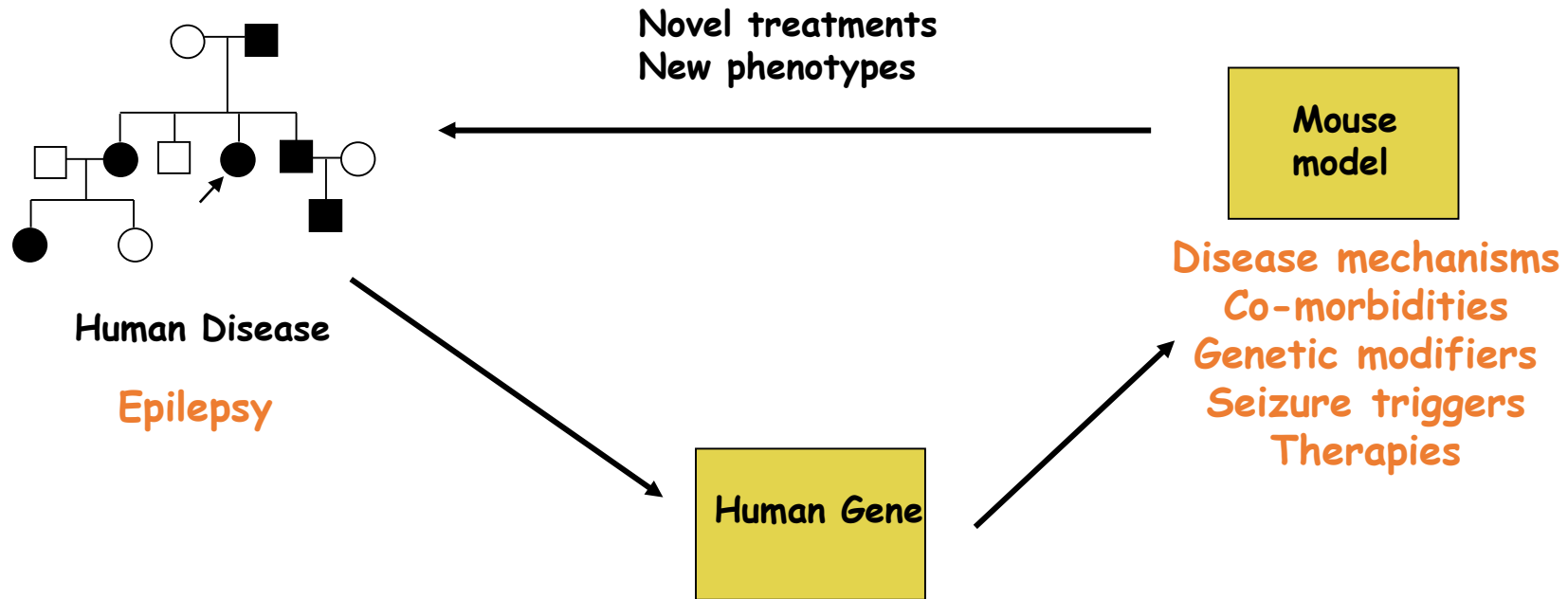


Mechanisms of *SLC6A1* dysfunction

SLC6A1 Connect Roundtable
November 29th, 2018

Andrew Escayg, Ph.D.
Dept. of Human Genetics
Emory University

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EGL Genetic
Molecular Genetics

PATIENT INFORMATION
Patient Name: Egl Genetics
DOB or Birth: 3/15/2017
Person Sex: Female
Case Reference #: 133-434
Project ID: T330345

Reviewed By: **MS, MD - Scott**
Q23.1 - Cerebellum
Reviewed Date/Time: 3/15/17

SUMMARY
Variant(s) detected:
PTPN11 NM_00204.3 c.2092C>T

RESULTS AND INTERPRETATION
Sequence analysis of the coding s
Gene: **PTPN11** MIM: **116275** Disease (link)
PTPN11 116275 Noonan syndrome
LEOPARD syndrome
Mitochondrial

The detection of a PTPN11 pathogenic variant in this individual, however, does not result in a diagnosis of Noonan syndrome.

PTPN11 NM_00204.3 c.2092C>T
The variant c.2092C>T is a missense variant for PTPN11.

