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A cross-sectional study of the neuropsychiatric phenotype of *CACNA1C*-related disorder

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Abstract

Background: *CACNA1C* encodes the voltage gated L-type calcium channel Cav1.2. A specific gain of function pathogenic variant in *CACNA1C* causes Timothy syndrome type 1 (TS1) with cardiac long QT syndrome, syndactyly, and neuropsychiatric symptoms. Our previous work found that the TS1 mutation alters neuronal activity-dependent signaling and interneuron migration. Recent case series highlighted a broader spectrum of *CACNA1C*-related disorder (CRD) that includes isolated cardiac disease, isolated neurologic deficits, and TS, but it is unknown how the clinical presentation of other CRD variants relate to neural defects. We surveyed individuals with CRD to define the neuropsychiatric and developmental phenotype in an effort to guide future research into the role of calcium channels in neural development.

Methods: Caregivers of and individuals with CRD completed an online survey of pre- and perinatal events, cardiac events, developmental milestones, neuropsychiatric symptoms, and neuropsychiatric diagnoses. Multiple Mann-Whitney tests were used for comparison of categorical values and Fisher's exact test for comparison of categorical variables between participants with and without cardiac arrhythmia.

Results: Twenty-four participants with germline *CACNA1C* variants including TS1 completed the survey. The most common neuropsychiatric symptoms and/or diagnoses were developmental delay in 92%, incoordination in 71%, hypotonia in 67%, autism spectrum disorder in 50% (autistic features in 92%), seizures in 37.5%, and attention deficit hyperactivity disorder in 21% of participants. There were no significant differences in symptoms between participants with and without arrhythmia.

Conclusions: In our CRD cohort there was an increased prevalence of multiple neuropsychiatric symptoms compared with the general population. These findings indicate the key role of Cav1.2 in brain development and the clinical importance of screening and therapeutically addressing neuropsychiatric symptoms in all individuals with CRD. Future directions include deep phenotyping of neuropsychiatric symptoms and efforts to relate these symptoms to cellular defects.

Keywords: calcium channel, neurodevelopmental delay, autism spectrum disorder, psychiatric disorders

Highlights:

- *CACNA1C*-related disorder (CRD) is a rare genetic disorder with a high risk of neurodevelopmental disorders and risk of cardiac arrhythmia.
- Among 24 individuals with CRD we found high rates of developmental delay, neurologic, and psychiatric symptoms, including a 37.5% prevalence of epilepsy.
- Whether or not an individual had cardiac symptoms did not change these risks.
- These findings suggest that individuals with CRD should be screened and receive care for these common symptoms.

Introduction

Voltage gated L-type calcium channels (LTCC) are critical for transforming chemical to electrical signals in the nervous system and other electrically excitable tissues.^{1,2} Common variants in *CACNA1C*, which encodes the α -subunit of the $Ca_v1.2$ LTCC, are among the most replicable associations in genome-wide association studies of autism spectrum disorder (ASD),³ bipolar disorder,³⁻⁶ depression,³ and schizophrenia.^{3,6-8} The dominant gain of function variant p.G406R in exon 8A of *CACNA1C* causes Timothy syndrome type 1 (TS1), a syndromic form of cardiac long QT interval syndrome (LQTS) with syndactyly, multi-system involvement, and a high prevalence of neurodevelopmental symptoms.⁹ In the original TS1 case series 80% of participants had ASD, thus TS1 has the highest prevalence of syndromic ASD.⁹

Heterozygous microdeletions including *CACNA1C* are associated with intellectual disability and are not known to be associated with cardiac or syndromic features, suggesting channel loss of function via reduced expression may also impact neurodevelopment.¹⁰⁻¹² De novo variants in *CACNA1C*, along with other genes, were enriched in a population with developmental delay.¹³ Additional *CACNA1C* variants were recently identified in a broad spectrum of both cardiac and neurodevelopmental disorders including: typical and atypical syndromic TS,^{9,14-19} structural heart disease with LQTS,²⁰⁻²² isolated long or short QTS,²³⁻²⁶ and isolated neurologic symptoms.²⁷⁻³⁰ This is now referred to as the *CACNA1C*-related disorder (CRD) spectrum that encompasses a range of clinical features caused by pathogenic variants predicted to both increase and decrease channel function.^{31,32}

A better understanding of the epidemiology, clinical spectrum, and biological phenotype of highly penetrant monogenic disease models such as CRD would help unravel how calcium signaling affects neuropsychiatric disorders more broadly and how modulation of calcium

pathways could be leveraged for therapeutic purposes. A major gap in the field is that neuropsychiatric symptoms are not well characterized in the CRD literature beyond the early TS1 cohort and very recent neurologic-only CRD cohort.^{9,27} Case reports of CRD-related arrhythmia report few if any neuropsychiatric symptoms and do not evaluate a temporal relationship with cardiac arrest or other hypoxic risk factors. Individuals with CRD have never been systematically assessed for the prevalence of developmental, neurologic, and psychiatric symptoms, in part due to the rarity of this disorder. Moreover, there is neither clinical nor neuroscience research into whether treating channel dysfunction impacts neuropsychiatric outcomes.

Methods

Participants were enrolled in the ongoing study of neuropsychiatric and developmental symptoms in CRD, which was approved by the Stanford University Institutional Review Board. Participants were identified via the Timothy Syndrome Foundation, the Timothy Syndrome Alliance, and communication with medical providers. Legally authorized representatives gave written informed consent for their children and/or dependents.

