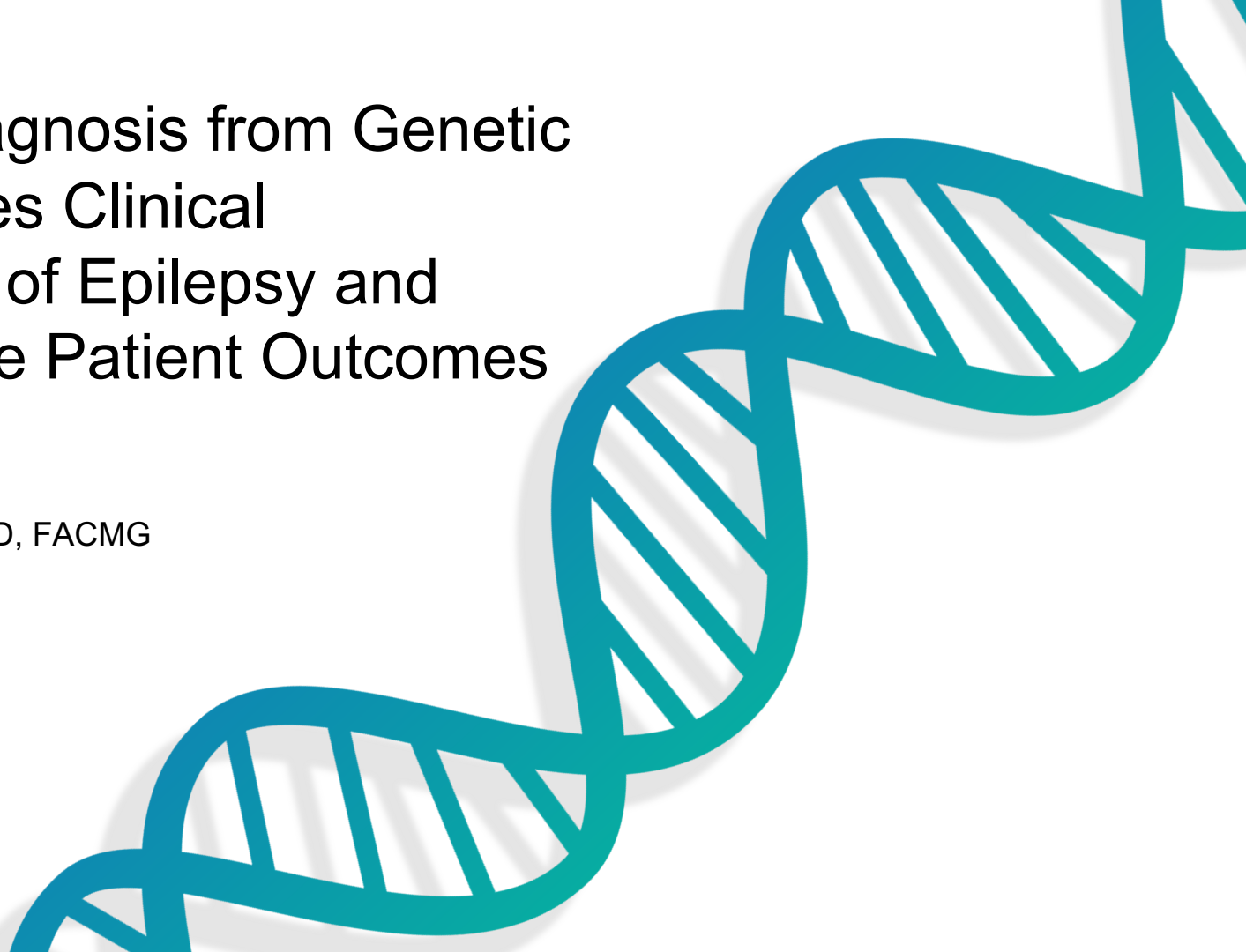


# Molecular Diagnosis from Genetic Testing Guides Clinical Management of Epilepsy and Helps Improve Patient Outcomes

Dianalee McKnight, PhD, FACMG  
Medical Affairs Director  
Invitae



# Learning Objectives

- List examples of patient management changes that can be made by clinicians based on a molecular diagnosis of epilepsy.
- Summarize evidence showing that epilepsy testing can be associated with positive health outcomes.

# The state of genetics today

It takes on average 4.8 years for a rare disease patient to receive an accurate diagnosis\*

- Cost barriers
- Reimbursement challenges
- Lack of comfort with genetics

\*Global Genes website: [https://globalgenes.org/rare-facts/?gclid=EAlalQobChMIupHBhrPr6AIV1BatBh0bCgH6EAAAYASAAEgJwzPD\\_BwE](https://globalgenes.org/rare-facts/?gclid=EAlalQobChMIupHBhrPr6AIV1BatBh0bCgH6EAAAYASAAEgJwzPD_BwE)



# Benefits of identifying a genetic etiology for patients with early-life epilepsy

- **High diagnostic yield<sup>1</sup>**
  - **Genetic testing of patients with early-life epilepsy has diagnostic yield of 30%–40%**
- **Direct medical care<sup>2,3</sup>**
  - **Prevent unnecessary studies/procedures**
  - **Surveillance for anticipated symptoms**
  - **Inform medications (indicated vs. contraindicated)**
- **Avoidance of a diagnostic odyssey<sup>3</sup>**
  - **Save money and time for family**
  - **Reduce overall healthcare costs**

<sup>1</sup>Berg AT, et al. *JAMA Pediatr.* 2017;171(9):863–71;

<sup>2</sup>Truty R, et al. *Epilepsia Open.* 2019;4(3):397–408.

<sup>3</sup>Oates S, et al. *NPJ Genom Med.* 2018; 3:13.

# Benefits of identifying a genetic etiology for patients with early-life epilepsy (cont.)

- **Family planning<sup>1</sup>**
  - Determine recurrence risk for parents
  - Determine risk for siblings
- **Family support**
  - Find advocacy/support groups centered around same rare disease<sup>1</sup>
  - Be part of a community of families with similar concerns
- **Clinical trials and research studies<sup>1</sup>**
  - Early access to therapies
  - Cutting-edge disease management
  - Natural history studies

→ We anticipate that all of these benefits may ultimately result in better outcomes for the patient.

<sup>1</sup>Berg AT, et al. *JAMA Pediatr.* 2017;171(9):863–71;

# Many studies demonstrate the clinical utility and costs savings of utilizing a gene panel for patients with epilepsy

Accepted: 3 April 2018

DOI: 10.1111/epi.14087

FULL-LENGTH ORIGINAL RESEARCH

Epilepsia

## A population-based cost-effectiveness study of early genetic testing in severe epilepsies of infancy

Katherine B. Howell<sup>1,2,3</sup> | Stefanie Eggers<sup>3</sup> | Kim Dalziel<sup>3,4</sup> | Jessica Riseley<sup>3</sup> |  
Simone Mandelstam<sup>2,3,5,6,7</sup> | Candace T. Myers<sup>8</sup> | Jacinta M. McMahon<sup>9</sup> |  
Amy Schneider<sup>9</sup> | Gemma L. Carvill<sup>10</sup> | Heather C. Mefford<sup>8</sup> | the Victorian Severe Epilepsy  
of Infancy Study Group | Ingrid E. Scheffer<sup>1,2,7,9</sup> | A. Simon Harvey<sup>1,2,3</sup>

ARTICLE

## Diagnostic yield of genetic tests in epilepsy

A meta-analysis and cost-effectiveness study

Iván Sánchez Fernández, MD, MPH, Tobias Loddenkemper, MD, Marina Gaínza-Lein, MD,  
Beth Rosen Sheidley, MS, CGC, and Annapurna Poduri, MD, MPH

*Neurology*® 2019;92:e1-e11. doi:10.1212/WNL.0000000000006850

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FULL-LENGTH ORIGINAL RESEARCH



## Diagnostic yield of genetic testing in epileptic encephalopathy in childhood

\*†Saadet Mercimek-Mahmutoglu, \*Jaina Patel, \*Dawn Cordeiro, \*Stacy Hewson, ‡David Callen,  
§Elizabeth J. Donner, §Cecil D. Hahn, \*†Peter Kannu, §Jeff Kobayashi, †§Berge A. Minassian,  
§Mahendranath Moharir, \*Komudi Siriwardena, §Shelly K. Weiss, \*†Rosanna Weksberg, and  
§O. Carter Snead III

*Epilepsia*, 56(5):707–716, 2015  
doi: 10.1111/epi.12954

ORIGINAL RESEARCH ARTICLE

Front. Neurol., 13 September 2019 | <https://doi.org/10.3389/fneur.2019.00988>



## Diagnostic Yield of Epilepsy Panel Testing in Patients With Seizure Onset Within the First Year of Life

✉ Se Song Jang<sup>1</sup>, ✉ Soo Yeon Kim<sup>1</sup>, ✉ Hunmin Kim<sup>2</sup>, ✉ Hee Hwang<sup>2</sup>, ✉ Jong Hee Chae<sup>1</sup>, ✉ Ki Joong Kim<sup>1</sup>, ✉ Jong-Il Kim<sup>3,4,5</sup> and ✉ Byung Chan Lim<sup>1</sup>

