





Fig. 2. Amyloid plaques in the brain of an early-onset Alzheimer's patient carrying a mutation in the *SORL1* gene

In total, we identified 92 rare genetic variants in patients and control individuals. Rare genetic variants in *SORL1* were more common in the patients (8.8% carried a rare *SORL1* variant) than in control individuals (5.6%). Remarkably, we observed a specific type of genetic variants, known as protein truncating variants, exclusively in patients. These protein truncating variants are predicted to lead to either a heavily altered gene product, or to loss of protein, thus abolishing the protective effect of the *SORL1* protein.

Most carriers of these mutations had – except for their early onset - a typical course of disease, with memory problems dominating their symptomatology. However, patients with a *SORL1* mutation were more likely to have a familial history of Alzheimer's disease, further implicating a role of *SORL1* in the heritability of AD.

In conclusion, while most rare genetic variants in the *SORL1* gene are either neutral or mildly risk-increasing, variants predicted to lead to loss of functional protein are likely to be stronger risk factors that may contribute to familial occurrence of AD. This implicates that *SORL1* plays an important role in the development of Alzheimer's disease, and could possibly be a target for novel therapies. Further research is needed to see if carrying these mutations can predict the future development of the disease, and if genetic tests of *SORL1* can be used in clinical practice.

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## Publication

[A comprehensive study of the genetic impact of rare variants in SORL1 in European early-onset Alzheimer's disease.](#)

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