A novel mutation in L1CAM causes a mild form of L1 syndrome: a case report

Maarten Otter1,2, Marijke Wevers3,4, Marline Pisters1,†, Rolph Pfundt5, Yvonne Vos6, Rutger Jan Niewelstein7 & Constance Stumpel8

1Department of Community Mental Health in Learning Disabilities, Trajectum, Zutphen, The Netherlands
2Forensic Psychiatric Department, STEVIG/Dichterbij, Oostrum, The Netherlands
3Department of Clinical Genetics, University Medical Center, Utrecht, Utrecht, The Netherlands
4Department of Clinical Genetics, Radboud University Medical Center, Nijmegen, The Netherlands
5Department of Human Genetics, Radboud University Medical Center, Nijmegen, The Netherlands
6Department of Genetics, University Medical Center, Groningen, Groningen, The Netherlands
7Department of Radiology and Nuclear Medicine, University Medical Center, Utrecht/Wilhelmina Children’s Hospital, Utrecht, The Netherlands
8Department of Clinical Genetics and School for Oncology and Developmental Biology, Maastricht University Medical Center, Maastricht, The Netherlands

Correspondence
Maarten Otter, Department of Community Mental Health in Learning Disabilities, Trajectum, Zutphen, The Netherlands.
Tel: +31-88-9296703;
Fax: +31-575-576777;
E-mail: maartenotter@gmail.com

Present address
†Dr. Leo Kannerhuis, Jonkheer Nedermeijer van Rosenthalweg 16, 6862 ZV, Oosterbeek, The Netherlands

Key Clinical Message
Clinical geneticists, neurologists, psychiatrists, and other healthcare providers can learn from this case report that patients with a behavioral phenotype that includes a mild learning disability may also require a thorough examination, including brain MRI and whole-exome sequencing.

Keywords
Behavioral phenotype of genetic syndromes, corpus callosum hypogenesis, L1CAM mutation, X-linked mental retardation.

Introduction
Mutations in the gene encoding L1 cell adhesion molecule (L1CAM) are phenotypically characterized by X-linked hydrocephalus (OMIM #307000) or—in milder cases—corpus callosum agenesis or hypogenesis (OMIM #304100) [1]. The variability in clinical presentation reflects different functions of L1CAM. Hydrocephalus is a very complex disorder. Mutations in L1 as a cell adhesion molecule play a role in the pathogenesis of hydrocephalus. The corpus callosum hypogenesis reflects dysfunction as a neural recognition molecule through disorders in axonal growth and guidance [2].

The L1CAM gene is located on the X chromosome at Xq28 and is expressed primarily in the nervous system, where it plays important roles in neuronal development, including the guidance of neurite outgrowth, neuronal cell migration, axon bundling, synaptogenesis, myelination, neuronal cell survival, and long-term potentiation [3].
The patients with hydrocephalus have a worse prognosis [4]. This is in agreement with recent evidence that some forms of hydrocephalus and the inborn neurological impairment are inseparable phenomena. This is because both fetal onset hydrocephalus and abnormal neurogenesis have a common origin: a primary pathology of the embryonic ventricular zone, involving neural stem cells and multiciliated ependymal cells. Junctional failure and loss of neural stem cells underpin the etiology of neuronal migration defects at specific regions of brain. Thus, neonates are born with a neurodevelopmental deficit that derivative surgery cannot solve [5, 6].

Here, we report a patient with a generally mild physical presentation but with a more distinct behavioral phenotype.

Clinical Description

The proband (referred to here as “Patient A”) is a 56-year-old man born in a family with two brothers from nonconsanguineous healthy parents. One of his two brothers has a learning disability of unknown severity; the other does not. Their mother had 14 siblings; three of her brothers had a learning disability. Patient A’s parents died around the age of 75. Patient A’s daughter developed epilepsy at the age of 6; she initially experienced absence seizures, and later developed grand mal seizures. For the past 2 years, she has taken valproate and has been seizure-free. Patient A’s son is healthy.

The proband sought help from professional caregivers due to anxiety and depression following several major life events. He was fired 10 years ago and was unable to find a new job until he found employment at a sheltered workplace as a postal employee. In addition, he experienced the deaths of his parents and his parents-in-law. Patient A also experienced difficulty remembering essential information during work and at home. He lives with his wife and two adult children in the center of a large city in the Netherlands, where he was born. The patient requested a psychological and neurological evaluation because he experienced a subdural hematoma with cerebral contusion following an unexpected fall when he was 28 years of age. Although the subdural hematoma was caused by an epileptic seizure, as he frequently experienced unexpected falls.

The results of a psychological examination revealed overall difficulties with cognitive functioning, including impaired memory, attention, and executive functions. The patient’s clinical history and test results revealed that psychological problems appeared at an early age, leading to social, emotional, and behavioral problems. Therefore, combined with his intellectual challenges, the patient lacked sufficient opportunities to learn the appropriate skills needed to cope during complex, stressful situations. These developmental issues, limited coping skills, and limited opportunities to express himself emotionally likely predisposed the patient to develop depression, particularly when experiencing stress due to a lack of structure in his daily life, thereby having a strongly disrupting effect on daily functioning and activities (Table 1).

Physical examination revealed a healthy adult male with the following parameters: height, 177.5 cm (–1 SD); weight, 77.6 kg (+1.3 SD); occipito-frontal circumference, 57.3 cm (–0.33 SD); left ear length, 7.4 cm (75th–97th percentile); left hand length, 21.1 cm (above the 97th percentile); and mildly adducted thumbs. The patient presented with no gait abnormalities, and a neurological examination was unremarkable.

