

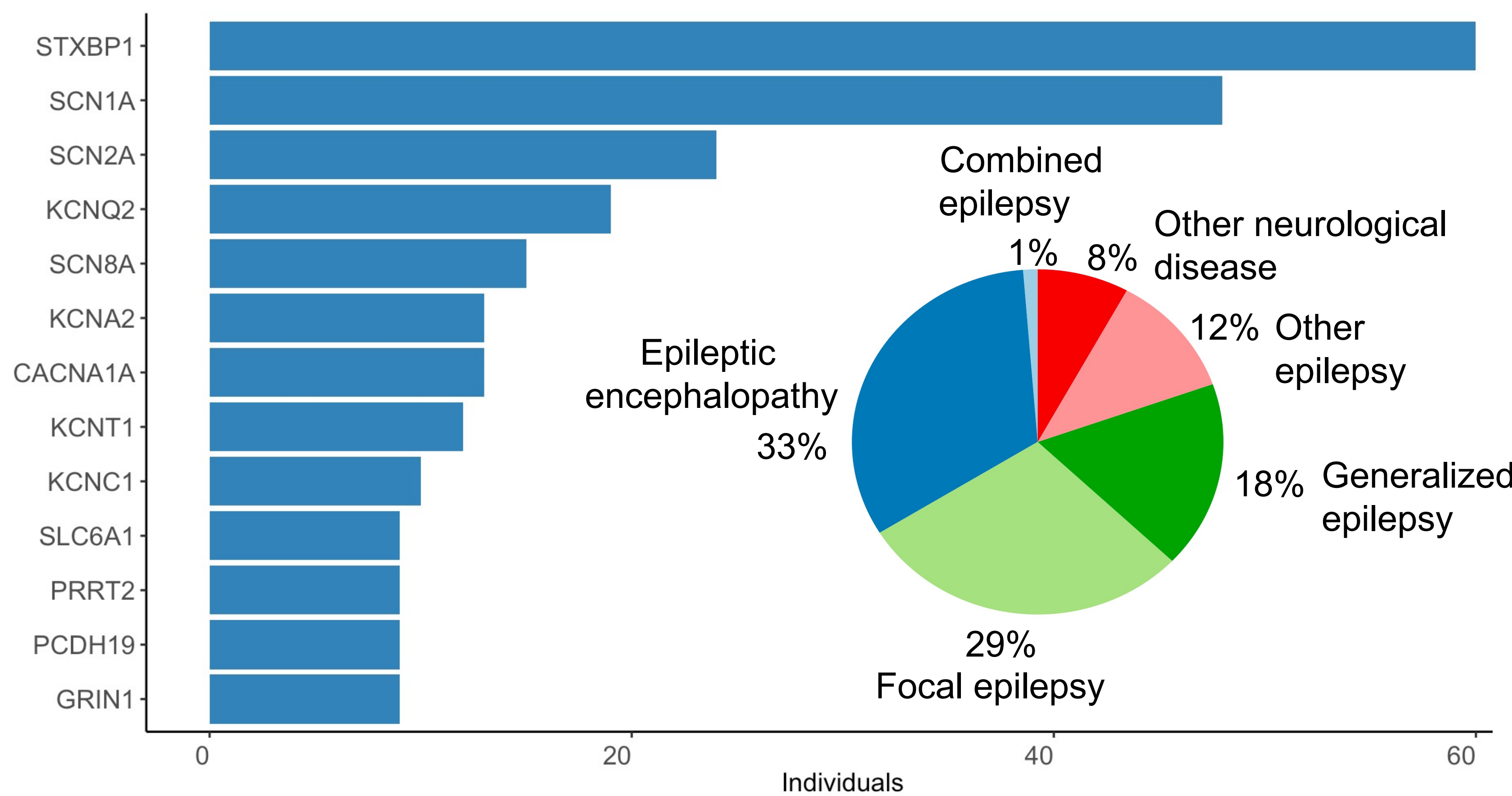
# A data-driven analysis of changes in anti-seizure medications after genetic diagnosis in 1,598 individuals with genetic epilepsies