Data collection was via a secure online survey hosted in RedCap. Survey questions covered genetic variants, pre- and perinatal comorbidities that influence neurodevelopment, cardiac and medical complications, developmental milestones, neurologic symptoms, psychiatric symptoms, and medications. The survey questions are available upon request. One participant was able to complete the survey for him/herself; all other participants' surveys were completed by caregivers. If answers were unknown (particularly for developmental milestone ages), data was

excluded from analysis. For developmental milestones, participants were excluded from analysis if they were not old enough to have reasonably achieved that milestone. For neuropsychiatric symptoms and diagnoses all 24 participants were considered in analysis. While not all individuals would have been old enough at time of enrollment or age of death to have developed and/or manifested some of these symptoms, to exclude participants by age could over-estimate the statistical effect.

Medical records were obtained from caregivers as available and included verification of *CACNA1C* variants for all participants; insufficient complete records were available to analyze standardized scales such as IQ scores or neuroimaging reports.

Descriptive statistical analysis included percentages for categorical variables and mean and range for continuous variables. The multiple Mann-Whitney tests were used for comparison of continuous values and Fisher's exact test for comparison of categorical variables between participants with and without cardiac involvement. Bonferroni correction for α was computed for each of the developmental, neurologic, and psychiatric comparisons. With our cohort size we had 80% power to detect a difference of 37% between the groups as a two-sided χ^2 test with 5% significance with presumed 15% prevalence.

Results

Demographics

Twenty-four survey participants with CRD had germline variants in *CACNA1C* (**Table 1**). One participant with mosaicism is not reported in this data although this individual reported

symptoms that overlap with the cohort. Participants were from the United States and the United Kingdom. Unfortunately, three participants were deceased prior to study enrollment at a range of 1.6 to 6 years old. Mean age in the cohort at time of completion of survey was 9.8 years (range 0.4 years to 37 years) including the deceased participants and 10.6 years among surviving participants. Nine of the participants (37.5%) were female. Two participants were monozygotic twins, reported here as separate individuals as their symptoms were not completely concordant.

We investigated genotype-phenotype correlation by considering the following sub-populations of CRD. We grouped participants by genotype and presence of LQTS at baseline, which is a well-defined major clinical criterion for TS. Seven participants have TS1 (defined as p.G406R in exon 8A), two have TS type 2 (TS2, defined as p.G406R in exon 8), three have other LQTS CRD (defined as a variant outside of p.G406R plus LQTS at baseline), and 12 have non-LQTS CRD (defined as a variant outside the p.G406R locus and absence of LQTS at baseline). Given the small number of TS2 and other LQTS CRD participants, analyses were performed as a comparison between all LQTS CRD (TS1, TS2, and other LQTS CRD) and non-LQTS CRD participants.

Table 1: CRD cohort demographics and symptoms.

ID	Genotype	Phenotype	Hypoxic event	Developmental delay	Neurologic symptoms	Psychiatric diagnoses
1	p.G406R exon 8A	TS1	Y	Y	Y	Y
2	p.G406R exon 8A	TS1	Y	Y	Y	N
3	p.G406R exon 8A	TS1	Y	Y	Y	Y
4 *	p.G406R exon 8A	TS1	Y	Y	Y	Y
5 *	p.G406R exon 8A	TS1	Y	Y	Y	N
6	p.G406R exon 8A	TS1	Y	Y	Y	Y
7	p.G406R exon 8A	TS1	Y	Y	Y	Y
8	p.G406R exon 8	TS2	Y	Y	Y	N
9	p.G406R exon 8	TS2	Y	Y	Y	Y

10	p.V1363L	Other LQTS CRD	N	Y	Y	N
11	p.C1021R	Other LQTS CRD	N	Y	Y	N
12	p.G402S	Other LQTS CRD	Y	Y	Y	Y
13	p.E2062X	Non-LQTS CRD	Y	Y	Y	Y
14	p.V403M	Non-LQTS CRD	N	Y	Y	Y
15	p.A1521P	Non-LQTS CRD	N	Y	Y	Y
16	Deletion of exon 3	Non-LQTS CRD	N	N	N	Y
17	p.L658P	Non-LQTS CRD	N	Y	Y	N
18	p.L207R	Non-LQTS CRD	N	Y	Y	Y
19	p.V1167A	Non-LQTS CRD	N	Y	Y	N
20 *	p.V1363M	Non-LQTS CRD	N	Y	Y	N
21	p.G1173_Q1175del	Non-LQTS CRD	N	Y	Y	Y
22	p.R1332Q	Non-LQTS CRD	Y	Y	N	Y
23	p.R1332Q	Non-LQTS CRD	N	Y	N	Y
24	p.T941I	Non-LQTS CRD	N	Y	Y	N

* Indicates deceased participant. Abbreviations: CRD: *CACNA1C*-related disorder; LQTS: long QT syndrome; TS1: Timothy syndrome type 1; TS2: Timothy syndrome type 2.

Developmental risk factors

Overall, 54.2% of participants reported a major hypoxic event, defined as perinatal respiratory distress, perinatal hypoxic ischemic encephalopathy, or cardiac arrest (**Table 2**).

Major hypoxic events tended to be more common in LQTS CRD than non-LQTS CRD participants (83.3% vs. 25%, Fisher's exact test $P = 0.012$, Bonferroni corrected for $\alpha = 0.00167$ for developmental comparisons). [Because most participants' medical records were not available to determine the extent of hypoxic injury and neurodevelopmental symptom severity before and after the events, we were unable to use reported hypoxic events as an outcome measure in our stratified analysis.](#)

Prematurity (birth before 36 weeks gestational age) was reported in 29.2% of all participants. Premature birth was not significantly more common in LQTS CRD than non-LQTS CRD participants (33.3% vs. 25%, Fisher's exact test $P = 1$).

Table 2: Percent of CRD participants with commonly reported neuropsychiatric symptoms.

