

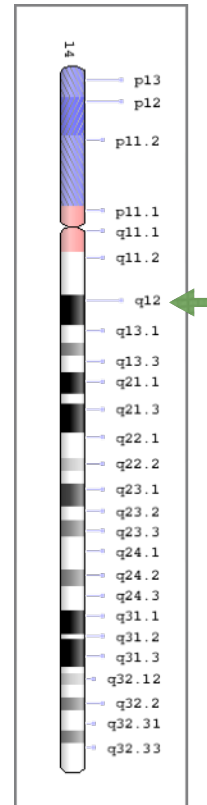


DELETION 14Q12

Deletion 14q12 (FOXG1-Syndrome)

The wide implementation of molecular cytogenetics which started ~15 years ago drastically changed our perception of what is known as a “chromosomal disorder”. Routine cytogenetics can recognize only relatively large deletions or duplications, and most syndromes reported in the “pre-molecular” era involve abnormalities affecting at least 10 Mb of genetic material. Such large segments usually contain several disease-causing genes. The introduction of molecular methods showed that basically each chromosome carries several morbid genes, and the malfunction of any of these genes may cause a specific genetic (chromosomal) disorder. Chromosome 14 is no exception.

The segment 14q12 (an area of the long arm located close to the centromere) contains the FOXG1 gene. This tiny gene (it has only one exon) encodes a developmental transcription factor with repressive activity important for the development of the telencephalon (brain)¹. If this gene does not work it causes a wide spectrum of neurodevelopmental problems. Patients with FOXG1 syndrome reveal microcephaly (sometimes congenital, but sometimes postnatal), poor eye contact, absent language, irritability, sleep disturbances, frequent gastro-esophageal reflux, seizures starting in infancy, and early-onset movement problems. The movement disorder may include generalized chorea, dystonia, facial dyskinesias and myoclonic jerks². These manifestations are similar to the so-called Rett syndrome and before the implementation of molecular techniques, such patients were considered to have a congenital variant of Rett syndrome³ (in classic Rett syndrome, caused by a mutation of the MECP2 gene, patients start their development as normal, but show a loss of acquired skills after several months). An MRI examination



of patients with FOXP1 syndrome shows agenesis or hypoplasia of the corpus callosum in almost all patients. Other defects include delayed myelination and a simplified gyral pattern. These manifestations are basically the same for patients with mutations of the FOXP1 gene and for patients with deletions of this gene⁴. Currently there are at least 60 publications detailing patients who have FOXP1 syndrome caused by 14q12 deletion. The segment 14q12 does not contain many genes. The frequent involvement of the neighboring NOVA1 gene does not produce any additional abnormalities. Several patients with deletions in close proximity to FOXP1 and not directly affecting the gene had the same manifestations as typical FOXP1 deletion patients. Most likely, deletions in these cases affect areas containing elements controlling activity of the FOXP1 gene.

There are no defects outside the nervous system in patients with 14q12 deletions. Of course, if deletions are relatively large and involve more distal segments of 14q (14q13 or even 14q21) the patients may show defects in other systems or other manifestations unrelated to FOXP1 syndrome.

All known 14q12 deletions causing FOXP1 syndrome are sporadic. However, the possibility of an unrecognized low-level mosaicism in one parent cannot be excluded.

REFERENCES

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² Caporali C, Signorini S, De Giorgis V. et al. (2018). Early-onset movement disorder as diagnostic marker in genetic syndromes: three cases of the FOXP1-related syndrome. *Europ J Paediatr Neurology* v. 22, 336-330.

³ Venugopal VS, Dutta UR, Tallapaka K. et al. (2018). Whole exome sequencing identifies a novel 5 Mb deletion at 14q12 region in a patient with global developmental delay, microcephaly and seizures. *Gene* v. 673, 56-60.

⁴ Lu G, Zhang Y, Xia H. et al. (2022). Identification of a de novo mutation of the FOXP1 gene and comprehensive analysis for molecular factors in Chinese FOXP1-related encephalopathies. *Front Molec Neuroscience* v. 15:1039990.