

# Delayed SCN1A Diagnosis: A Case Series of Three Families

HARVARD MEDICAL SCHOOL TEACHING HOSPITAL

Emma Sexton BS, Lacey Smith MS, CGC, Hyun Yong Koh MD, PhD, Devon Knight BS, Rebecca Pinsky MSN, RN, CPNP, Chellamani Harini MD, Ann Marie Bergin MB, ScM, MRCP (UK), Beth Rosen Sheidley MS, CGC, and Annapurna Poduri MD, MPH

## Introduction

- The SCN1A gene encodes the alpha 1 subunit of the sodium channel.<sup>1</sup>
- Variants in SCN1A have been well-described and are related to a variety of epilepsy phenotypes ranging from genetic epilepsy with febrile seizures plus (GEFS+) to developmental and epileptic encephalopathies, such as Dravet Syndrome.<sup>1</sup>
- Recommendations for diagnosis and management of Dravet Syndrome are available and obtaining a genetic diagnosis can impact clinical management.<sup>2</sup>
- Although next generation sequencing has a high diagnostic yield<sup>3</sup>, many children with early onset and childhood epilepsies do not undergo early diagnostic testing.

# Objective

 To assess the reasons for, and potential clinical impact of, delayed genetic diagnosis of SCN1A-related epilepsy.

## **Methods**

- We enrolled over 500 probands and their available, biological parents in the Children's Rare Disease Cohorts Initiative (CRDCI) at Boston Children's Hospital.<sup>4</sup>
- Proband inclusion criteria included having a diagnosis of epilepsy with unknown etiology and normal brain magnetic resonance imaging.
- Whole exome sequencing was completed for probands and their biological parents.
- Phenotypic data and clinical history notes were reviewed using the electronic medical record, and data was organized and managed using REDCap.

#### References

- ¹ Scheffer, I. E., & Nabbout, R. (2019). SCN1A-related phenotypes: Epilepsy and beyond. Epilepsia, 60 Suppl 3, S17–S24.
  ² Wirrell, E. C., et al. (2017). Optimizing the Diagnosis and Management of Dravet Syndrome: Recommendations From a North American Consensus Panel. Pediatric neurology, 68, 18–34.e3.
- <sup>3</sup> Costain, G. et al. (2019). Clinical Application of Targeted Next-Generation Sequencing Panels and Whole Exome Sequencing in Childhood Epilepsy. Neuroscience, 418, 291–310.
- Sequencing in Commission Epileps, Neuroscience, 418, 291–310.

  \*\*Rockowitz, S., et al. (2020). Children's rare disease cohorts: an integrative research and clinical genomics initiative.

  \*Npj Genomic Medicine, 5(1), 29.

**Table 1.** Epilepsy diagnosis, age at seizure onset and genetic diagnosis, and total time passed from age at seizure onset to age at genetic diagnosis for each participant.

Participant	Epilepsy Diagnosis	Age at Seizure Onset	Age at Genetic Diagnosis	Time Elapsed
1	Dravet	9 months	4 years	3.4 years
2	Dravet	6 months	10 years	8.3 years
3	Dravet	6 months	11 years	9.8 years
4	Benign Occipital Lobe Epilepsy	4 years	7 years	3.9 years

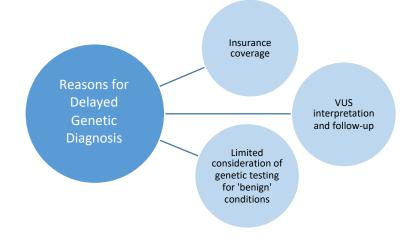


Figure 1. Three factors contributing to delayed genetic diagnosis for the four participants.

Table 2. Genotypic information and ACMG classification for participant variants in SCN1A.

Pa	rticipant	Variant	Protein Change	Classification	Inheritance
	1	c.664C>T	p.R222X	Pathogenic	De novo
	2	c.5495C>A	p.A1832E	Likely Pathogenic	Maternal (mosaic)
	3	c.5495C>A	p.A1832E	Likely Pathogenic	Maternal (mosaic)
	4	c.5066T>C	p.M1689T	Likely Pathogenic	De novo

## Results

- We identified pathogenic or likely pathogenic SCN1A variants in 4 children with epilepsy (Table 2).
- Time elapsed between age at seizure onset and age at genetic diagnosis of SCN1A variants ranged from 3.4 to 9.8 years (Table 1).
- Genetic testing was attempted clinically for Participants 1, 2, and 3 prior to enrollment in the CRDCI.
  - For Participant 1, despite clinical suspicion of a SCN1A-related etiology, clinical testing was not obtained due to insurance coverage.
  - Participants 2 and 3 are biological brothers. A clinical gene panel was pursued for Participant 3 (the older sibling) and a variant of uncertain significance (VUS) was identified in SCN1A. Segregation analysis was not pursued at the time.
  - Clinical genetic testing was not pursued for Participant 4 as the healthcare provider did not consider his relatively benign phenotype to be suggestive of a genetic etiology.
- Reasons for delayed diagnosis for this cohort include: 1) insurance coverage, 2) lack of provider familiarity with variant classification, VUS interpretation, and the need for re-analysis, and 3) limited clinical consideration for genetic testing for relatively 'benign' epilepsy phenotypes (Figure 1).

### Conclusions

- Definitive diagnosis of SCN1A-related epilepsy varied considerably in the 4 cases identified through this institutional sequencing initiative.
- Delayed diagnosis can impact an individual and family's ability to participate in research and clinical trials, receive counseling regarding risk recurrence, and engage with advocacy groups and community resources.
- Recommendations include early consideration of genetic testing for
  patients with unexplained epilepsy, and routine re-analysis of VUS and
  additional testing strategies for those who remain 'unsolved' after initial
  evaluation.

## Acknowledgements

We would like to thank the research participants and their families, and the Epilepsy Genetics Program at Boston Children's Hospital.