Symptom (%)	All CRD (n = 24)	LQTS CRD (n = 12)	Non-LQTS CRD (n = 12)	Fisher's exact P-value
Hypoxic event	54.2	83.3	25	0.01
Learning or intellectual disability	70.8	66.7	75	1
Incoordination	70.8	66.7	75	1
Dysphagia	62.5	58.3	66.7	1
Hypotonia	66.7	50	83.3	0.19
Seizures, epilepsy	37.5	41.7	33.3	1
Any psychiatric dx	62.5	58.3	66.7	1
ASD dx	50	50	50	1
Aggression, self-injurious behavior	29.2	33.3	25	1
ADHD dx	20.8	16.7	25	1
Anxiety dx	16.7	16.7	16.7	1

Percent of participants with reported neurologic and psychiatric symptoms for all participants and genetic subsets. Fisher's exact test two tailed P-value was calculated on the number of participants with and without cardiac arrhythmia. Abbreviations: ADHD: attention deficit hyperactivity disorder; ASD: autism spectrum disorder; CRD: *CACNA1C*-related disorder; dx: diagnosis (indicates report of a formal diagnosis); LQTS: long QT syndrome.

Developmental phenotype

Developmental delay was present in 91.7% of all participants, with 91.7% of participants reporting motor delay and 75% reporting language delay (**Supplemental Table 1**). Seventeen percent of participants were not yet able to walk and/or speak and other language and motor milestones were similarly affected. There was a trend that non-LQTS CRD participants were less likely to be able to learn to ride a bicycle or have imaginative play, but there were no significant differences between participants with and without LQTS in the ability to achieve any of the milestones after multiple comparison correction.

Among participants who achieved developmental milestones, the average age tended to be higher than typical development on the Denver II Developmental Scale across motor, language, and social domains and across all subtypes of CRD (**Figure 1A, B**).³³ Participants with non-LQTS CRD tended to be more delayed on development of first spoken word and independent sitting compared to participants with LQTS CRD (**Supplemental Table 2**).

Neurologic phenotype

Neurologic symptoms of any kind were reported in 95.8% of all participants (**Table 2, Supplemental Table 3**). The most common symptoms included learning disability and/or intellectual disability, incoordination, dysphagia, hypotonia, and altered sensitivity to noxious stimuli (**Figure 1C**). Seizures or epilepsy were reported in 37.5% of all participants. Of note, while intellectual and learning disabilities were common, 33.3% of participants reported highly accurate memory or other above-average skills for age.

No symptoms were significantly different between participants with or without arrhythmia. There was a trend that imbalance and abnormal extraocular movements were more common in non-LQTS CRD.

Psychiatric phenotype

A spectrum of psychiatric symptoms and diagnoses were reported. Sixty-three percent of participants reported a formal diagnosis of at least one psychiatric disorder (**Table 2, Supplemental Table 4**). The most commonly reported diagnoses were ASD in 50% of all participants, attention deficit hyperactivity disorder (ADHD) in 20.8%, and anxiety disorder in 16.7% (**Figure 1D**). When asked about symptoms, 91.7% of participants reported at least one

behavioral feature associated with ASD, 54.2% reported a shorter attention span, 29.2% reported aggressive behavior towards self or others, and 25% reported a specific phobia. There were no significant differences in self-reported symptoms nor in reported formal diagnoses between participants with and without LQTS.

Of note, one older participant with TS1 was the only individual with diagnoses of obsessive-compulsive disorder, oppositional defiant disorder, depression, mania and/or bipolar disorder, and schizophrenia, in addition to ASD and anxiety disorder.

Discussion

CACNA1C-related disorder is a syndromic cause of cardiac arrhythmia that also impacts neurodevelopment. We surveyed a cohort of 24 participants with CRD (out of approximately 200 individuals in the literature, global incidence unknown) to catalog more broadly what types of developmental, neurologic, and psychiatric symptoms they experienced and found a high prevalence of all symptoms.

Ninety-two percent of participants reported motor and/or language developmental delay and 96% reported at least one neurologic symptom. Some of the most common neurologic symptoms reported were learning or intellectual disability, incoordination, concerns for feeding safety, hypotonia, and altered sensory sensitivity. Seizures or epilepsy were reported in 37.5% suggesting that education about seizure semiology and safety could be beneficial for individuals with CRD in addition to detailed screening.

The most commonly reported psychiatric symptoms were features of ASD in 92% of participants with a formal diagnosis of ASD in 50%. Other commonly reported symptoms included short attention span, aggressive behavior, and specific phobias (never before reported in

this disorder). Of note, while 63% of participants reported at least one formal psychiatric diagnosis, many reported symptoms but lacked a formal diagnosis, raising concern that these individuals may not have received psychiatric clinical evaluation or support for symptom management.

There are no significant differences in neuropsychiatric symptoms between participants with and without LQTS despite a trend toward a difference in the history of major hypoxic events. This suggests that the neuronal calcium channel alteration is the primary etiology for CRD neuropsychiatric symptoms and that cardiac-related hypoxic injury is a secondary contributing factor. These findings highlight the essential role of Cav1.2 in neurodevelopment.

Our data align with and reinforce findings from the contemporaneously conducted Rodan *et al.* study examining neurologic-only CRD.²⁷ We found comparable prevalence of neuropsychiatric symptoms even in the absence of cardiac symptoms. Together, these results demonstrate that *CACNA1C* plays an important role in neurodevelopment, cognition, and behavioral control. Our cohort is gene-centric, not symptom-centric, permitting a less biased ascertainment of the prevalence of neuropsychiatric symptoms. Thus, our data help confirm prior literature on the presence and spectrum of neuropsychiatric and developmental disorders across all subtypes of CRD.^{9,27-30}

Two prior publications reported on the same individual with atypical TS (with LQTS) and bipolar disorder who experienced episodic psychosis that improved on verapamil, as well as subjective improvement in cognition on this medication.^{34,35} We report one older participant who received multiple psychiatric diagnoses. These results suggest a need for longitudinal follow up to determine if other individuals with CRD develop bipolar disorder, OCD, and/or psychosis.