Ref:
1. Chikama et al., Eur J Hum Genet
2. Chikama et al., Eur J Hum Genet
3. Schmitt et al., Hormones
4. Aki et al., Indian J Pediatr

RECOMMENDATIONS
• These results must be interpreted in the context of the clinical history and physical examination.
• Genetic counseling is recommended.
• EGL Genetics offers targeted testing for this variant.

EGL GENETICS
eglgenetics.com

Most referrals are children with severe epilepsy



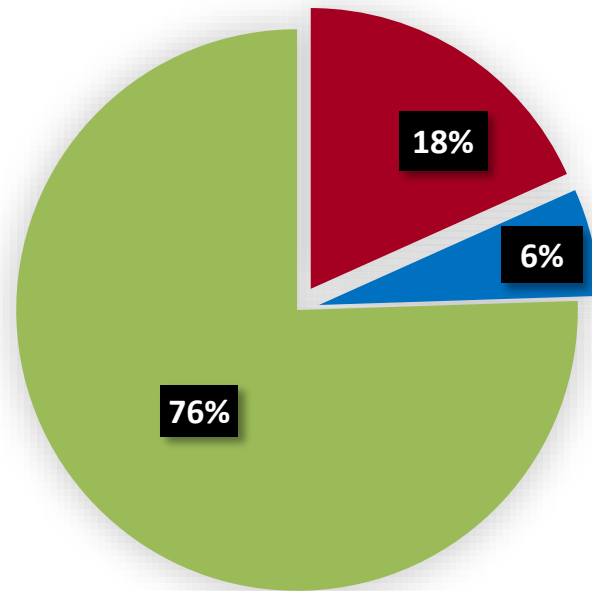
Demographics	
Average Age	7.53 yrs
Median Age	5.6 yrs
Maximum	74 yrs
Minimum	0.2 yrs

Phenotype listed	%
Developmental delay	31.2
Movement disorders	14.4
Hypotonia	13.0
Epileptic encephalopathy	12.6
Intellectual disability	7.4
Autism	5.6
Developmental regression	4.7
Microcephaly	4.7
Family History	4.2