<sup>1</sup>Department of Pediatrics, Seoul National University College of Medicine, Seoul National University Children's Hospital, Seoul, South Korea

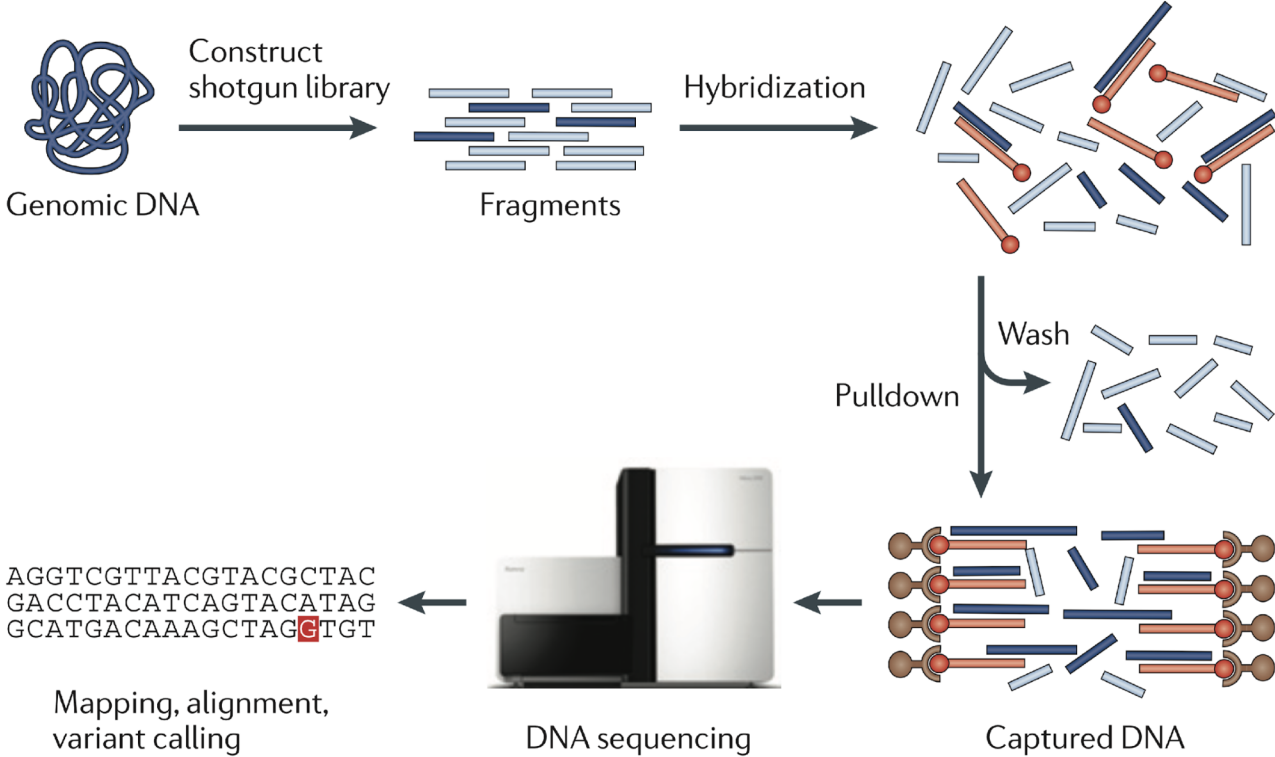
<sup>2</sup>Department of Pediatrics, Seoul National University College of Medicine, Seoul National University Bundang Hospital, Bundang-gu, South Korea

<sup>3</sup>Department of Biomedical Sciences, Seoul National University Graduate School, Seoul, South Korea

<sup>4</sup>Department of Biochemistry and Molecular Biology, Seoul National University College of Medicine, Seoul, South Korea

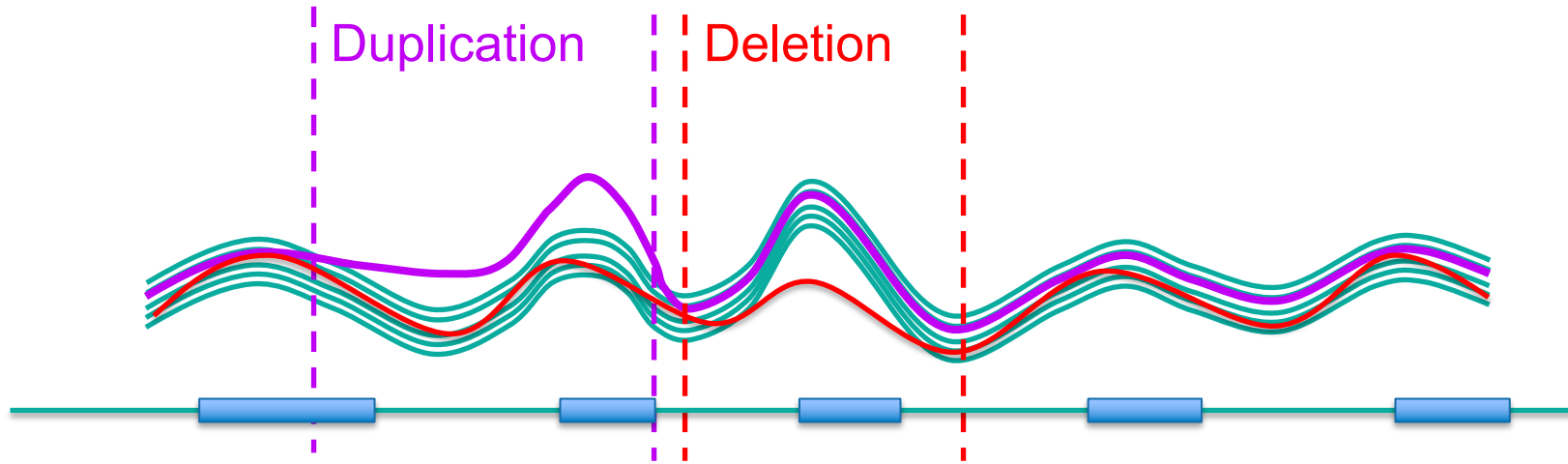
<sup>5</sup>Medical Research Center, Genomic Medicine Institute, Seoul National University, Seoul, South Korea

# Next Generation Sequencing



Source: Bamshad MJ, et al. *Nat Rev Genet.* 2011;12(11):745–55.

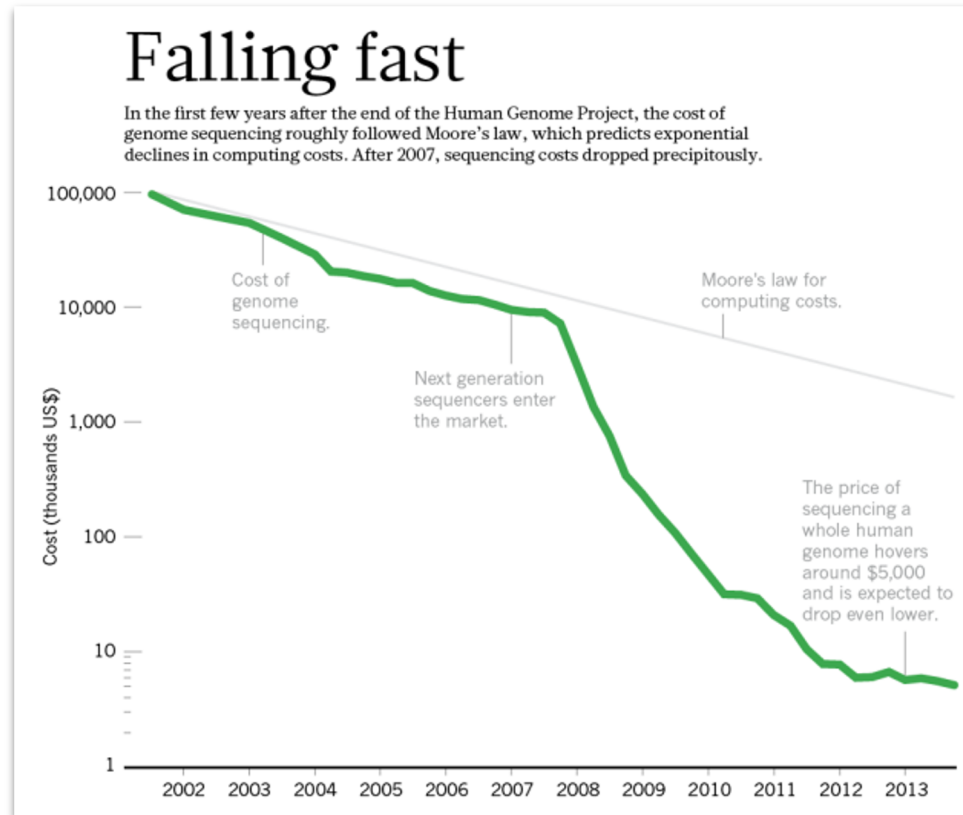
# Copy number detection by next-generation sequencing



- The depth profile is non-uniform but reproducible.
- Look for deviations with respect to *baseline samples*.
- Perform this evaluation at the assay level to be able to detect deletions/duplications down to single-exon resolution across the panel.