There are notable recent advances from animal and human models of CRD.³⁶ Heterozygous *Cacna1c* knock out rats had altered associative learning,³⁷ as well as alterations in network oscillations and hippocampal plasticity markers, which were rescued by the neurotrophin receptor agonist LM22B-10.³⁸ A knock-in mouse model of TS2 displayed restricted and repetitive behaviors.³⁹ In parallel, 2D neurons and cortical organoids derived from individuals with TS1 had altered gene expression, increased neurotransmitters (norepinephrine and dopamine), and aberrant differentiation.⁴⁰ In forebrain assembloids, TS1-derived interneurons demonstrated abnormal migration that was related to increased myosin light chain phosphorylation and GABA-A receptor sensitivity.^{41,42} It remains to be determined how these cellular level abnormalities can cause network level alterations that manifest as developmental and psychiatric symptoms seen in rodent models and individuals with CRD. We hope that continued research from clinical studies partnered with basic neuroscience will hopefully yield answers about the role of *CACNA1C* in typical development, the pathophysiology of CRD, and insight into treatments.

Limitations

We acknowledge that our study has limitations stemming from cohort size and survey-based data acquisition methods. Our sample size was inherently limited by participant availability since CRD is a rare genetic disorder, which meant we were only powered to detect a large difference between subgroups. This also prevented a confirmatory replication study and more comprehensive statistical sub-analyses such as comparisons by hypoxic exposure and genotype-phenotype correlations. [Caregiver-reported hypoxic brain injury was common in this cohort and](#)

could contribute to neuropsychiatric symptoms; future studies should use medical records to assess symptoms before and after hypoxic episodes to disentangle what symptoms are attributable to hypoxic injury as opposed to CRD primarily. We were further limited to participants with sufficient English fluency to complete the survey. We classified participants by self-reported LQTS at baseline when there is a spectrum of QT interval involvement that may present later in life or under exercise stress.

This study was designed as a preliminary evaluation to determine if the subjective observations by caregivers and medical professionals that individuals with CRD often had developmental and neuropsychiatric symptoms was in fact correct. This study therefore involves a participant or caregiver-report based survey which, while based on neurodevelopmental and psychiatric symptom inventories, does not incorporate validated diagnostic tools. We recognize that this limits the accuracy of the data. Moreover, a survey approach may incorporate several forms of bias such as selection/ascertainment, recall, self-reporting, and/or diagnostic access biases. We are addressing some of these limitations via an ongoing deeper phenotyping study incorporating standardized assessment measures within this cohort.

Conclusions

In a self- or caregiver-reported survey of 24 individuals with *CACNA1C*-related disorder we found pervasive developmental, neurologic, and psychiatric symptoms, including a 37.5% prevalence of epilepsy and 50% prevalence of ASD. These data are in alignment with prior publications on the risk of neurodevelopmental disorders in CRD and expand the spectrum of symptoms in this disorder. There were no significant differences in symptom prevalence between participants with and without long QT syndrome, suggesting that there are more similarities than

distinctions when it comes to the impact of *CACNA1C* variants on the brain. In the absence of genotype-phenotype correlation, individuals with CRD should be screened and monitored for a broad spectrum of neuropsychiatric symptoms throughout their lifespan in order to properly diagnose, treat, and support maximal development. These findings will also inform research into the role of *CACNA1C* and calcium channels in neurodevelopment and neural function. [Future directions include a prospective study with clinical neurologic and psychiatric evaluations incorporating validated diagnostic tools in order to more accurately assess and follow neurodevelopmental phenotypes in individuals with CRD.](#)

Figure Legends

Figure 1:

(A) Mean age in months or (B) years at which participants achieved developmental milestones.

All participants are depicted next to participants with and without LQTS. There were no significant differences between subgroups. Age at achievement in 50% of typical population denoted by purple lines derived from the Denver II Developmental Scale.³³ Error bars show standard error of the mean.

(C) Percent of participants who reported the most common neurologic symptoms. There were no significant differences between subgroups.

(D) Percent of participants who reported the most common psychiatric diagnoses. There were no significant differences between subgroups.

Abbreviations: ADHD: attention deficit hyperactivity disorder; ASD: autism spectrum disorder; CRD: *CACNA1C*-related disorder; dx: diagnosis; ID: intellectual disability; LD: learning

disability; OCD: obsessive compulsive disorder; ODD: oppositional defiant disorder; Scz: schizophrenia; sx: symptoms.

List of abbreviations

ADHD: attention deficit hyperactivity disorder; ASD: autism spectrum disorder; CRD: *CACNA1C*-related disorder; LQTS: long QT interval syndrome; LTCC: L-type calcium channel; OCD: obsessive compulsive disorder; TS: Timothy syndrome; TS1: Timothy syndrome type 1; TS2: Timothy syndrome type 2.

Additional Files:

Supplemental Table 1: Percent of participants with delays or inability to achieve developmental milestones.

Supplemental Table 2: Mean age at which participants achieved developmental milestones.

Supplemental Table 3: Percent of participants with reported neurologic symptoms.

Supplemental Table 4: Percent of participants with reported psychiatric symptoms and diagnoses.

Declarations

1. Ethics approval and consent to participate: Ethics approval was provided by the Stanford University Institutional Review Board (Protocol 53610). Participants in the USA were consented via telephone call and written consent, while participants outside the USA completed an online written consent. Legally authorized representatives consented for minors or for adult participants without capacity.

