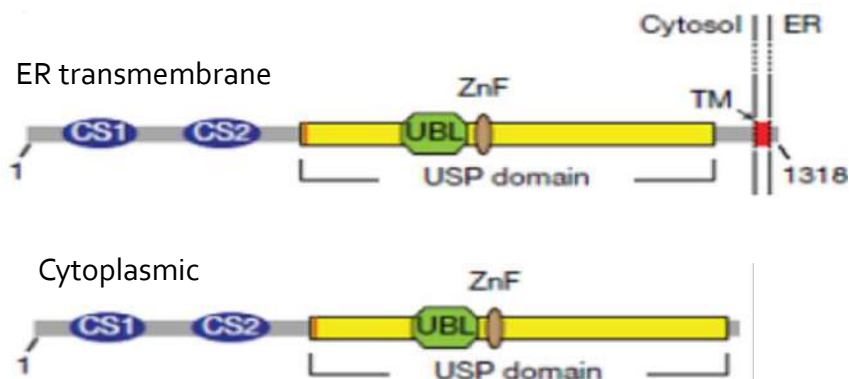


Figure 4. Scheme of USP19 protein.

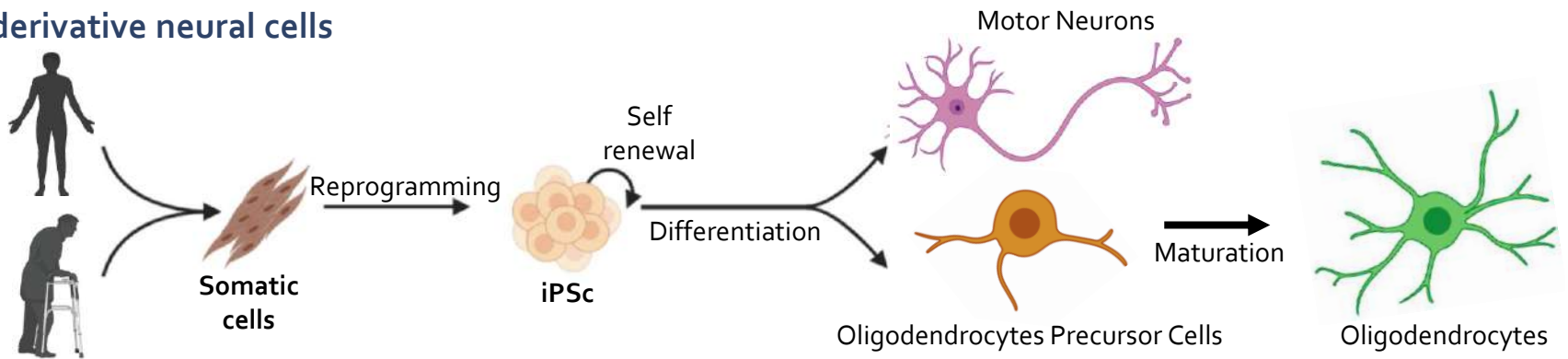
- Hydrolysis the bonds forms by ubiquitin
- Removing of poly ubiquitin chains from misfolded proteins
- 2 major isoforms: cytoplasmic and ER transmembrane

Lee et al., 2016

MODELS ● **iPSc generation and their derivative neural cells**

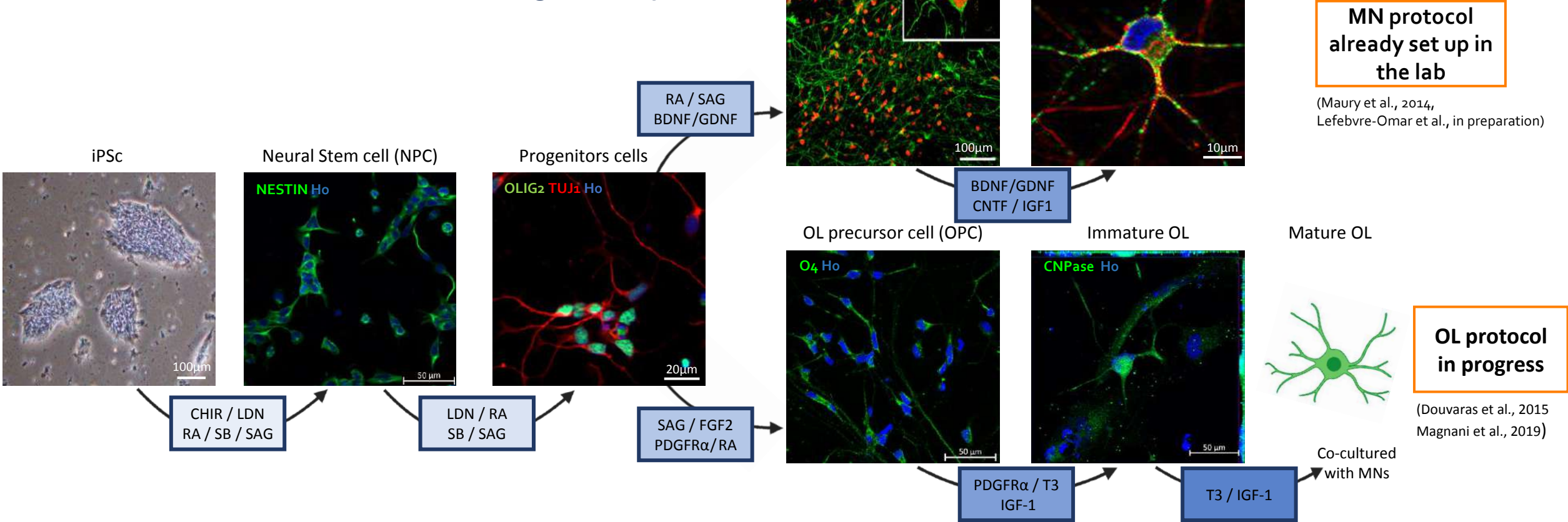
Age and sex matched healthy **controls**
Isogenic controls

