Megalencephaly and Epileptic Encephalopathy: Bad Luck or a Common Pathway?

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Abstract

Megalencephaly is a neuronal migration disorder characterized by an abnormally large brain. Numerous associated syndromes and various molecular mutations have been identified as an etiology for megalencephaly, however, SCN2A mutations have not been previously described. This report highlights a case of a term male megalencephalic neonate who presents with intractable seizures, who was found to have SCN2A gene variant that has now been identified as pathogenic. This patient expands our knowledge of the phenotypic spectrum of SCN2A mutations by adding consideration for macroscopic brain findings. Currently, we have no direct link between SCN2A mutations and megalencephaly, but our patient highlights the potential overlap in disease processes. It is possible that the biochemical disturbance associated with abnormal neuronal migration also affects the neuronal circuitry, thus increasing the propensity for electrical dysfunction and manifesting as seizures.

Keywords: Megalencephaly; Epileptic encephalopathy; SCN2A mutations; Neuronal migration

Introduction

Bilateral megalencephaly is a rare malformation of the central nervous system that affects 1 out of 50 children in the general. It is a disorder of abnormal brain proliferation and neuronal cell migration, defined as brain weight more than 2 standard deviations above the agerelated mean [1]. The PI3K-AKT pathway mutation is known to cause two clinically distinct syndromes: megalencephaly-capillary malformation (MCAP) and megalencephaly-polydactyly-polymergyria-hydrocephalus (MPPH). MCAP is characterized by cutaneous vascular malformations, connective tissue dysplasia, and somatic overgrowth which distinguish it from MPPH. Seizure is not a classic presentation of either syndrome [2]. We report a novel SCN2A gene variant association in a neonate with megalencephaly and intractable seizures.

Case Presentation

A term, small for gestational age, non-dysmorphic male infant was born to non-consanguineous parents at 37 weeks gestation via Cesarean section for breech presentation. His birth weight was 2422 g, length 45.8 cm head circumference 33 cm and Apgar scores were 5 at 1 minute, 9 at 5 minutes. Physical examination was normal. He received routine delivery room care and was admitted to the newborn nursery. Around 24 hours of age, the infant was admitted to the Neonatal Intensive Care Unit (NICU) for the management of hypoglycemia. While in the NICU, he developed multiple bradycardic episodes which progressively increased in frequency, prompting further evaluation.

Blood work was significant for hyponatremia with sodium of 122 with normal renal function, calcium, phosphorus, magnesium, CBC, differential and platelet counts. EKG showed normal sinus rhythm. Head ultrasound was normal. Electroencephalography (Figures 1 and 2) confirmed malignant migrating partial seizures correlating with clinical bradycardic events that were refractory to multiple anti-epileptic medications including: two Phenobarbital loading doses at 20 mg/kg/dose with maintenance dosing at 4 mg/kg/day twice daily, Keppra 60 mg/kg/day divided twice daily, Leucovorin 5 mg twice daily, Topamax 10 mg/kg, Pyridoxal-5-phosphate 20 mg every 6 hours, Bumetanide infusion at 0.1 mg/kg/hour, and continuous versed infusion running at 1.5 mg/kg/hour. He was also given fosphenytoin a single loading dose of 20 mg/kg/dose and subsequent load of 10 mg/kg/dose when seizures persisted. Ultimately, further fosphenytoin doses were held due to supratherapeutic levels.

He required intubation and ventilatory support for respiratory depression in the setting of multiple medications required for treatment of his intractable seizures. MRI demonstrated no structural abnormalities but multiple areas of bilateral punctate white matter injury with increased T1 signal, some with restricted diffusion (Figures 3 and 4). CSF studies were unremarkable. Due to the presence of persistent seizures, refractory to escalating doses of anti-epileptic medications, the possibility of a genetic epilepsy syndrome was considered. Despite maximum medical intervention the infant’s status deteriorated. After discussion with family about the guarded long-term outcome, the infant’s care was redirected to comfort measures only. Infant was terminally extubated and died on day of life 10.

Autopsy reported megalencephaly with neonatal brain weight of 500 grams, olfactory bulb hypertrophy and dysplasia, multifocal dysplastic areas of cortex and widespread subacute destructive processes in the hippocampal complex, dorsal thalamus, optic nerve, cerebellar purkinje layer with evidence of focal white matter infarcts (Figures 5 and 6). Serum epilepsy panel revealed SCN2A mutation. Whole exome sequencing performed on brain tissue reported infant was heterozygous for a de novo L210R pathogenic variant in the SCN2A gene.