2. Consent for publication: Participants consented to limited, anonymized information sharing for publication and research.
3. Availability of data and materials: Most of the data generated and analyzed during this study are available in the tables and supplemental files. Full data are available from the corresponding author on reasonable request to protect the individual privacy of the participants.
4. Competing interests: JFGU, KWT, SPP declare no competing interests. RJL has received funding from Tome Bio for unrelated consulting work. JH has received funding from Takeda Pharmaceutical Company for unrelated work.
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6. Authors' contributions: RJL, KWT, JFGU, JH, JAB, and SPP conceived and designed the study. RJL, KWT, and JAB generated the survey. KWT, RJL, JFGU, JH, and SPP

recruited participants and RJJ consented all participants and analyzed the data. All authors read and approved the final manuscript.

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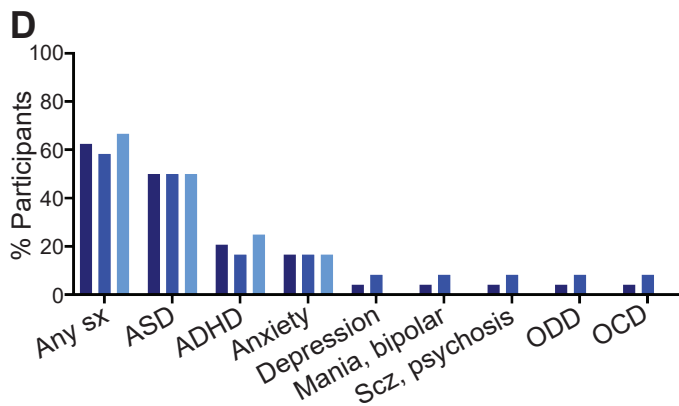
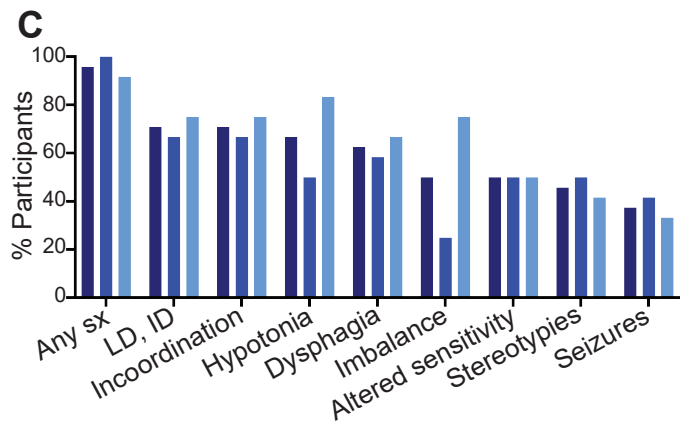
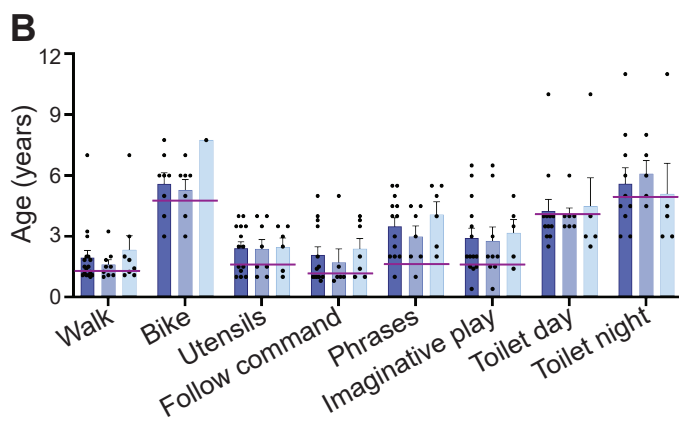
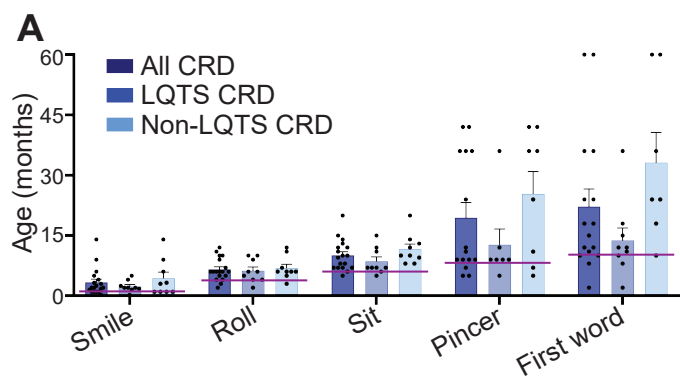
References