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## Introduction

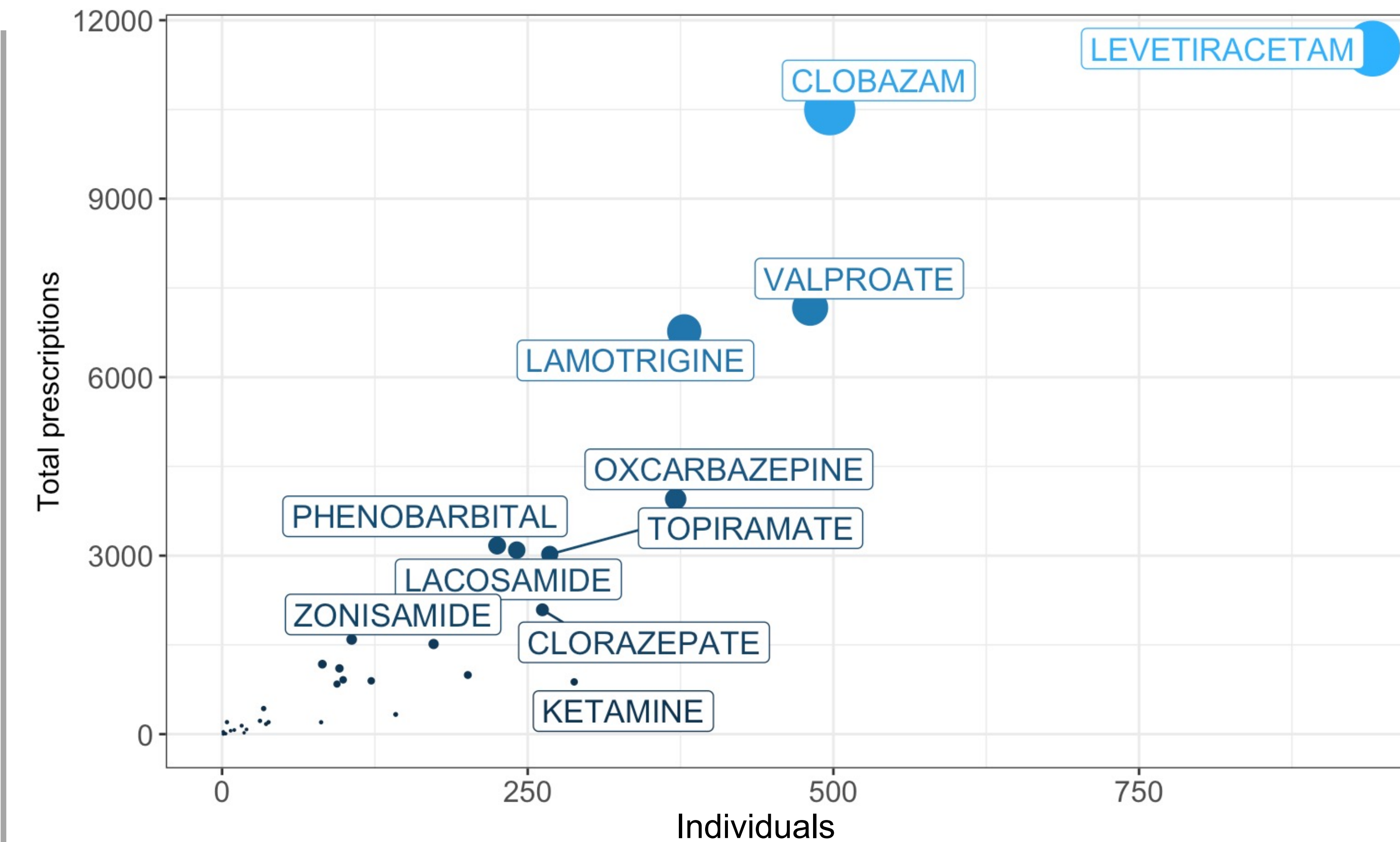
- Genetic etiologies are identified in >30% of epilepsies.
- Precision medicine approaches aim to tailor medication choices to specific genetic etiologies.
- Whether recommended treatment strategies for individual genetic epilepsies are translated into clinical practice remains unknown.
- Medication prescriptions from the electronic medical records (EMR) can be leveraged to track longitudinal anti-seizure medication (ASM) use.