In addition, an MRI examination of the brain revealed hypogenesis of the corpus callosum, particularly in the splenium (Fig. 1), as well as signs of prior traumatic injuries to the right frontal brain parenchyma and mild, slightly asymmetric dilatation of the lateral ventricles due to underdevelopment and/or loss of white matter (Fig. 2); the loss of white matter in the right hemisphere may have been due in part to the prior traumatic brain injury.

A genetic examination was also performed. SNP array analysis revealed a normal male karyotype. Whole-exome sequencing (WES) was performed essentially as reported previously [7]. Exome capture was performed using the Agilent SureSelect Human All Exon v4 enrichment kit (Agilent Technologies, Santa Clara, CA), and WES was performed using the Illumina HiSeq platform (BGI, Copenhagen, Denmark). Data were analyzed using the BWA (read alignment) and GATK (variant calling) software packages. Variants were annotated using an in-house-developed pipeline and prioritized using an in-house-designed “variant interface” with manual curation. Putative causative variants were confirmed by Sanger sequencing. WES revealed the missense mutation c.998C>T (p.Pro333Leu) in the proband’s L1CAM gene (NM_000425.3). Segregation analysis revealed that the same mutation was present in the proband’s brother with a learning disability; in contrast, the proband’s other brother—who did not have developmental abnormalities—did not carry this variant. It was not possible to obtain
genetic material for testing the proband’s deceased mother—who would have been an obligate carrier of the variant—or her brothers with mental disability.

This missense variant causes the substitution of a highly conserved nucleotide (phyloP: 5.29) and amino acid residue in the immunoglobulin-like domain of the L1CAM gene and protein, respectively. This mutation has not been described previously in any SNP database, including dbSNP and the EXAC database (http://exac.broadinstitute.org/); moreover, several mutation prediction tools predict that this variant is likely to be pathogenic.

Discussion

In this clinical report, we present a male patient with a novel, putative pathogenic variant in the L1CAM gene. This variant affects the structure of the fourth immunoglobulin-like domain in the protein, as Pro333 is believed to be one of the key residues required for proper folding of this domain [8]. Therefore, this variant likely disrupts interactions between both L1-L1 proteins and L1-non-L1 proteins, interactions that are important for protein function. Interestingly, a different variant in the same amino acid (p.Pro333Arg) was described in a patient with X-linked hydrocephalus; this patient died before the age of 1 [4]. As we mentioned before, some forms of hydrocephalus are not just ventriculomegaly, but the tip of the iceberg of a complex neurodevelopmental disorder [5, 6].

In the family reported here, the c.998C>T variant was present in both the proband and his affected brother, but
not in the proband’s unaffected brother. Both the patient and his affected brother presented with a rather mild clinical course. In most families with L1 syndrome, the affected boys die before, during, or soon after birth; however, both interfamily variability and intrafamily variability have been reported [9]. Basel-Vanagaite et al. described two half-brothers who had a mutation in the L1CAM gene and a hypoplastic corpus callosum, but no other characteristics of L1 syndrome [1]. Epilepsy has also been reported in a minority of patients with L1 syndrome [10].

In our study, the proband presented with a hypoplastic corpus callosum and some minor clinical features of L1 syndrome. In this case, the patient’s behavioral phenotype appeared to be more severe and distinct than his physical phenotype. Because social delays and behavioral abnormalities have been described in patients with corpus callosum abnormalities, an integrative examination of these functions should be included in the diagnostic protocol [11].

The value to this family from obtaining this diagnosis should not be underestimated. For example, several family members are now comforted by the fact that they—and their children—do not have an increased risk of recurrence. In addition, other family members—particularly the proband’s adolescent daughter—now have the opportunity to seek clinical genetic advice when deciding whether to have children. Moreover, this family has learned that they should not expect the proband to achieve a higher level of functioning, and they have learned not to overestimate his abilities as either a husband or a father. After receiving this genetic diagnosis, his primary clinical goal—namely to help resolve his depressive symptoms—has waned.

Based on these findings, clinicians should be aware of the value of performing neuroimaging in persons with developmental disabilities, and they should offer their patients the opportunity to seek an etiological diagnosis, including genetic testing. Psychiatrists—particularly those who work in the field of intellectual disabilities/learning disabilities—should be made aware of the options available regarding diagnostic procedures based on clinical genetics and neuroimaging. These procedures can provide evidence-based assistance to patients and their families, including preventive management, thereby improving quality of life.

Authorship

MO: main author, was responsible for writing and reviewing the manuscript, literature review, and provided direct neuropsychiatric care to the patient. MW: participated in study design, coordination, and data analysis, especially concerning clinical genetic aspects, and provided direct clinical genetic care to the patient and his family. MP: performed the psychological assessment and wrote Table 1. RP: performed the WES and analyzed the molecular data and co-authored the description of the molecular details in the manuscript. YV: analyzed the molecular data and co-authored the description of the molecular details in the manuscript. RJN: reviewed the MRI of the brain and provided the figures. CS: involved in overall supervision of the writing of the manuscript.

Acknowledgement for publication

This study makes use of data shared through the PhenomeCentral repository. Funding for PhenomeCentral was provided by Genome Canada and Canadian Institute of Health Research (CIHR).

Conflict of Interest

The authors declare to have no conflict of interest.
References