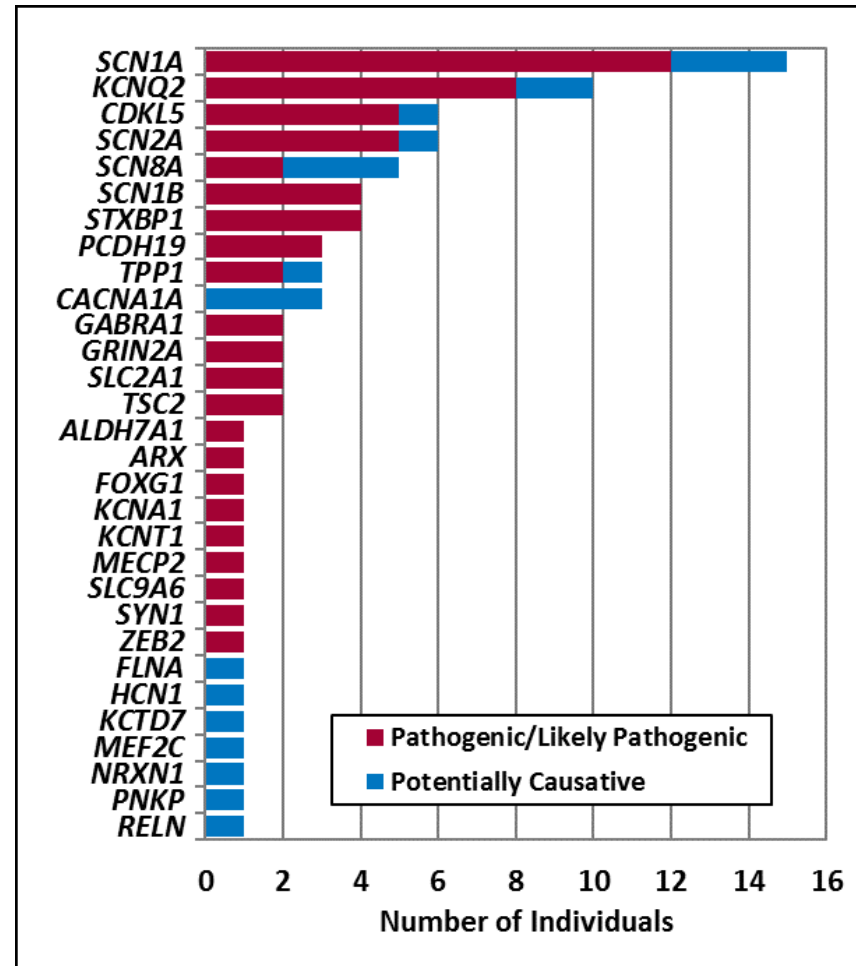
Diagnostic yield of the ESD panel

339 consecutive patients screened at EGL on the epilepsy panel

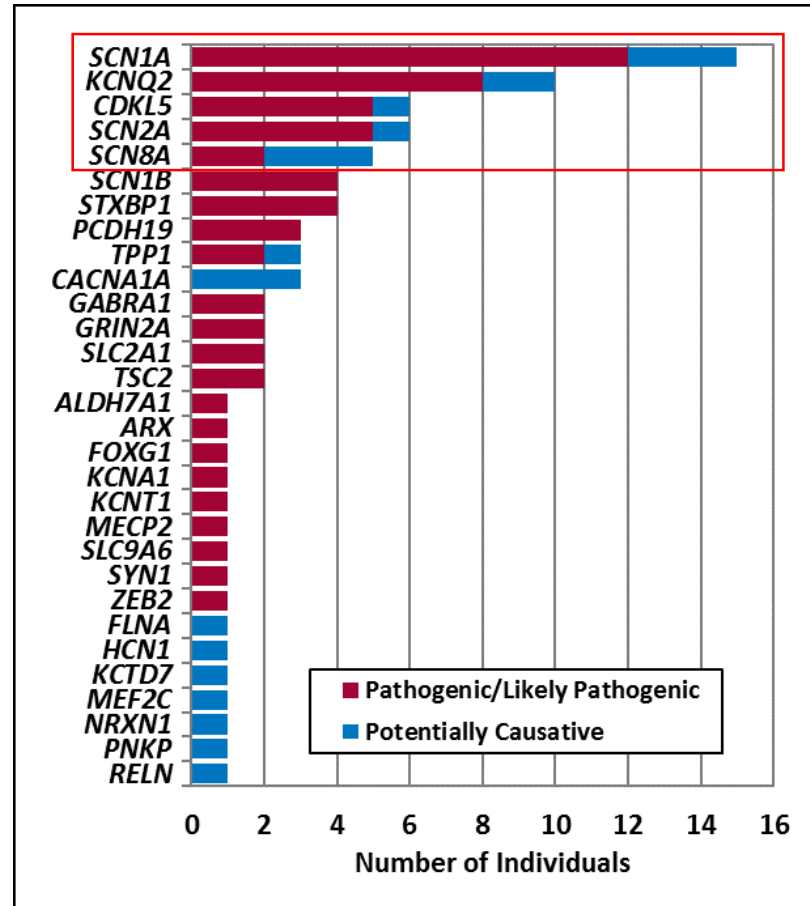
Variant Classification



- Pathogenic Variant
- Potentially Causative VUS
- VUS or Benign

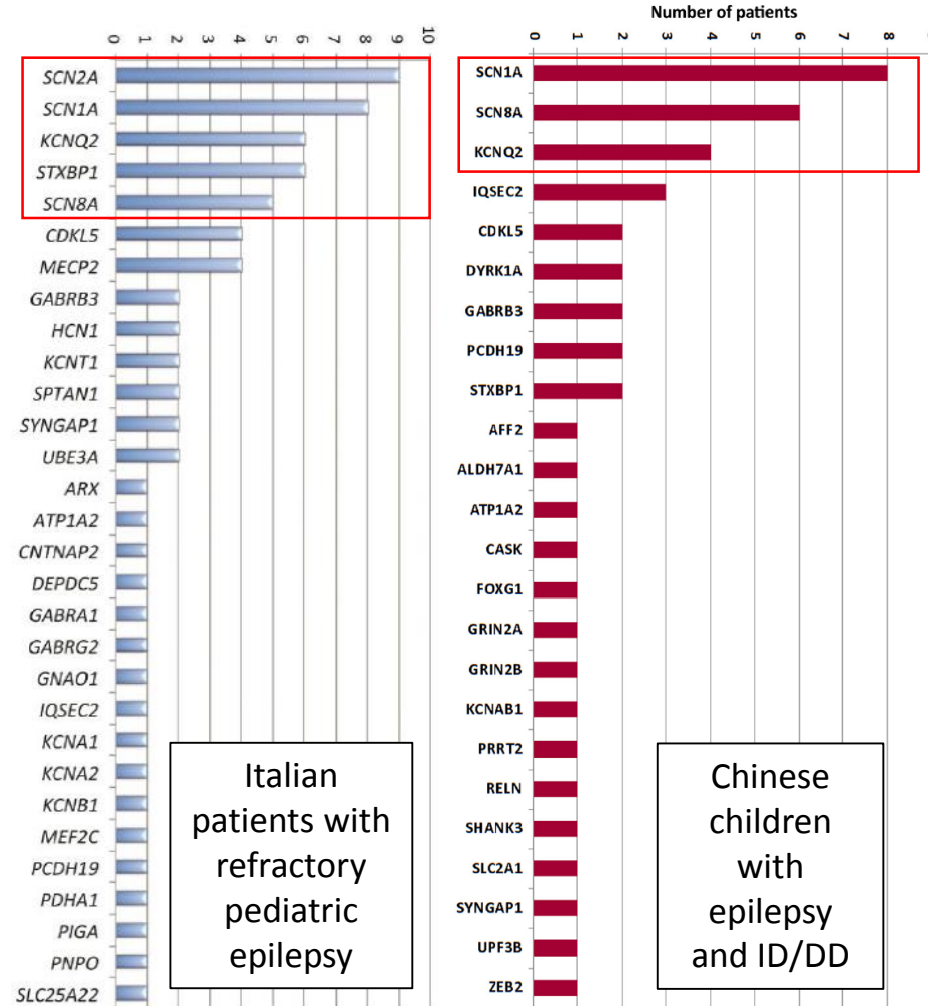


Comparable diagnostic yield of the ESD panel



Butler et al. (2017) *Ped. Neurol.*

N = 339



Parrini et al. (2017) *Hum Mutat.*

N = 349

Zhang et al. (2015) *PLoSOne*

N = 253

BRIEF COMMUNICATION***SLC6A1* variants identified in epilepsy patients reduce γ -aminobutyric acid transport****8 mutations from 460 referrals**

**Kari A. Mattison^{1,2} | Kameryn M. Butler^{1,2} | George Andrew S. Inglis^{1,2} |
Oshrat Dayan³ | Hanna Boussidan³ | Vikas Bhambhani⁴ | Bryan Philbrook⁵ |
Cristina da Silva⁶ | John J. Alexander^{1,6} | Baruch I. Kanner³ | Andrew Escayg^{1,2}**