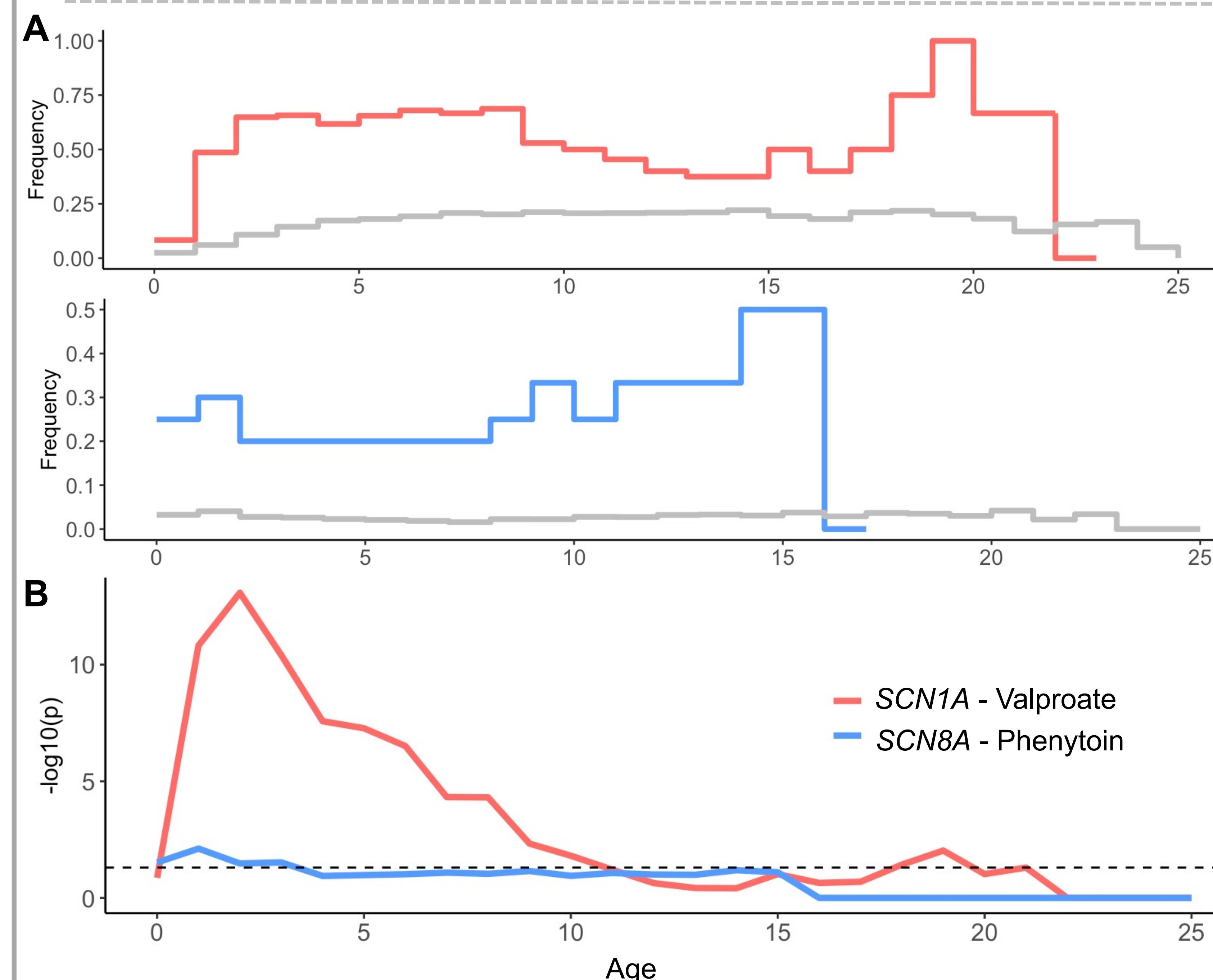
**Fig 1. Overview of genetic etiologies and epilepsy phenotypes in the cohort.** The most common genetic etiologies in the cohort were *STXBP1* ( $n=60$ ), *SCN1A* ( $n=48$ ), and *SCN2A* ( $n=24$ ).

