

Poster presentation

Prenatal diagnosis of L1CAM gene mutations in X-linked hydrocephalus

Mami Yamasaki*¹, Tomoko shohuda², Hiroaki Sakamoto³,
Masahiro Nonaka¹ and Yonehiro Kanemura²

Address: ¹Department of Neurosurgery, Osaka National Hospital, National hospital organization, 2-1-14 Hoenzaka, Chuo-ku, osaka city, Osaka, 540-0006, Japan, ²Institute for clinical research, Osaka National Hospital, National hospital organization, 2-1-14 Hoenzaka, Chuo-ku, osaka city, Osaka, 540-0006, Japan and ³Department of Pediatric Neurosurgery, Osaka City General Hospital 2-13-22 Miyakojima-hondouri, Miyakojima-ku, osaka city, Osaka, 534-0021, Japan

Email: Mami Yamasaki* - yamasaki@onh.go.jp

* Corresponding author

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Background

X-linked hydrocephalus (XLH) (severe type of human L1 syndrome) is now known to be due to mutations in the gene for the neural cell adhesion molecule L1. We performed prenatal diagnosis of L1CAM gene mutations in 5 families. We evaluated effective methods and discussed contribution for the prenatal diagnosis of XLH.

Materials and methods

We performed a nation-wide L1 gene analysis of patients with hydrocephalus and identified L1 gene mutations in 36 families. In these families, five obligate carriers were pregnant subsequently and want to perform the L1CAM gene analysis of their fetuses. Genomic DNA was extracted from chorionic villus biopsy (CVB) at from 10 to 15 weeks' gestations. Amplification of the exons and the exon-intron boundaries of the L1 gene was performed by polymerase chain reaction (PCR). Purified PCR amplification products were directly sequenced using the ABI BigDye™ Terminator Cycle Sequencing Ready Reaction Kit (Applied Biosystems) and analyzed with a capillary DNA sequencer ABI PRISM® 310 Genetic Analyzer.

Results

1. Two fetuses were male and three were female. L1CAM gene in two males did not have mutations. In three

females, one did not carry the mutation in L1CAM gene and two female fetuses had the same L1CAM gene mutation as his mother. 2. Five obligate carriers continued their pregnancy and delivered normal babies.

Conclusion

Prenatal L1 gene analyses are useful for the prenatal diagnosis of X linked hydrocephalus.