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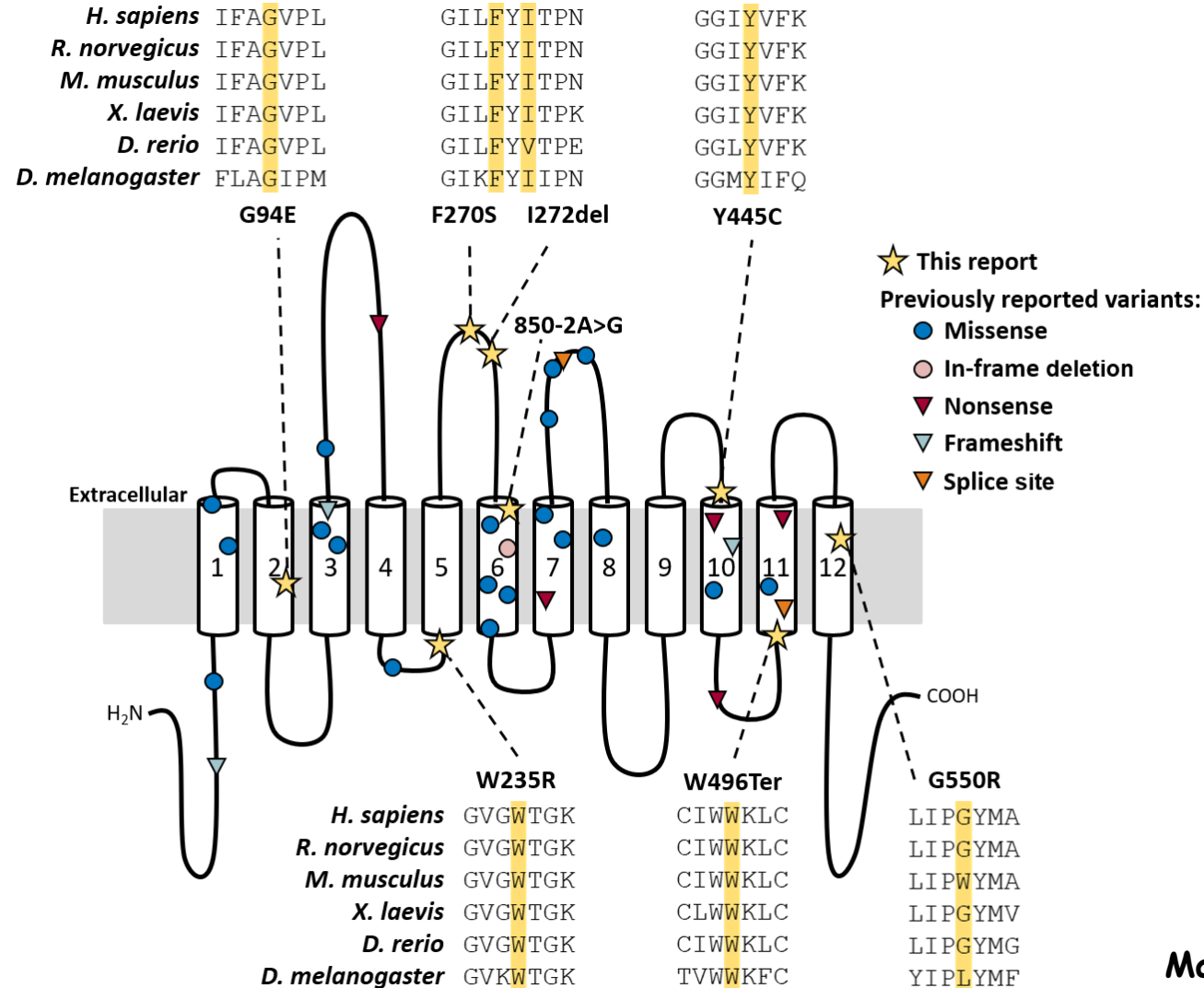
⁵Department of Pediatric Neurology, Emory University, Atlanta, Georgia

⁶EGL Genetics, Tucker, Georgia

Summary

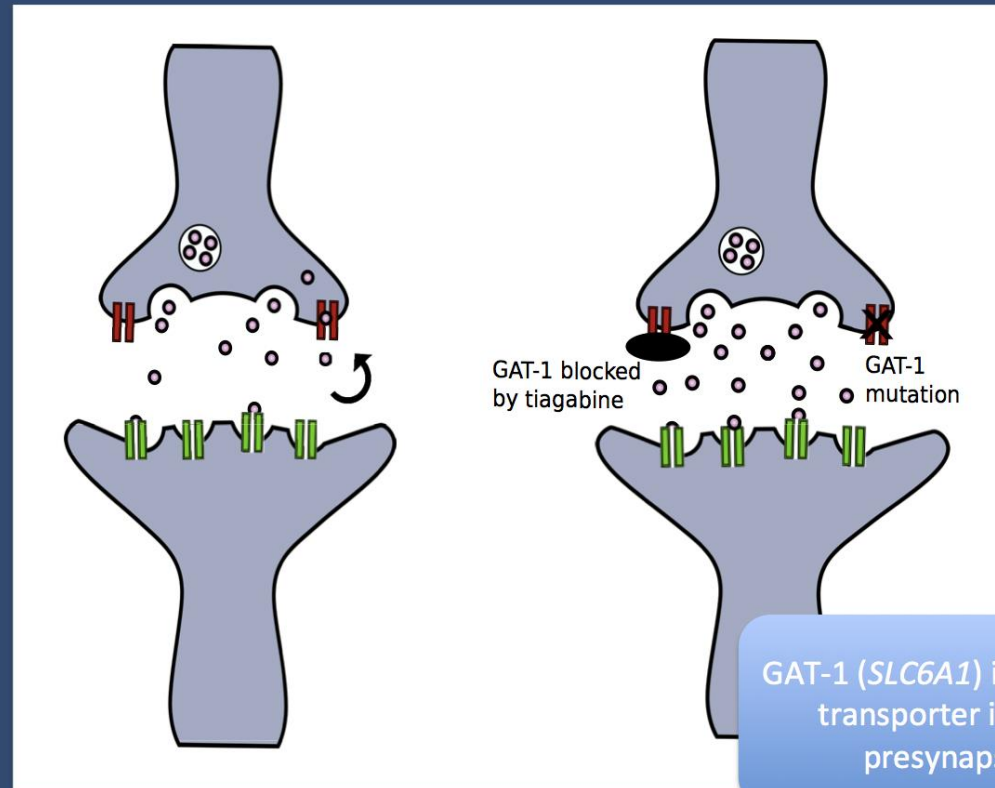
Previous reports have identified *SLC6A1* variants in patients with generalized epilepsies, such as myoclonic-atonic epilepsy and childhood absence epilepsy. However, to date, none of the identified *SLC6A1* variants has been functionally tested for an effect on GAT-1 transporter activity. The purpose of this study was to determine the incidence of *SLC6A1* variants in 460 unselected epilepsy patients and to evaluate the impact of the identified variants on γ -aminobutyric acid (GABA) transport. Targeted resequencing was used to screen 460 unselected epilepsy patients for variants in *SLC6A1*. Five missense variants, one in-frame deletion, one nonsense variant, and one intronic splice-site variant were identified, representing a 1.7% diagnostic yield. Using a [³H]-GABA transport assay, the seven identified exonic variants were found to reduce GABA transport activity. A minigene splicing assay revealed that the splice-site variant disrupted canonical