# Cost of DNA sequencing dropping rapidly



# Evolution of large public databases

## 1000 Genome (2010)

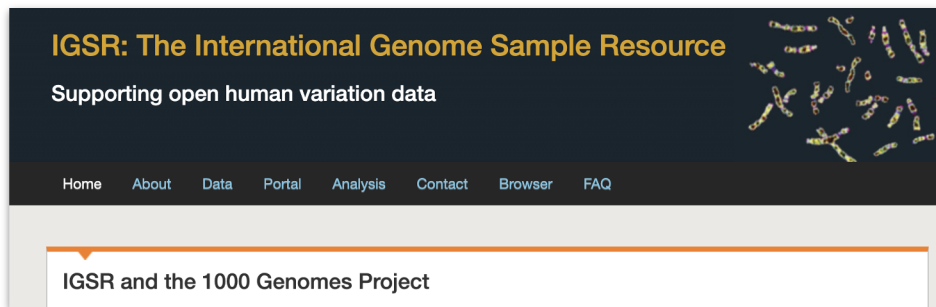
First large public database of genome data from “control” individual

## ExAC (2014)

Second large public database  
60,000 exomes

## gnomAD (2017)

Current largest public database  
v2: 125,748 exomes and 15,708 whole genomes  
v3: 71,702 whole genomes



**IGSR: The International Genome Sample Resource**  
Supporting open human variation data

Home About Data Portal Analysis Contact Browser FAQ

IGSR and the 1000 Genomes Project



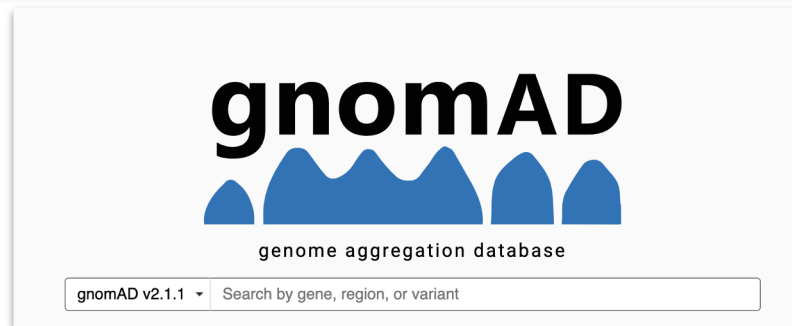
NATURE | EDITORIAL

عربي

### ExAC project pins down rare gene variants

Catalogue of genetic information from some 60,000 people reveals unexpected surprises — and highlights the need to make genomic data publicly accessible to aid studies of rare diseases.

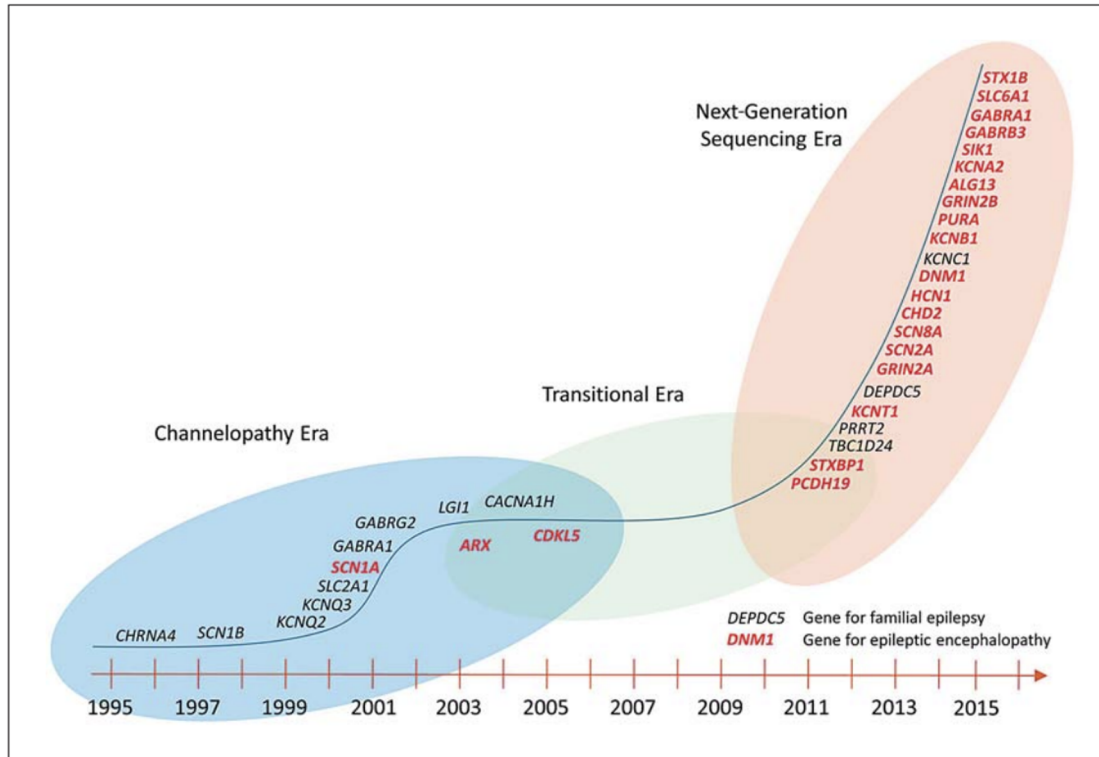
17 August 2016



**gnomAD**  
genome aggregation database

gnomAD v2.1.1 Search by gene, region, or variant

# Gene discovery for epilepsy



**Review Article**

**Molecular  
Syndromology**

Mol Syndromol 2016;7:172-181  
DOI: 10.1159/000448530

Published online: August 20, 2016

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**Understanding Genotypes and  
Phenotypes in Epileptic Encephalopathies**

Ingo Helbig<sup>a, c</sup> Ahmad N. Abou Tayoun<sup>b</sup>

Divisions of <sup>a</sup>Neurology and <sup>b</sup>Genomic Diagnostics, Department of Pathology and Laboratory Medicine, The Children's Hospital of Philadelphia, University of Pennsylvania Perelman School of Medicine, Philadelphia, Pa., USA; <sup>c</sup>Department of Neuropediatrics, Christian Albrechts University of Kiel and University Medical Center Schleswig-Holstein (UKSH), Kiel, Germany

Helbig I, Tayoun AA. Understanding Genotypes and Phenotypes in Epileptic Encephalopathies. *Mol Syndromol*. 2016;7(4):172-181.

# Study of genetic testing utility in epilepsy

FULL-LENGTH ORIGINAL RESEARCH

Epilepsia Open®

Open Access

## **Possible precision medicine implications from genetic testing using combined detection of sequence and intragenic copy number variants in a large cohort with childhood epilepsy**

Rebecca Truty<sup>1</sup> | Nila Patil<sup>2</sup> | Raman Sankar<sup>2</sup> | Joseph Sullivan<sup>3</sup> | John Millichap<sup>4</sup>  |  
Gemma Carvill<sup>5</sup> | Ali Entezam<sup>1</sup> | Edward D. Esplin<sup>1</sup> | Amy Fuller<sup>1</sup> | Michelle Hogue<sup>1</sup> |  
Britt Johnson<sup>1</sup> | Amirah Khouzam<sup>1</sup> | Yuya Kobayashi<sup>1</sup> | Rachel Lewis<sup>1</sup> |  
Keith Nykamp<sup>1</sup> | Darlene Riethmaier<sup>1</sup> | Jody Westbrook<sup>1</sup> | Michelle Zeman<sup>1</sup> |  
Robert L. Nussbaum<sup>1,6</sup> | Swaroop Aradhya<sup>1</sup> 

# Methods

- NGS panel consisting of up to **183 genes**
- Simultaneous detection of sequence variants and exon-level copy number variants (deletions and duplications)
- **~9769 patients** were referred for testing for all genes or a subset (e.g., panels for early infantile epileptic encephalopathy or Rett/Angelman spectrum)
  - Unselected cohort
  - ~90% where under the age of 18 at time of testing
- Definition of **precision medicine implications** based on published literature and curation by expert clinicians with long-standing experience in epilepsy