## Methods

- ASM prescription data was extracted from 1,598 individuals with known or presumed genetic epilepsies and binned monthly.
- Longitudinal prescription patterns were assessed across 25 ASMs with a total EMR observation time of 7,872 years.
- ASMs significantly associated with individual genetic etiologies were compared to recommended guidelines.

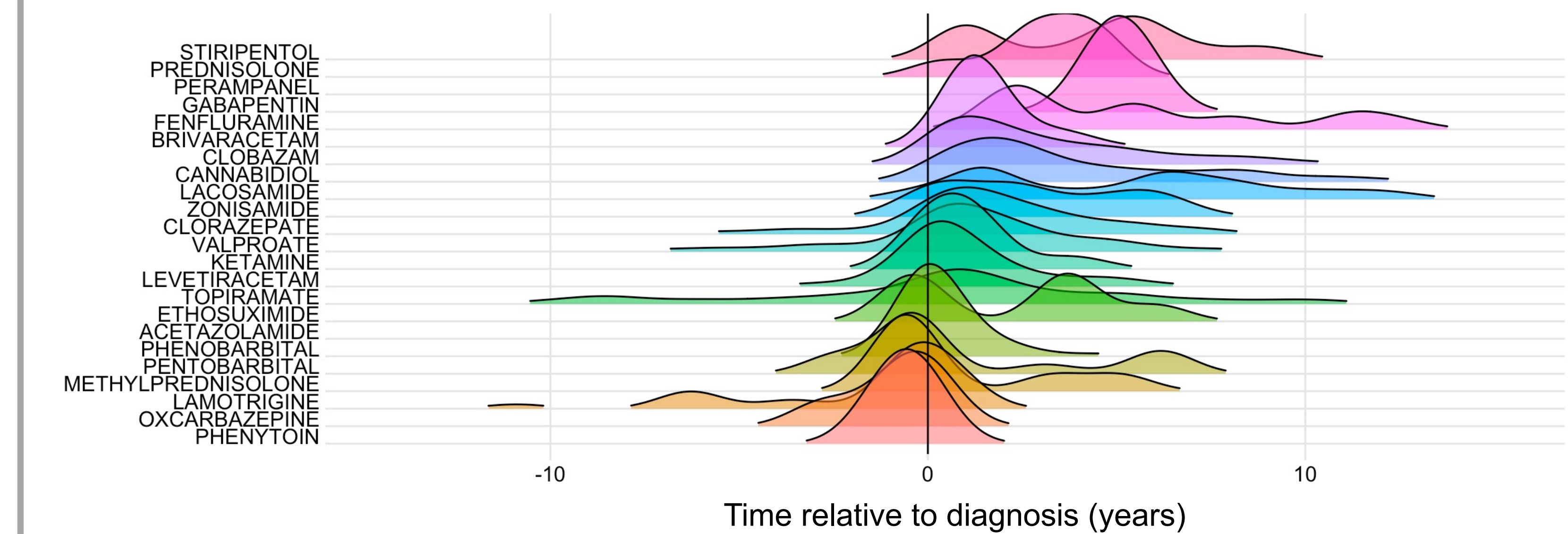


**Fig 2. Distribution of ASMs in the overall cohort.** Levetiracetam ( $n=11,523$ ), clobazam ( $n=10,495$ ), and valproate ( $n=7,167$ ) were the most common ASMs prescribed in the cohort.

