

Is there a biochemical or molecular switch from healthy to symptomatic phenotype in Amyotrophic Lateral Sclerosis?



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BACKGROUND

ALS is a fatal neurodegenerative disease, characterized by the loss of MNs¹.10% of ALS cases are associated with mutations in specific genes (familial, fALS) and patients display diverse phenotypic expression patterns and may exhibit different progression rate. Although the effects of the mutations are not completely clear, they share some common pathways: glutamate excitotoxicity, structural and functional abnormalities of mitochondria, oxidative stress, intracellular aggregates, activation of microglia.

EXPERIMENTAL PLAN



d LP, et al., Shneider NA. (2001). Amyotrophic lateral sclerosis. N Engl J Med; J., & Zhang, Z. (2019). Structural properties and interaction partners of familial ALS-SOD1 mutants. Frontiers in Neurology:

3) Mitsuzawa, S. et al. (2018). TARDBP p.G376D mutation, found in rapid progressive familial ALS, induces mislocalization of TDP-43. ENeurologicalSci.

AIMS

Comparison between healthy and symptomatic fibroblasts with mutations in SOD1² and TARDBP³ genes to investigate the existence of a molecular or biochemical switch from sane to affected phenotype, thanks phenotypical, biochemical, to and molecular analysis.



Although there are no significant differences in ROS levels, healthy individuals have higher GSH levels compared to symptomatic ones and this could represent a protective mechanism (p value: *p<0,05; **p<0,01;***p<0,001).

SOD

TARDBP

NEXT STEPS: Seahorse analysis to study bioenergetic parameters (Glycolysis VS OXPHOS), since the normal process of ETC is perturbed in ALS patients, causing less production of ATP.

TARDBP

SOD