# Epilepsy genes related to precision medicine

Strong		Emerging	
<b>ALDH7A1</b>	Pyridoxine-dependent epilepsy	<b>ALDH5A1</b>	Succinic semialdehyde dehydrogenase deficiency
<b>CSTB</b>	Unverricht and Lundborg PME	<b>ATP1A3</b>	Alternating hemiplegia of childhood
<b>EPM2A</b>	Lafora disease	<b>DEPDC5</b>	Familial focal epilepsy
<b>FOLR1</b>	Cerebral folate transport deficiency	<b>GLRA1</b>	Hyperekplexia
<b>GAMT</b>	Cerebral creatine deficiency syndrome	<b>GNAO1</b>	Early infantile epileptic encephalopathy
<b>GATM</b>	Cerebral creatine deficiency syndrome	<b>GOSR2</b>	Progressive myoclonic epilepsy
<b>KCNQ2</b>	Early infantile epileptic encephalopathy	<b>GRIN1</b>	Epilepsy with ID and hyperkinesia
<b>NHLRC1</b>	Lafora disease	<b>GRIN2A</b>	Focal epilepsy, speech impairment & ID
<b>PNPO</b>	Pyridoxamine 5'-PO <sub>4</sub> oxidase deficiency	<b>GRIN2B</b>	EIEE & ID
<b>POLG</b>	Mitochondrial DNA depletion; PEO	<b>KCNQ3</b>	Benign infantile epilepsy, developmental disability
<b>SCN1A</b>	Dravet syndrome	<b>KCNT1</b>	Early infantile epileptic encephalopathy
<b>SCN2A</b>	Early infantile epileptic encephalopathy	<b>NGLY1</b>	Congenital disorder of deglycosylation
<b>SCN8A</b>	Early infantile epileptic encephalopathy	<b>PCDH19</b>	Sex-limited EIEE
<b>SLC2A1</b>	Glucose transporter deficiency	<b>PIGA</b>	Congenital anomalies-hypotonia-seizures syndrome
<b>SLC6A8</b>	Cerebral creatine deficiency syndrome	<b>PRRT2</b>	Episodic kinesigenic dyskinesia & seizures
<b>TPP1</b>	Neuronal ceroid lipofuscinosis 2 (CLN2)	<b>QARS</b>	Progressive microcephaly with seizures
<b>TSC1</b>	Tuberous sclerosis	<b>SLC6A1</b>	Myoclonic-atonic epilepsy
<b>TSC2</b>	Tuberous sclerosis		

Legend:

Genes associated with biochemical disorders

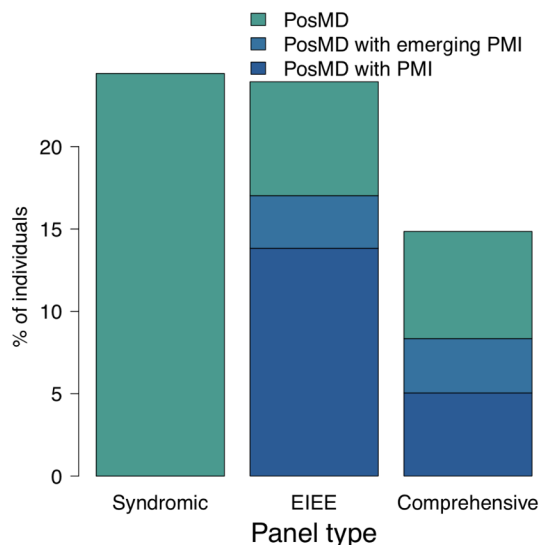
Genes associated with disorders for which there are anti-epileptic drug contraindications

Genes associated with disorders for which there are indications for using specific anti-epileptic drugs.

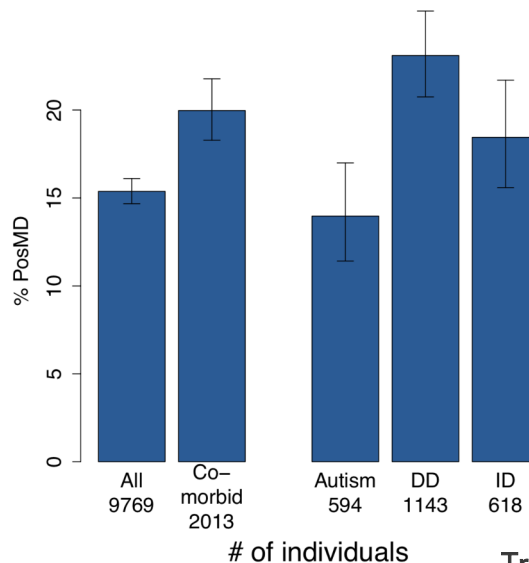
# Diagnostic yield of genetic testing in epilepsy

- Diagnostic **yield of 15-25%**, depending on panel used
- Syndromic and EIEE have higher yield compared to Comprehensive panel
- **33% of individuals** with positive reports had results with **precision medicine implications**

PosMD yield by panel



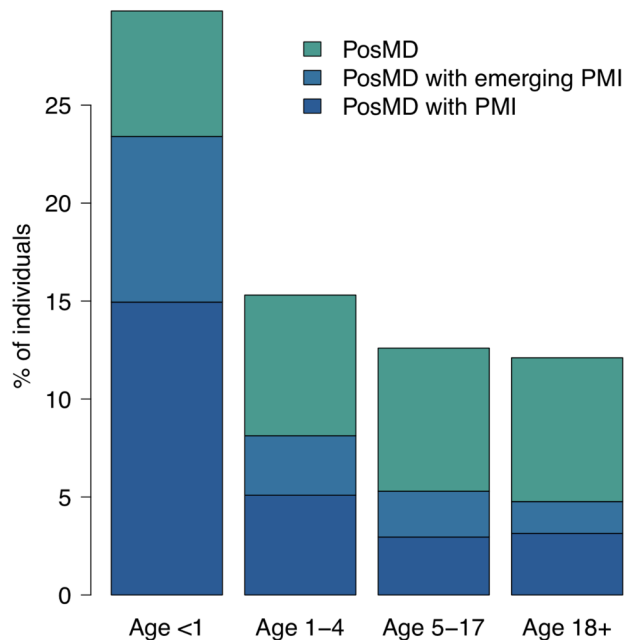
PosMD yield by co-morbidity



# Early molecular diagnosis is important in epilepsy

- Children with epilepsy in **first year of life** have the **highest diagnostic yield, at 30%**
- Nearly **half** of those with seizures in their first year of life and with positive reports have results with **precision medicine implications**
- 66% of all positive results with precision medicine implications were in children **younger than 5 years**

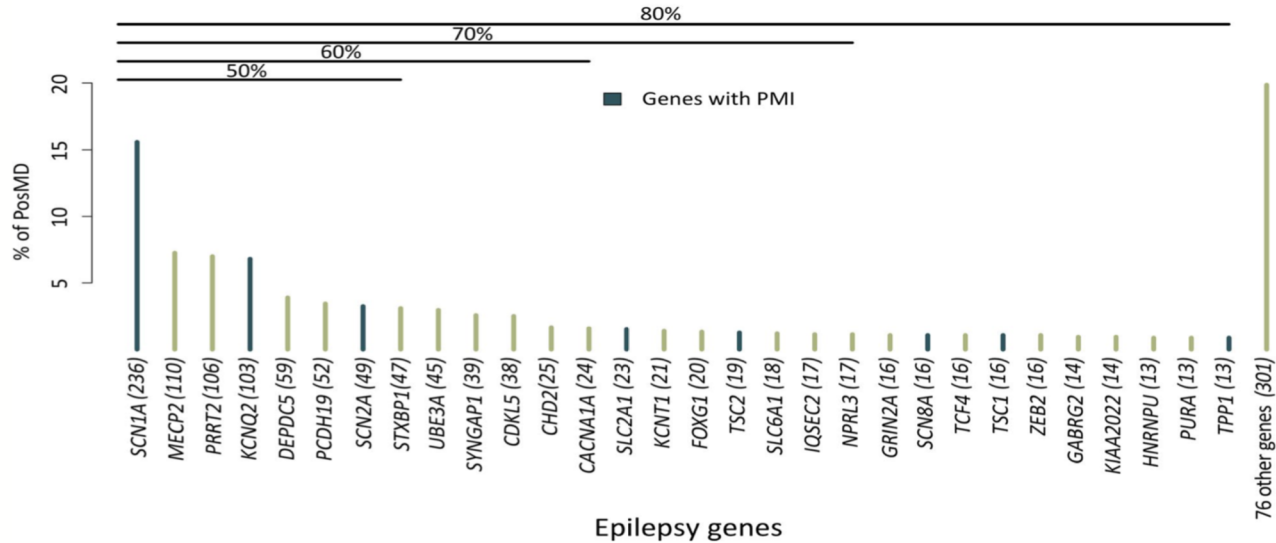
PosMD by age at testing





# Critical genes involved in epilepsy

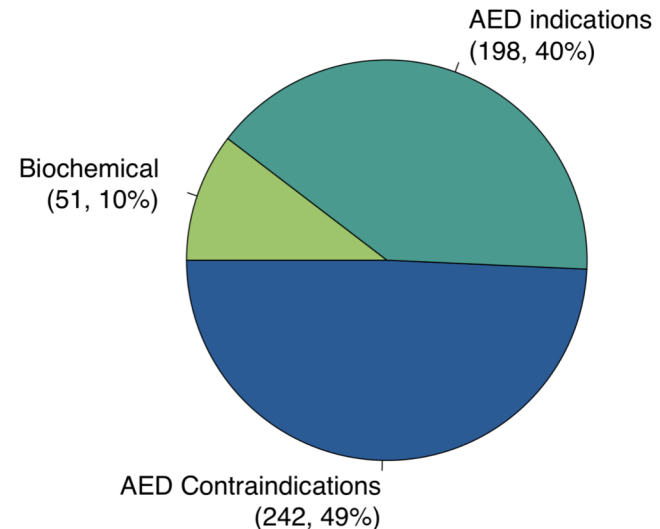
- **80% of positive molecular diagnoses came from only 30 genes** while the remaining 20% were from 76 other genes.
- *SCN1A* has the highest yield
- Corroborated by another large genetic testing study (Lindy et al., *Epilepsia* 2018)