1. Flavell SW, Greenberg ME. Signaling mechanisms linking neuronal activity to gene expression and plasticity of the nervous system. *Annu Rev Neurosci.* 2008;31:563-590. doi:10.1146/annurev.neuro.31.060407.125631
2. Greer PL, Greenberg ME. From synapse to nucleus: calcium-dependent gene transcription in the control of synapse development and function. *Neuron.* 2008;59(6):846-860. doi:10.1016/j.neuron.2008.09.002
3. Cross-Disorder Group of the Psychiatric Genomics Consortium. Identification of risk loci with shared effects on five major psychiatric disorders: a genome-wide analysis. *Lancet.* 2013;381(9875):1371-1379. doi:10.1016/S0140-6736(12)62129-1
4. Green EK, Grozeva D, Jones I, et al. The bipolar disorder risk allele at CACNA1C also confers risk of recurrent major depression and of schizophrenia. *Mol Psychiatry.* 2010;15(10):1016-1022. doi:10.1038/mp.2009.49
5. Ferreira MAR, O'Donovan MC, Meng YA, et al. Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. *Nat Genet.* 2008;40(9):1056-1058. doi:10.1038/ng.209
6. Moskvina V, Craddock N, Holmans P, et al. Gene-wide analyses of genome-wide association data sets: evidence for multiple common risk alleles for schizophrenia and bipolar disorder and for overlap in genetic risk. *Mol Psychiatry.* 2009;14(3):252-260. doi:10.1038/mp.2008.133
7. Schizophrenia Working Group of the Psychiatric Genomics Consortium. Biological insights from 108 schizophrenia-associated genetic loci. *Nature.* 2014;511(7510):421-427. doi:10.1038/nature13595
8. Hamshere ML, Walters JTR, Smith R, et al. Genome-wide significant associations in schizophrenia to ITIH3/4, CACNA1C and SDCCAG8, and extensive replication of associations reported by the Schizophrenia PGC. *Mol Psychiatry.* 2013;18(6):708-712. doi:10.1038/mp.2012.67
9. Splawski I, Timothy KW, Sharpe LM, et al. Ca(V)1.2 calcium channel dysfunction causes a multisystem disorder including arrhythmia and autism. *Cell.* 2004;119(1):19-31. doi:10.1016/j.cell.2004.09.011
10. Mio C, Passon N, Baldan F, et al. CACNA1C haploinsufficiency accounts for the common features of interstitial 12p13.33 deletion carriers. *Eur J Med Genet.* 2020;63(4):103843. doi:10.1016/j.ejmg.2020.103843
11. Quintela I, Eirís J, Gómez-Lado C, et al. Copy number variation analysis of patients with intellectual disability from North-West Spain. *Gene.* 2017;626:189-199. doi:10.1016/j.gene.2017.05.032

12. Roberts JL, Hovanes K, Dasouki M, Manzardo AM, Butler MG. Chromosomal microarray analysis of consecutive individuals with autism spectrum disorders or learning disability presenting for genetic services. *Gene*. 2014;535(1):70-78. doi:10.1016/j.gene.2013.10.020
13. Kaplanis J, Samocha KE, Wiel L, et al. Evidence for 28 genetic disorders discovered by combining healthcare and research data. *Nature*. 2020;586(7831):757-762. doi:10.1038/s41586-020-2832-5
14. Splawski I, Timothy KW, Decher N, et al. Severe arrhythmia disorder caused by cardiac L-type calcium channel mutations. *Proc Natl Acad Sci U S A*. 2005;102(23):8089-8096; discussion 8086-8088. doi:10.1073/pnas.0502506102
15. Fröhler S, Kieslich M, Langnick C, et al. Exome sequencing helped the fine diagnosis of two siblings afflicted with atypical Timothy syndrome (TS2). *BMC Med Genet*. 2014;15:48. doi:10.1186/1471-2350-15-48
16. Gillis J, Burashnikov E, Antzelevitch C, et al. Long QT, syndactyly, joint contractures, stroke and novel CACNA1C mutation: expanding the spectrum of Timothy syndrome. *Am J Med Genet A*. 2012;158A(1):182-187. doi:10.1002/ajmg.a.34355
17. Krause U, Gravenhorst V, Kriebel T, Ruschewski W, Paul T. A rare association of long QT syndrome and syndactyly: Timothy Syndrome (LQT 8). *Clin Res Cardiol*. 2011;100(12):1123-1127. doi:10.1007/s00392-011-0358-4
18. Boczek NJ, Miller EM, Ye D, et al. Novel Timothy syndrome mutation leading to increase in CACNA1C window current. *Heart Rhythm*. 2015;12(1):211-219. doi:10.1016/j.hrthm.2014.09.051
19. Hennessey JA, Boczek NJ, Jiang YH, et al. A CACNA1C variant associated with reduced voltage-dependent inactivation, increased CaV1.2 channel window current, and arrhythmogenesis. *PLoS One*. 2014;9(9):e106982. doi:10.1371/journal.pone.0106982
20. Boczek NJ, Ye D, Jin F, et al. Identification and Functional Characterization of a Novel CACNA1C-Mediated Cardiac Disorder Characterized by Prolonged QT Intervals With Hypertrophic Cardiomyopathy, Congenital Heart Defects, and Sudden Cardiac Death. *Circ Arrhythm Electrophysiol*. 2015;8(5):1122-1132. doi:10.1161/CIRCEP.115.002745
21. Boczek NJ, Best JM, Tester DJ, et al. Exome sequencing and systems biology converge to identify novel mutations in the L-type calcium channel, CACNA1C, linked to autosomal dominant long QT syndrome. *Circ Cardiovasc Genet*. 2013;6(3):279-289. doi:10.1161/CIRCGENETICS.113.000138
22. Landstrom AP, Boczek NJ, Ye D, et al. Novel long QT syndrome-associated missense mutation, L762F, in CACNA1C-encoded L-type calcium channel imparts a slower inactivation tau and increased sustained and window current. *Int J Cardiol*. 2016;220:290-298. doi:10.1016/j.ijcard.2016.06.081