SLC6A1 Variants Identified in Epilepsy Patients

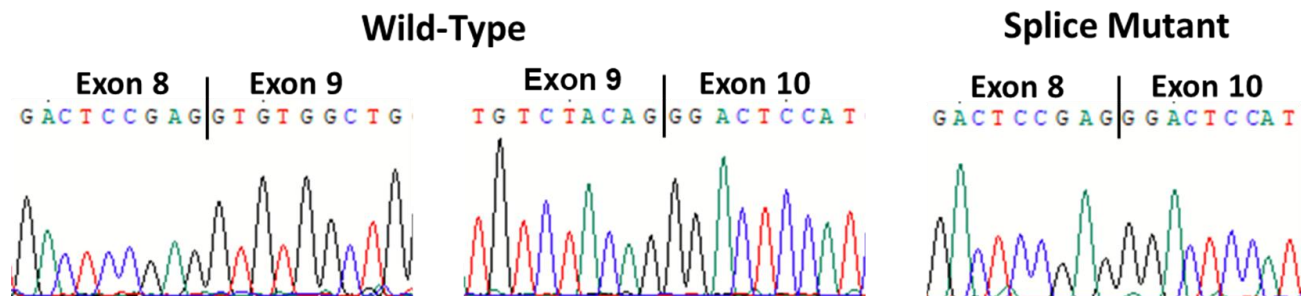
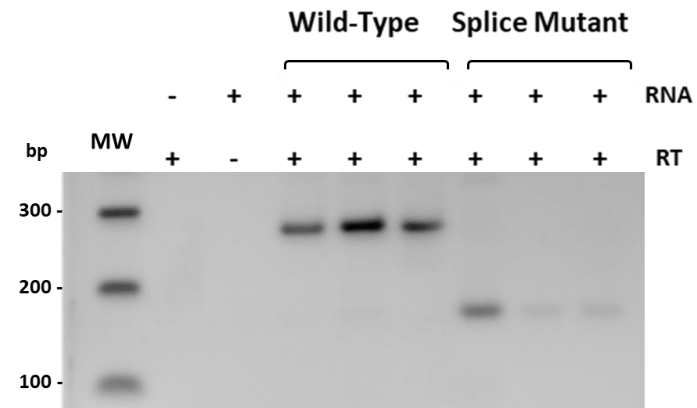
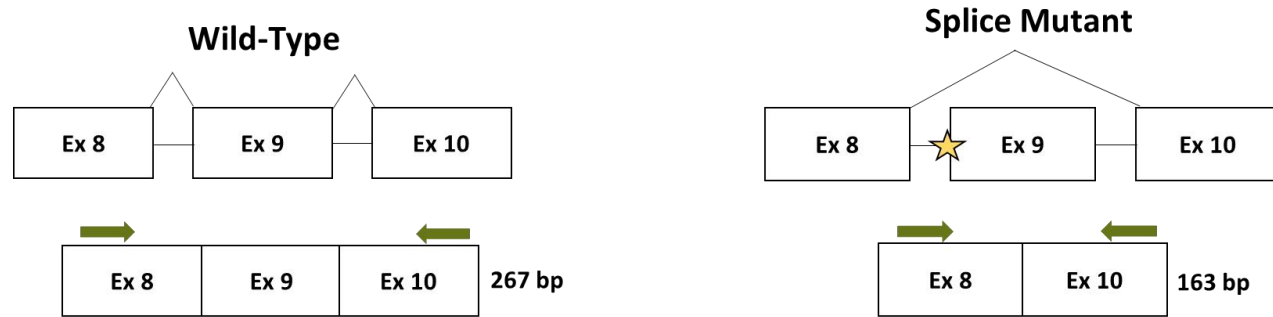


