

In the last few years, there has been a rapid advance in our knowledge of genetic causes for ALS. Furthermore, the relationship between the genetic subtypes and the pathological subtypes as well as clinical phenotype has become more and more clear. In addition to superoxide dismutase 1 (SOD1), mutations in the genes coding for TAR DNA-Binding Protein (TARDBP), fused in sarcoma (FUS), Ubiquilin2 (UBQLN2), C9ORF72 and several others are closely associated with typical clinical phenotype. Figure 1 provides up-to-date findings of genetic

defects in ALS and the underlying mechanisms for the cause and pathogenesis of the disease.

Genetics of fALS

The causative genes have been identified in almost 5-10% of all fALS cases to date [4,5]. Among those 20% of fALS cases are caused by the mutation in SOD1 gene, 4-5% of fALS cases are the results of mutations in TARDBP and FUS genes, more than 30% of fALS cases are associated with C9ORF72 mutations and the rest are