Discussion

When considering megalencephaly, discovering the underlying etiology can often be challenging. Although previous reports have implicated mutations in the PI3K-AKT pathway, we now know there is significant heterogeneity in etiology and presentation. Over the

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Figure 1: Electroencephalography (EEG) demonstrating migrating partial seizures.

Figure 2: Electroencephalography (EEG) demonstrating abnormal interictal period.
Figure 3: T1 sagittal and axial MRI images demonstrating multiple punctate areas of increased T1 signal consistent with white matter injury.

Figure 4: Gross anatomy of brain: The 500-gram formalin fixed neonatal brain is depicted below. The leptomeninges are thin and transparent. There is no appreciable vascular congestion. No evidence of brain edema or herniation is found. Some of the gyri appear thick, but not pachygyric. There is no abnormality patterning. The vessels distribute normally at the base of the brain.
past century, numerous associated syndromes and many molecular mutations have been identified as causes for megalencephaly but SCN2A mutations have not been previously reported [1-3].

The SCN2A gene encodes the alpha subunit of the voltage gated sodium channel which is involved with neuronal firing [4]. SCN2A variants are inherited in an autosomal dominant manner. Pathogenic variants causing encephalopathy are typically de novo, while inherited variants often result in milder phenotypes and may exhibit incomplete penetrance [5]. To our knowledge, the L210R variant in the SCN2A gene has not been classified as a pathogenic variant prior to this patient.
On review of large scale human genomic datasets, we know that the L210R variant is not observed in large population cohorts [6]. The L210R variant is a non-conservative amino acid substitution, which is likely to impact secondary protein structure as these residues differ in polarity, charge, size and/or other properties. This substitution in p. Leu210Arg (L210R) (CTG>GCG): c.629 T>G in exon 6 of the SCN2A gene is predicted to be within the extracellular loop between the S3 and S4 transmembrane segments of the first homologous domain. Additionally, missense variants in nearby residues that have a gain-of-function effect lead to a spectrum of SCN2A-related disorders, ranging from benign familial infantile-onset seizures (BFNIS) to epileptic encephalopathies, including Ohtahara syndrome and epilepsy of infancy with migrating focal seizures (EIMFS) [4,6,7]. Therefore, in consideration of the available literature in combination with the clinical context, we conclude L210R as a pathogenic variant and the likely cause of refractory seizures in our patient. In our case, there was a lack of phenotypic diagnostic clues given the infant’s normal prenatal anatomy scan, normal physical exam and non-specific brain imaging. Therefore, if the clinical suspicion is high, genetic testing via the comprehensive epilepsy panel should be considered for a definitive diagnosis.

To date, there is sparse data on an association between SCN2A-related disorders and megalencephaly. There is no comprehensive overview of possible SCN2A phenotypes, and the genotype-phenotype correlation remains poorly understood. In general, there are 4 recognized phenotype groups: 1) benign neonatal or infantile epilepsies, 2) neurodevelopmental/neuropsychiatric disorders, including schizophrenia, autism and intellectual disability, 3) Infantile Spasms, and 4) early onset epileptic encephalopathies. Severe neonatal epileptic encephalopathies can also be associated with cortical abnormalities including cortical dysplasia but megalencephaly has not been described [8]. Similarly, there have been case reports with severe non-SCN2A related epilepsy as a consequence of hemi-megalencephaly [7]. Our findings support a previous case described by Bernado of a neonate with SCN2A epileptic encephalopathy with severe cortical dysplasia. The diagnosis was heralded by a pre-natal MRI at 32 weeks which demonstrated dysplasia, corpus callosum dysgenesis and pachygyria. Our case highlights a severe presentation in the absence of abnormal prenatal and post-natal MRI findings. The identified variant further adds to our knowledge of the vast spectrum of SCN2A-related disorders and highlights the need for continued investigation and discovery.

**Conclusion**

Our findings support a previous case described by Bernado of a neonate with SCN2A epileptic encephalopathy with severe cortical dysplasia. The diagnosis was heralded by a pre-natal MRI at 32 weeks which demonstrated dysplasia, corpus callosum dysgenesis and pachygyria. Our case highlights a severe presentation in the absence of abnormal prenatal and post-natal MRI findings. The identified variant further adds to our knowledge of the vast spectrum of SCN2A-related disorders and highlights the need for continued investigation and discovery.

**References**