SLC6A1 Encodes the GAT 1 GABA Transporter

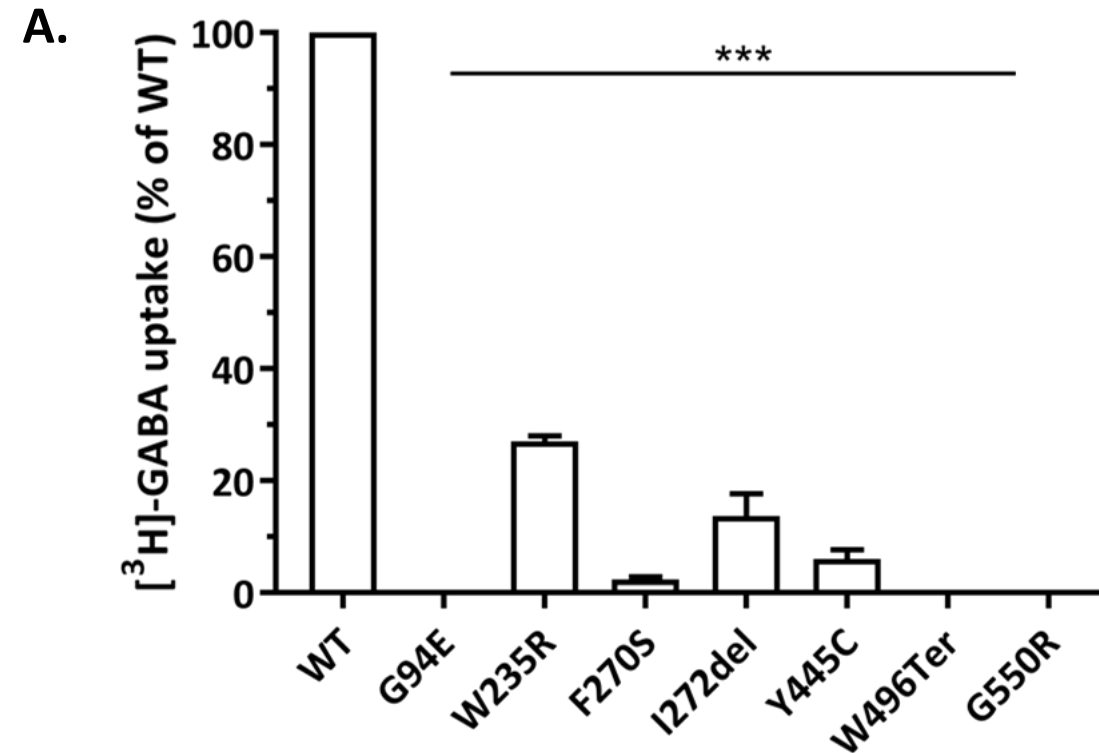
SLC6A1 mutations in Myoclonic Astatic Epilepsy (Carvill et al., 2015)



SLC6A1 c.850-2A>G variant causes exon skipping



GAT 1 mutations reduce GABA transport



No obvious genotype/phenotype correlation

Summary

- *GAT 1* mutations reduce *GABA* clearance.
- Likely to affect neuronal excitability via multiple mechanisms.
- No obvious genotype/phenotype correlation at this time.

Escayg Laboratory



Escayg Laboratory

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