# Categories of precision medicine implications

- Nearly **half of positive molecular diagnoses** with precision medicine implications were related to **contraindications** for anti-epileptic drugs (AED), largely due to variants in *SCN1A*.
- **10%** of molecular diagnoses were related to **biochemical disorders** with available treatments
- **40%** of molecular diagnoses invoked **indications** for specific AED (e.g., Vigabatrin for spasms in TSC)
- Another **21%** of individuals had positive molecular diagnoses in genes with **emerging associations** with precision medicine implications

PosMD with PMI



# Study of genetic testing utility for epilepsy in adults

ARTICLE

OPEN ACCESS

## Multigene Panel Testing in a Large Cohort of Adults With Epilepsy

Diagnostic Yield and Clinically Actionable Genetic Findings

Dianalee McKnight, PhD, Sara L. Bristow, PhD, Rebecca M. Truty, PhD, Ana Morales, MS, Molly Stetler, MS, M. Jody Westbrook, PhD, Kristina Robinson, PhD, Darlene Riethmaier, MS, Felipe Borlot, MD, Marissa Kellogg, MD, Sean T. Hwang, MD, Anne Berg, PhD, and Swaroop Aradhya, PhD

**Correspondence**

Dr. McKnight  
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*Neurol Genet* 2022;8:e650. doi:10.1212/NXG.0000000000000650

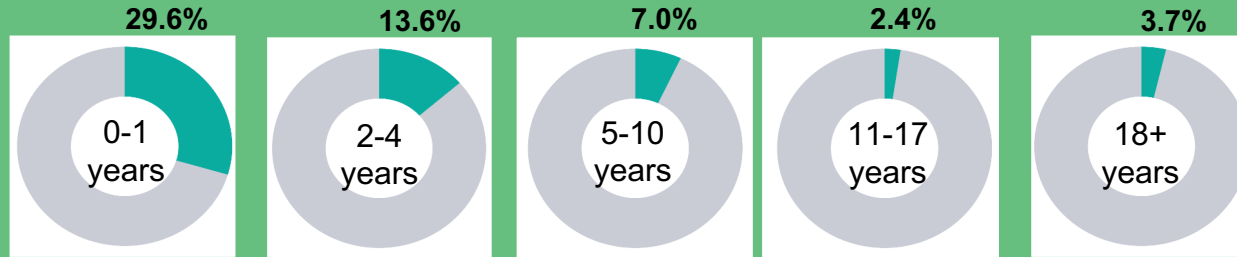
# Overall diagnostic yields for genetic testing in adults with epilepsy

2,008 individuals over 18 years of age were tested with multi-gene panels



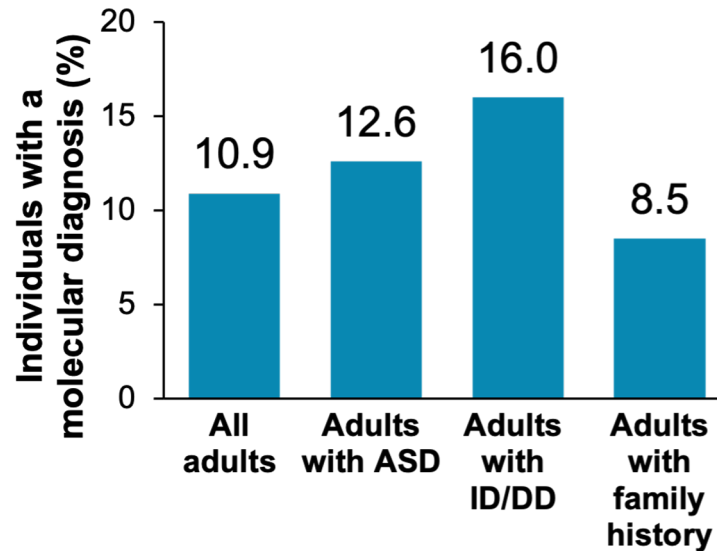
One in ten adults with epilepsy received a definitive molecular diagnosis

Seizure onset in infancy resulted in the highest diagnostic yield



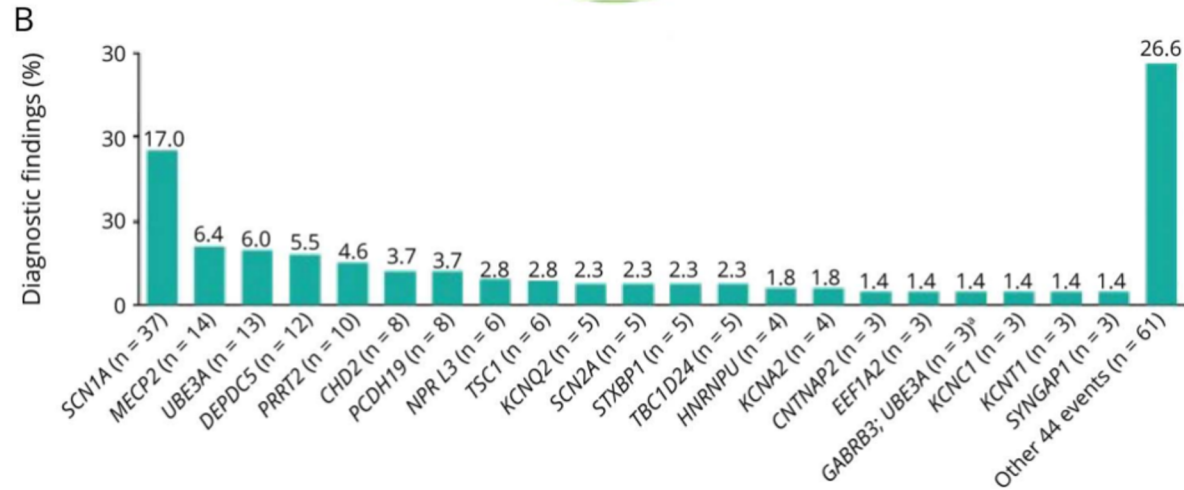
# Adults with epilepsy and comorbidities

Among investigated comorbidities, **adults with reported ID/DD** were most likely to have a molecular diagnosis (16.0%).



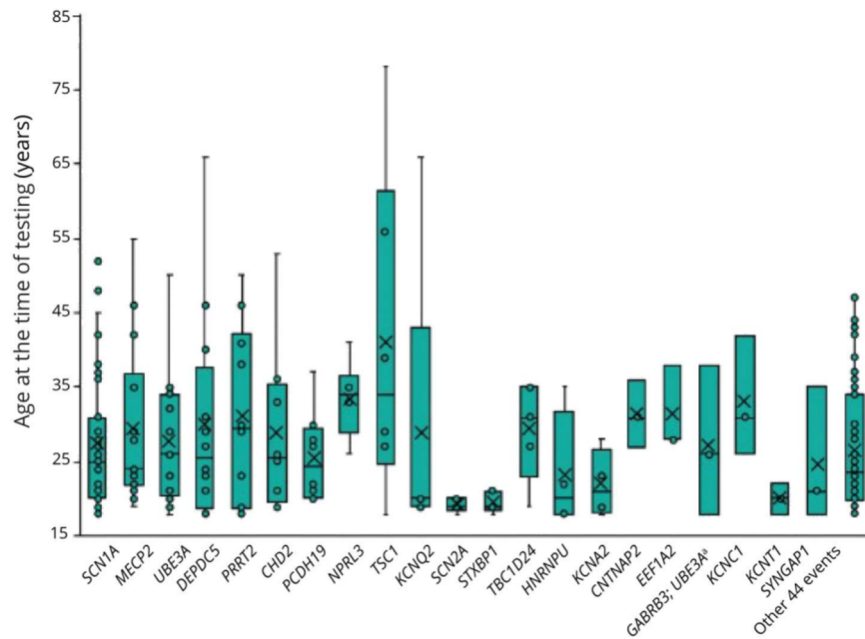
# Critical genes involved in epilepsy

- Just like previous studies, most findings are in under 30 genes
- *SCN1A* has the highest yield
- 11 of the 13 top genes in both the Truty et al. study (90% kids) and this study (100% adults) were the same