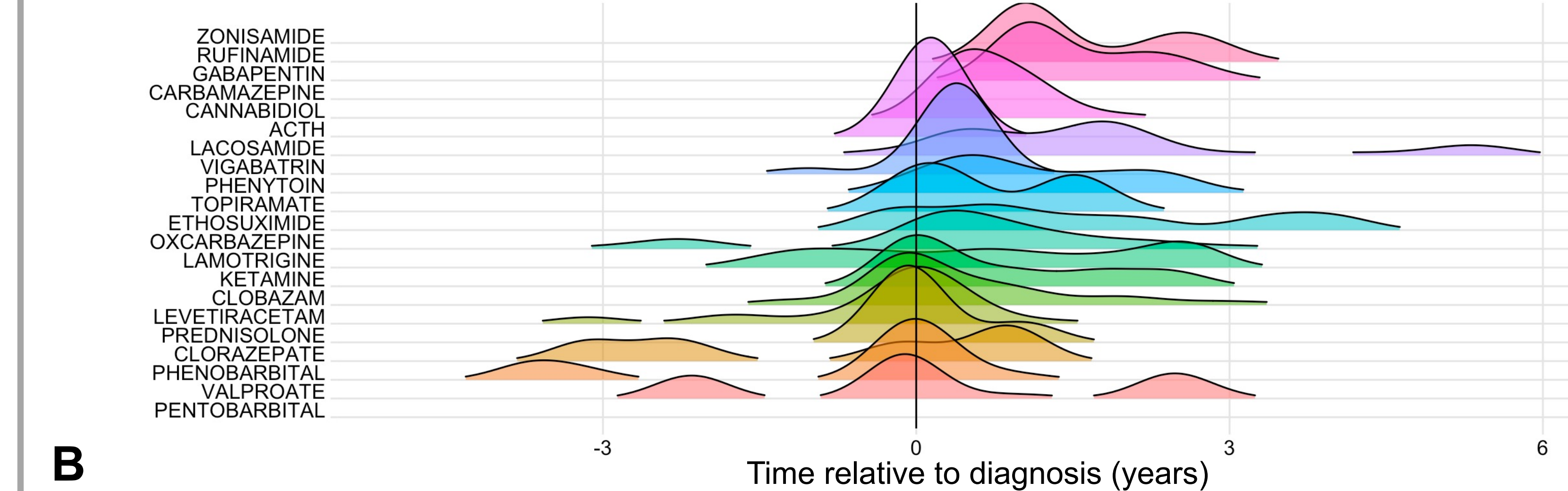


**Fig 3. (A) Frequencies of two select ASMs in *SCN1A* and *SCN8A*-related epilepsies compared to the frequency in the remainder of the cohort (grey). (B) Statistical significance of the difference in frequencies by age.**

## A 4,797 ASM prescriptions in 45 individuals with *SCN1A*



## 1,300 ASM prescriptions in 14 individuals with *SCN8A*



## B

	CBZ	CBD	LEV	TPM	QUIN	VPA	OXC	CZP	LCM	PRED	KET	PHB	PHT	LTG
<i>SCN1A</i>	107.92	81.55	4.61	0.64	-	1.14	0.14	2.51	0.00	-	3.01	3.74	0.07	0.02
<i>PCDH19</i>	100.00	-	10.06	0.00	-	-	-	1.85	-	-	-	0.18	-	0.67
<i>KCNT1</i>	10.70	100.00	0.11	1.62	100.46	0.54	0.00	0.11	0.02	-	13.81	0.36	0.00	-
<i>KCNQ2</i>	3.22	-	1.19	1.85	-	0.08	1.42	6.25	-	48.46	100.00	3.72	-	-
<i>SCN2A</i>	9.49	100.00	1.80	1.09	-	100.00	0.79	18.49	5.08	0.19	10.63	1.99	6.95	0.31
<i>SCN8A</i>	2.55	-	0.39	4.57	-	1.64	0.34	0.13	43.26	2.99	11.52	0.49	3.20	1.14
<i>STXBP1</i>	4.75	-	2.67	0.56	-	-	-	-	-	12.87	-	0.77	-	-
<i>PRRT2</i>	-	-	3.05	-	-	-	3.72	-	-	-	-	0.26	-	-

**Fig 4. (A) Shift in ASM prescription density in *SCN1A* and *SCN8A*-related epilepsies. (B) Odds ratios for ASM changes in temporal relation to genetic diagnosis.**

## Discussion

- The majority of ASMs agree with known recommendations for treatment, such as the initiation of sodium channel blockers in *SCN8A*-related epilepsies.
- A learning health systems approach can be applied to assess clinical decision making in the pediatric epilepsies.

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