23. Wemhöner K, Friedrich C, Stallmeyer B, et al. Gain-of-function mutations in the calcium channel CACNA1C (Cav1.2) cause non-syndromic long-QT but not Timothy syndrome. *J Mol Cell Cardiol.* 2015;80:186-195. doi:10.1016/j.yjmcc.2015.01.002
24. Mellor GJ, Panwar P, Lee AK, et al. Type 8 long QT syndrome: pathogenic variants in CACNA1C-encoded Cav1.2 cluster in STAC protein binding site. *Europace.* 2019;21(11):1725-1732. doi:10.1093/europace/euz215
25. Fukuyama M, Ohno S, Wang Q, et al. L-type calcium channel mutations in Japanese patients with inherited arrhythmias. *Circ J.* 2013;77(7):1799-1806. doi:10.1253/circj.cj-12-1457
26. Liu X, Shen Y, Xie J, et al. A mutation in the CACNA1C gene leads to early repolarization syndrome with incomplete penetrance: A Chinese family study. *PLoS One.* 2017;12(5):e0177532. doi:10.1371/journal.pone.0177532
27. Rodan LH, Spillmann RC, Kurata HT, et al. Phenotypic expansion of CACNA1C-associated disorders to include isolated neurological manifestations. *Genet Med.* Published online June 23, 2021. doi:10.1038/s41436-021-01232-8
28. Bozarth X, Dines JN, Cong Q, et al. Expanding clinical phenotype in CACNA1C related disorders: From neonatal onset severe epileptic encephalopathy to late-onset epilepsy. *Am J Med Genet A.* 2018;176(12):2733-2739. doi:10.1002/ajmg.a.40657
29. Iossifov I, O’Roak BJ, Sanders SJ, et al. The contribution of de novo coding mutations to autism spectrum disorder. *Nature.* 2014;515(7526):216-221. doi:10.1038/nature13908
30. Deciphering Developmental Disorders Study. Prevalence and architecture of de novo mutations in developmental disorders. *Nature.* 2017;542(7642):433-438. doi:10.1038/nature21062
31. Han D, Xue X, Yan Y, Li G. Dysfunctional Cav1.2 channel in Timothy syndrome, from cell to bedside. *Exp Biol Med (Maywood).* 2019;244(12):960-971. doi:10.1177/1535370219863149
32. Bauer R, Timothy KW, Golden A. Update on the Molecular Genetics of Timothy Syndrome. *Front Pediatr.* 2021;9:668546. doi:10.3389/fped.2021.668546
33. Frankenburg W, Dodds J. DENVER II Training Manual. Published online 1992.
34. Jacobs A, Knight BP, McDonald KT, Burke MC. Verapamil decreases ventricular tachyarrhythmias in a patient with Timothy syndrome (LQT8). *Heart Rhythm.* 2006;3(8):967-970. doi:10.1016/j.hrthm.2006.04.024
35. Gershon ES, Grennan K, Busnello J, et al. A rare mutation of CACNA1C in a patient with bipolar disorder, and decreased gene expression associated with a bipolar-associated common SNP of CACNA1C in brain. *Mol Psychiatry.* 2014;19(8):890-894. doi:10.1038/mp.2013.107

36. Moon AL, Haan N, Wilkinson LS, Thomas KL, Hall J. CACNA1C: Association With Psychiatric Disorders, Behavior, and Neurogenesis. *Schizophr Bull.* 2018;44(5):958-965. doi:10.1093/schbul/sby096
37. Moon AL, Brydges NM, Wilkinson LS, Hall J, Thomas KL. Cacna1c Hemizyosity Results in Aberrant Fear Conditioning to Neutral Stimuli. *Schizophr Bull.* Published online January 7, 2020:sbz127. doi:10.1093/schbul/sbz127
38. Tigaret CM, Lin TCE, Morrell ER, et al. Neurotrophin receptor activation rescues cognitive and synaptic abnormalities caused by hemizyosity of the psychiatric risk gene Cacna1c. *Mol Psychiatry.* 2021;26(6):1748-1760. doi:10.1038/s41380-020-01001-0
39. Bader PL, Faizi M, Kim LH, et al. Mouse model of Timothy syndrome recapitulates triad of autistic traits. *Proc Natl Acad Sci U S A.* 2011;108(37):15432-15437. doi:10.1073/pnas.1112667108
40. Paşca SP, Portmann T, Voineagu I, et al. Using iPSC-derived neurons to uncover cellular phenotypes associated with Timothy syndrome. *Nat Med.* 2011;17(12):1657-1662. doi:10.1038/nm.2576
41. Birey F, Andersen J, Makinson CD, et al. Assembly of functionally integrated human forebrain spheroids. *Nature.* 2017;545(7652):54-59. doi:10.1038/nature22330
42. Birey F, Li MY, Gordon A, et al. Dissecting the molecular basis of human interneuron migration in forebrain assembloids from Timothy syndrome. *Cell Stem Cell.* 2022;29(2):248-264.e7. doi:10.1016/j.stem.2021.11.011



Supplemental Table 1: Percent of participants with developmental delays or the inability to achieve age-appropriate developmental milestones.

Milestone	All CRD	LQTS CRD	Non-LQTS CRD	Fisher's exact P-value
Language delay (%)	75	66.7	83.3	0.64
Motor delay (%)	91.7	91.7	91.7	1
Unable to achieve milestone despite appropriate age (% , n)				
Rolling	8.3 (2/24)	8.3 (1/12)	8.3 (1/12)	1
Sitting	8.7 (2/23)	8.3 (1/12)	9.1 (1/11)	1
Walking	17.4 (4/23)	16.7 (2/12)	18.2 (2/11)	1
Bicycling	42.1 (8/19)	18.2 (2/11)	75 (6/8)	0.02
Pincer	8.7 (2/23)	8.3 (1/12)	9.1 (1/11)	1
Utensils	21.7 (5/23)	8.3 (1/12)	36.4 (4/11)	0.16
First word	17.4 (4/23)	8.3 (1/12)	27.3 (3/11)	0.32
Speak in phrases	18.2 (4/22)	8.3 (1/12)	30 (3/10)	0.29
Follow a command	17.4 (4/23)	8.3 (1/12)	27.3 (3/11)	0.32
Smile	4.2 (1/24)	8.3 (1/12)	0 (0/12)	1
Imaginative play	26.1 (6/23)	8.3 (1/12)	45.5 (5/11)	0.07
Toilet trained day	25 (5/20)	18.2 (2/11)	33.3 (3/10)	0.64
Toilet trained night	22.2 (4/18)	20 (2/10)	25 (2/8)	1

The percent of all participants and subsets with and without cardiac arrhythmia are shown for presence of language and motor delay and then inability to achieve developmental milestones. For the first two rows of developmental delay, n=24 for all CRD, n=12 for both LQTS and non-LQTS CRD. For other rows, n is adjusted for participants who are at least average age for that milestone on the Denver II Scale and indicated in parentheses as fraction unable to achieve despite appropriate age. Bonferroni multiple comparison correction $P < 0.00167$ for comparisons of development in **Supplemental Tables 1 and 2**. Fisher's exact test two tailed P-value was calculated on the number of participants with and without cardiac arrhythmia.