ALS patients mutated in :
• SOD1
• TDP-43
• FUS
• C9ORF72



PROTOCOLS

● **iPSc differentiation into motor neurons and oligodendrocytes**



RESULTS in MNs

USP19 mRNA and protein expression in progenitors and MNs, and its colocalization with endoplasmic reticulum

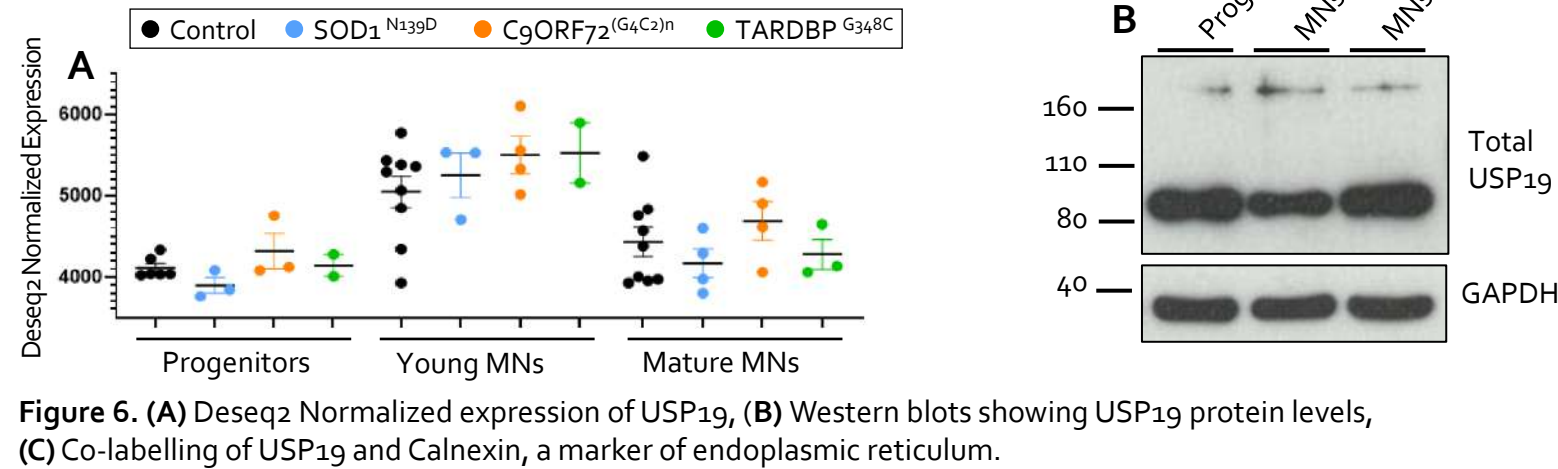
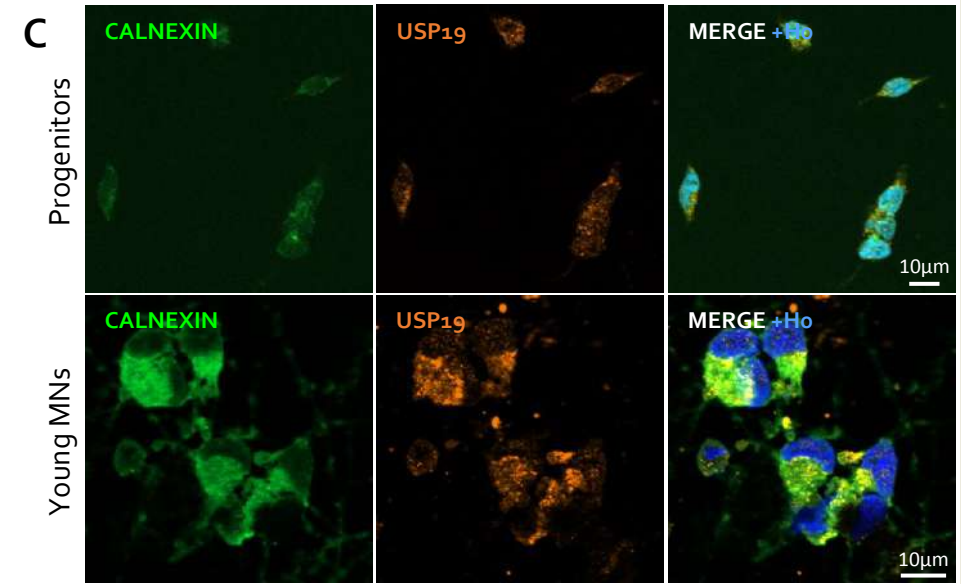