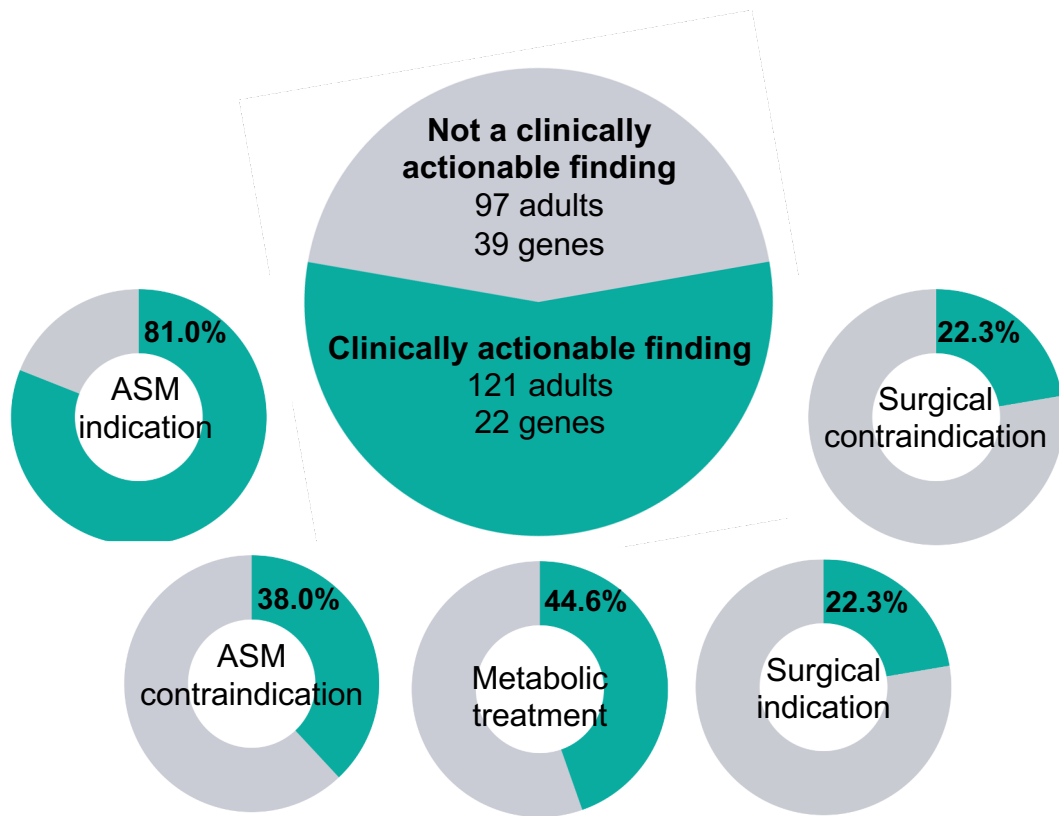


# Many adults with genetic epilepsy live well into adulthood

Many adults in their **50s, 60s, 70s and even 80s** were found to have genetic epilepsy



# Results with clinically actionable findings



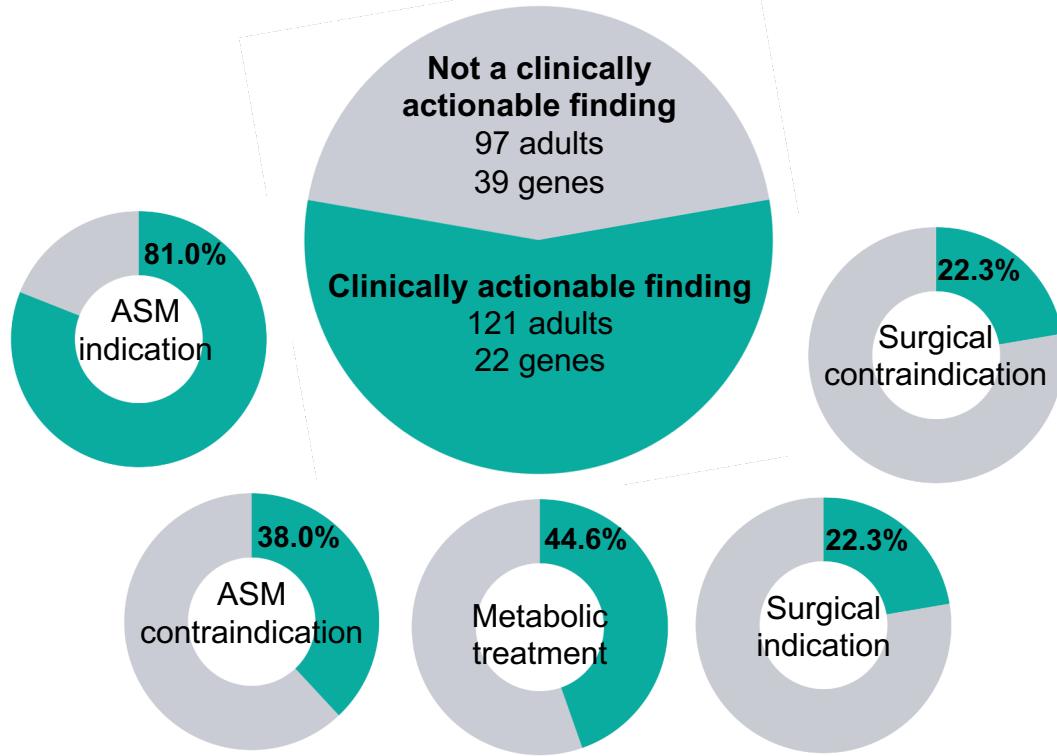
- **Over one-half** of adults with epilepsy had diagnostic findings in genes associated with at least one **specific treatment for seizure control**
- Genetic testing could have a direct impact on clinical management for many adult patients with epilepsy



**Table 3.** Genes associated with epilepsy that have clinically actionable implications

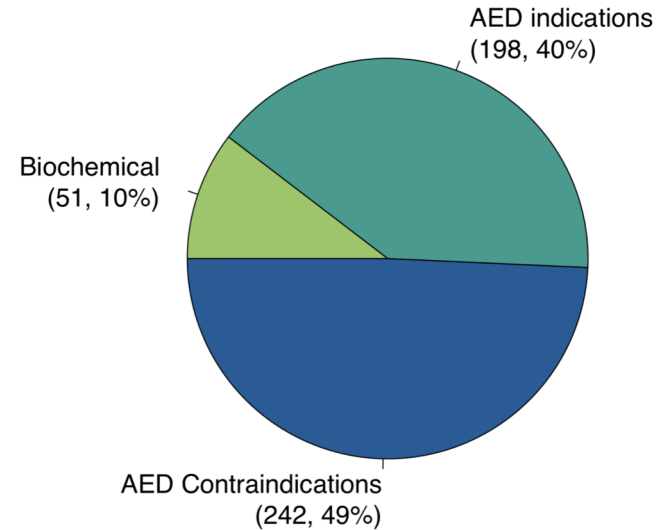
Gene	Associated syndrome or disorder	Inheritance	Clinical action	Potential clinical management options for daily maintenance of seizures*	Potential clinical management options for other manifestations	References (PMIDs)						
<i>ALDH7A1</i>	Pyridoxine-dependent epilepsy; folinic acid-responsive seizures	AR				Indicated: high resolution neuroimaging; epilepsy surgery for identifiable epileptogenic foci						
<i>ATPIA3</i>	Alternating hemiplegia of childhood type 2; dystonia 12; cerebellar ataxia, areflexia, pes cavus, optic atrophy and sensorineural hearing loss	AD	<i>DEPDC5</i>	Familial focal epilepsy with variable foci; autosomal dominant nocturnal frontal lobe epilepsy	AD	Surgical indications	26434565, 30782578, 27683934, 30093711					
<i>CACNA1A</i>	Developmental and epileptic encephalopathy; episodic ataxia type 2 (EA2); familial hemiplegic migraine type 1 (FHM1)	AD	<i>EPM2A</i>	Progressive myoclonus epilepsy, Lafora type	AR	ASM indicated ASM contraindicated	Indicated: VPA and BZDs as first-line;					
<i>CDKL5</i>	Developmental and epileptic encephalopathy; atypical Rett syndrome; Angelman-like syndrome	XLD	<i>GLRA1</i>	Hyperekplexia 1	AD/AR	ASM indicated	Autosomal dominant nocturnal frontal lobe epilepsy; developmental and epileptic encephalopathy	AD	ASM indications; Metabolic treatment	Indicated: quinidine, KD	26369628, 32167590, 31054119	
<i>CHD2</i>	Childhood-onset epileptic encephalopathy	AD	<i>KCNH2</i>	Long QT syndrome type 2; short QT syndrome	AD	Other	<i>KCNT1</i>	Autosomal dominant lateral temporal lobe epilepsy	AD	ASM indications	Indicated: PHT, CBZ, VPA	20301709, 7647791
			<i>KCNQ2</i>	Benign familial neonatal seizures; developmental and epileptic encephalopathy	AD	ASM indicated (subject to GC consideration)	<i>LGII</i>			Indicated: VPA and BZDs as first-line; LEV, ZNS, TPM, and PER as second-line; primidone, PB, piracetam, and ESM as third-line		
							<i>NHLRC1</i>	Progressive myoclonus epilepsy, Lafora type	AR	ASM indications; ASM contraindications	Contraindicated: PHT, LTG, CBZ, OXC	20301563, 25667898
							<i>NPRL3</i>	Familial focal epilepsy with variable foci	AD	Surgical intervention	Indicated: high resolution neuroimaging; epilepsy surgery for identifiable epileptogenic foci	26434565, 26285051
							<i>PCDH19</i>	Developmental and epileptic encephalopathy	XLD	ASM indications; Metabolic treatment	Indicated: clobazam, potassium bromide, PHT, KD	26820223, 23712037
							<i>PRRT2</i>	Episodic kinesigenic dyskinesia; benign familial infantile seizure 2; familial infantile convulsions with paroxysmal choreoathetosis	AD	ASM indications	Indicated: OXC, CBZ	28056630, 29334453, 32392383

# Two studies reported ~50% of positive results should be clinically actionable.....but are they in practice?



McKnight et al. Neurol Genet. 2022;8:e650

## PosMD with PMI



Truty et al. Epilepsia Open. Jul 2019



## **Molecular Diagnosis from Genetic Testing Guides Clinical Management of Epilepsy and Helps Improve Patient Outcomes**

**Dianalee McKnight, PhD, FACMG  
Invitae**

**December 6, 2021**



### **Collaborators:**

Ana Morales (Invitae)  
Kathryn E. Hatchell (Invitae)  
Sara Bristow (Invitae)  
Edward D. Esplin (Invitae)  
Chad Moretz (Invitae)  
Robert L. Nussbaum (Invitae)  
Swaroop Aradhya (Invitae)  
Felippe Borlot (University of Calgary)  
Kaitlin Angione (Children's Hospital Colorado)  
M. Scott Perry (Cook Children's Medical Center)  
Joshua Bonkowsky (University of Utah)  
Loreto Ríos-Pohl (Clínica Integral de Epilepsia)  
Anne Berg (Northwestern-Feinberg School of Medicine)  
**ELEVIATE consortium**

**75<sup>TH</sup>  
ANNUAL  
MEETING**

# Epilepsy Outcomes Study Objectives

- Cite data demonstrating that epilepsy genetic testing can lead to actionable results.
- List examples of patient management changes that are made by clinicians based on a molecular diagnosis of epilepsy.
- Discuss evidence showing that epilepsy testing can be associated with positive health outcomes.

# Rationale

- Genetic testing for epilepsy identifies a molecular etiology in up to 40% of cases
- ~50% of positive findings are in clinically actionable genes <sup>1 2</sup>
- Previous studies have demonstrated reduced healthcare costs for individuals with good versus poor seizure control (~\$14,000/year vs. ~\$23,000/year, respectively) due to decreased hospitalizations and emergency department visits <sup>3</sup>
- Limited information on how genetic information is used by clinicians to guide management and subsequent patient outcomes

→ **AIM:** We investigated changes in clinical management and patient outcomes after a definitive genetic diagnosis of epilepsy was identified

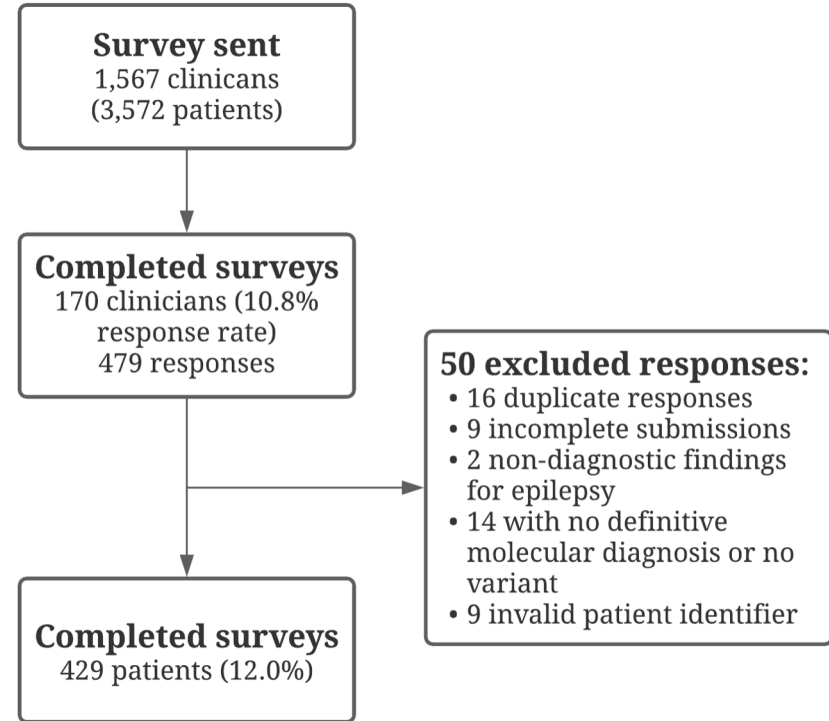
1. Truty R, et al. *Epilepsia Open*. 2019; 4(3):397–408)

2. McKnight et al. *Neurol Genet*. 2022;8:e650

3.. Cramer, Joyce A., et al. *Epilepsy & Behavior: E&B* 31 (February): 356–62.

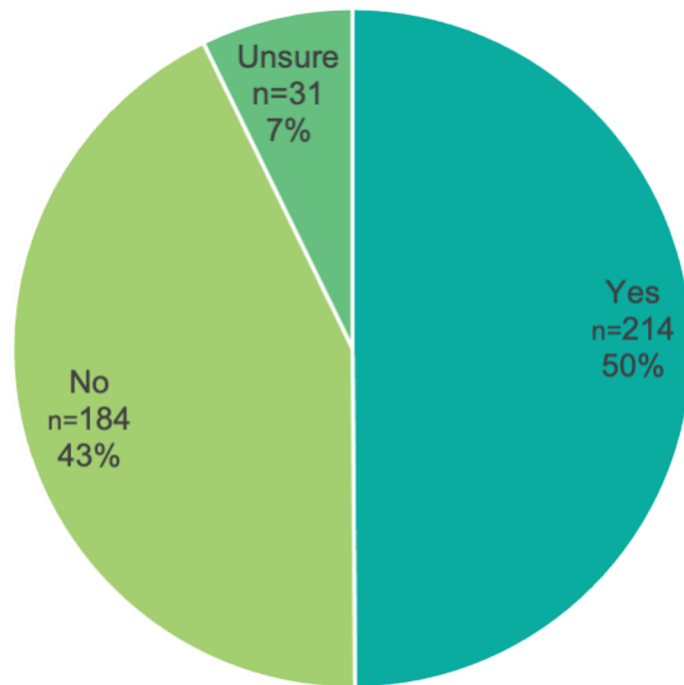
# Methods

- 1,567 clinicians who received a positive diagnostic finding for a patient with epilepsy were sent an invitation to participate
- Case report forms were collected between May-November 2020
- 170 clinicians completed case report forms for 429 patients with epilepsy and a genetic diagnosis
- Clinical specialties included:
  - Genetics (22.4%)
  - Pediatric neurology (17.1%)
  - Neurology (15.3%)
  - Epilepsy (7.6%)
  - Internal medicine (0.6%)
  - Multiple specialties (37.1%)



# Did the genetic finding influence a change in clinical management?

- **Half** of positive genetic diagnoses **led to** a change in clinical management.
- In **81.3%** of cases, providers changed clinical management **within 3 months** of receiving the genetic testing result.