Figure 6. (A) Deseq2 Normalized expression of USP19, (B) Western blots showing USP19 protein levels, (C) Co-labelling of USP19 and Calnexin, a marker of endoplasmic reticulum.



Expression of ER transmembrane USP19 in MNs

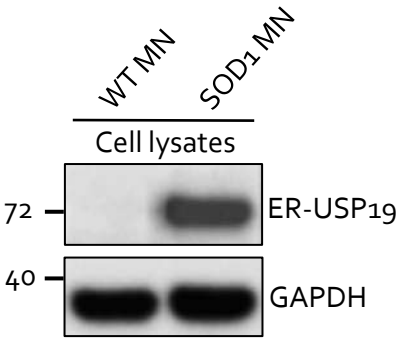


Figure 7. Western Blot showing the expression of transmembrane USP19 only in MNs SOD1, (GAPDH normalizer)

Pathological determinants in MNs

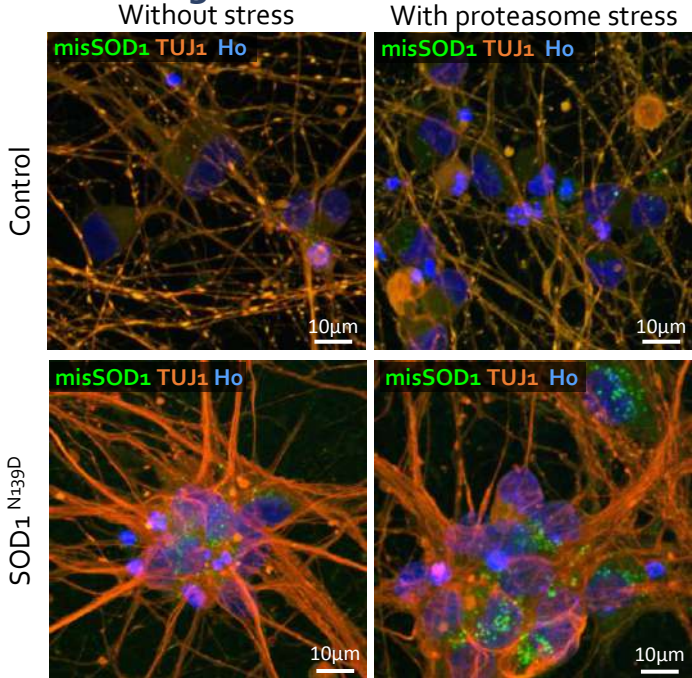


Figure 8. Immunolabelling of misfolded SOD1 in MNs

Secretion of SOD1 in the supernatant of control and SOD1 MNs

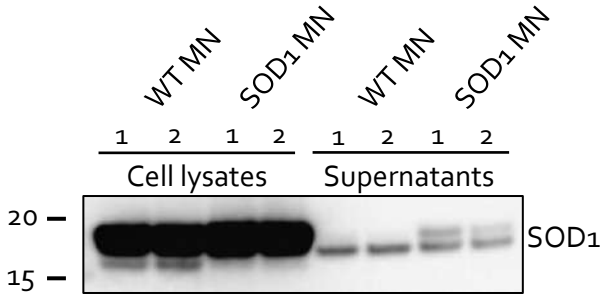


Figure 9. Western Blot of cell lysates and supernatants purified by ultracentrifugation (Théry et al., 2006) of MNs. SOD1 is expressed in cell lysates and is secreted in supernatants.

CONCLUSIONS

- USP19 is expressed in iPSc, progenitors and mature MNs and is expressed near the ER.
- ER transmembrane USP19 is detected exclusively in SOD1 MNs.
- Both control and mutant SOD1 MNs secrete SOD1 in the supernatant.
- A protocol to differentiate iPSc into OPCs and OLs is in progress, coculture will be done with MNs.

PERSPECTIVES

- Search for inclusions (SOD1, TDP₄₃, FUS and DPR) in ALS iPS-derived MNs, OPCs and OLs (normal or stress conditions).
- Analyze the secretion of pathological determinants in MNs, OPCs and OLs.
- Assess the specific USP19 pathway with USP19 siRNA.
- Analyze whether propagation occurs between MNs and OPCs or OLs.