# Changes that were implemented due to genetic finding

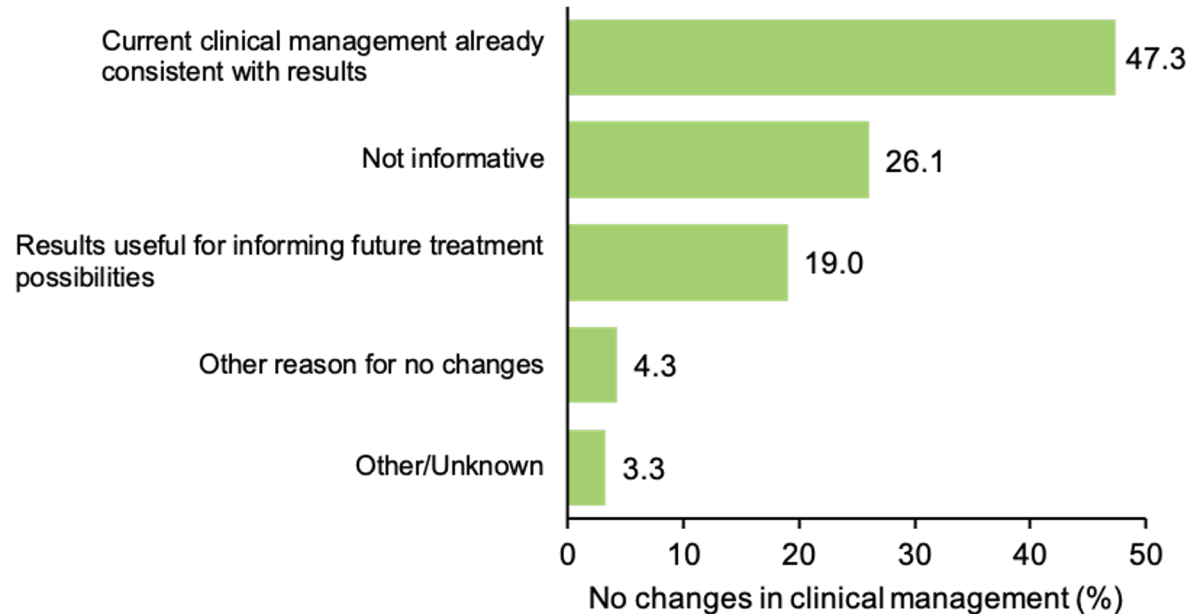
Most common change was to add a new medication.





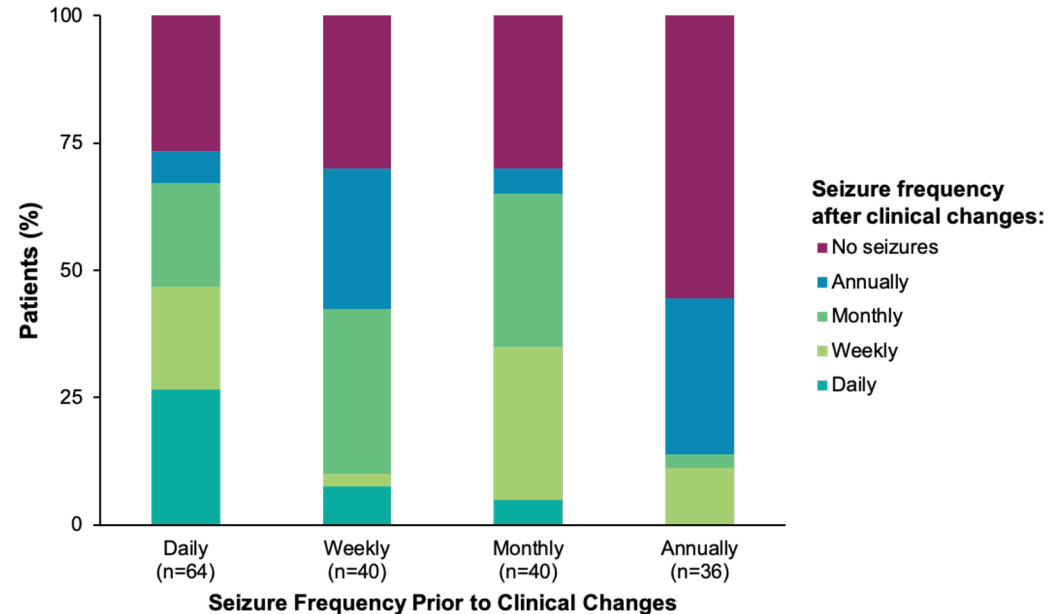
# Reasons for not changing management due to the genetic finding

Most common response was that clinical management was already consistent with results.



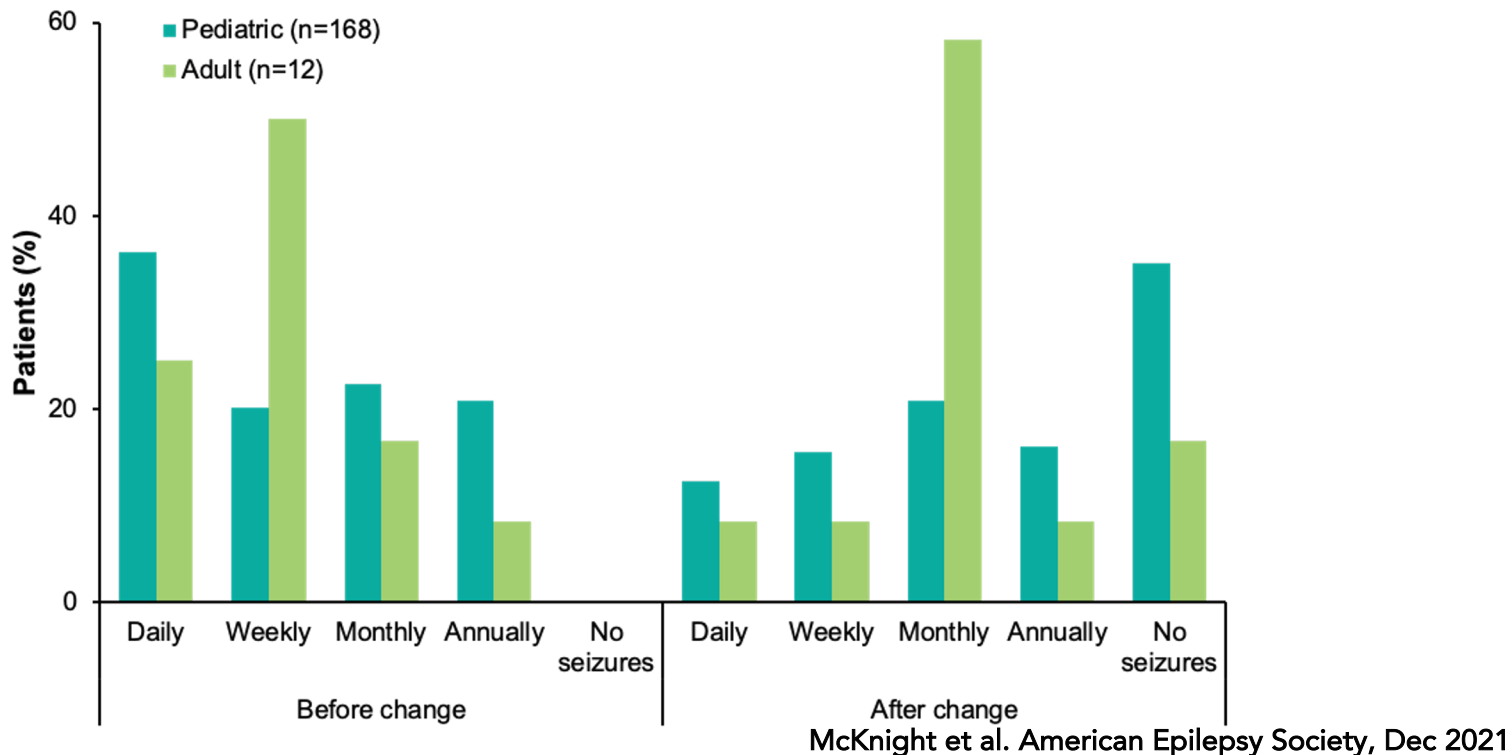
# Patient outcomes after treatment changes due to a molecular finding

- 75% reported positive outcomes (129/172)
- 65% reported reduced or elimination of seizures
- 23% reported other improvements (i.e. in behavior, development, academics, movement issues)
- 6% reported reduced medication side effects



# Positive outcomes observed in both adults and children

Similar trends in increased seizure control after treatment changes due to a diagnostic finding



# Conclusions

- **Genetic testing** for epilepsy identifies a genetic diagnosis in **up to 40%** of cases depending on many factors including age of seizure onset and presence of comorbidities.
- **~50%** of positive findings are in **clinically actionable genes**.<sup>1,2</sup>
- A genetic diagnosis appears to inform **changes in clinical management**.<sup>3</sup>
- Management changes informed by genetic information contributes to **improved seizure control** in **65% of cases** and other positive outcomes for many patients.<sup>3</sup>
- These results support growing evidence that genetic testing can **improve health outcomes**, which could also **reduce healthcare costs**.

1.Truty et al. Epilepsia Open. Jul 2019

2.McKnight et al. Neurol Genet. 2022;8:e650

3.McKnight et al. American Epilepsy Society, Dec 2021

Thank you