Abbreviations: CRD: *CACNA1C*-related disorder; LQTS: long QT syndrome; TS1: Timothy syndrome type 1; TS2: Timothy syndrome type 2.

Supplemental Table 2: Mean age at which participants achieved developmental milestones.

Mean age to achieve milestone	All CRD	TS1	TS2	Other LQTS CRD	LQTS CRD	Non-LQTS CRD	Mann-Whitney P-value
Rolling (mo)	6.6	6.0	6.0	7.0	6.2	6.9	0.51
Sitting (mo)	10.1	7.6	10.5	9.0	8.6	11.6	0.05
Walking (mo)	23.4	16.3	39.0	18.5	19.3	28	0.49
Bicycling (y)	5.6	5.0	6.0	5.5	5.3	7.8	n/a
Pincer (mo)	19.5	9.8	7.0	36.0	12.7	25.4	0.19
Utensils (y)	2.4	1.9	2.5	3.3	2.4	2.5	0.95
First word (mo)	22.3	11.4	19.0	14.5	13.8	33.1	0.02
Speak in phrases (y)	3.5	2.0	3.3	4.3	3.0	4.1	0.19
Follow a command (y)	2.1	1.0	1.2	5.0	1.7	2.4	0.16
Smile (mo)	3.3	1.7	2.8	3.5	2.3	4.4	0.71
Imaginative play (y)	2.9	2.9	1.0	4.3	2.8	3.2	0.62
Toilet trained day (y)	4.3	3.7	4.0	6.0	4.1	4.5	0.35
Toilet trained night (y)	5.6	6.0	8.0	5.3	6.1	5.1	0.17

Mean age at which participants achieved developmental milestones for all participants and subsets with and without cardiac arrhythmia. If age of achieving milestone was unknown or participant never achieved an age-appropriate milestone, data were excluded from analysis; of note, for two oldest participants, most data were unknown. Multiple Mann-Whitney test was calculated on the mean age of participants with and without cardiac arrhythmia and P-values reported. Bonferroni multiple comparison correction $P < 0.00167$ for comparisons of development in **Supplemental Tables 1 and 2**.

Abbreviations: CRD: *CACNA1C*-related disorder; LQTS: long QT syndrome; mo: age in months; n/a: not applicable, no value can be calculated due to sample size; TS1: Timothy syndrome type 1; TS2: Timothy syndrome type 2; y: age in years.

Supplemental Table 3: Percent of participants with reported neurologic symptoms.

Symptom (%)	All CRD (n = 24)	TS1 (n = 7)	TS2 (n = 2)	Other LQTS CRD (n = 3)	LQTS CRD (n = 12)	Non-LQTS CRD (n = 12)	Fisher's exact P-value
Any neurologic symptoms	95.8	100	100	100	100	91.7	1
Seizures or epilepsy	37.5	28.6	50	66.7	41.7	33.3	1
Very accurate memory	33.3	28.6	0	66.7	33.3	33.3	1
Language delay	75	57.1	50	100	66.7	83.3	0.64
Language production disorder	50	28.6	50	66.7	41.7	58.3	0.68
Receptive language disability	33.3	14.3	0	33.3	16.7	50	0.19
Learning or intellectual disability	70.8	57.1	50	100	66.7	75	1
Motor delay	91.7	85.7	100	100	91.7	91.7	1
Incoordination	70.8	57.1	50	100	66.7	75	1
Weakness	50	28.6	0	66.7	33.3	66.7	0.22
Hypotonia	66.7	28.6	100	66.7	50	83.3	0.19
Hypertonia, spasticity	20.8	14.3	0	33.3	16.7	25	1
Imbalance	50	14.3	0	66.7	25	75	0.04
Tremors	8.3	14.3	0	0	8.3	8.3	1
Muscle cramps	8.3	14.3	0	0	8.3	8.3	1
Contractures	4.2	0	0	33.3	8.3	0	1
Dysphagia	62.5	71.4	50	33.3	58.3	66.7	1
Abnormal extraocular movements, nystagmus	16.7	0	0	0	0	33.3	0.09
Stereotypies or repetitive movements	45.8	28.6	50	100	50	41.7	1
Echolalia or repetitive vocalizations	50	42.9	50	66.7	50	50	1
Tics	29.2	57.1	0	33.3	41.7	16.7	0.37

Headache	16.7	28.6	0	0	16.7	16.7	1
Sensory loss	8.3	14.3	0	0	8.3	8.3	1
Altered sensitivity to pain and/or temperature	50	42.9	50	66.7	50	50	1
Vision deficit	45.8	28.6	50	66.7	41.7	50	1
Hearing deficit	12.5	14.3	50	33.3	25	0	0.22
Poor sleep quality	50	71.4	50	66.7	66.7	33.3	0.22

Percent of participants with reported neurologic symptoms for all participants and genetic or cardiac subsets. Fisher's exact test two tailed P-value was calculated on the number of participants with and without cardiac arrhythmia. Bonferroni multiple comparison correction $P < 0.0019$.

Abbreviations: CRD: *CACNA1C*-related disorder; LQTS: long QT syndrome; TS1: Timothy syndrome type 1; TS2: Timothy syndrome type 2.

Supplemental Table 4: Percent of participants with reported psychiatric symptoms and diagnoses.

Symptom (%)	All CRD (n = 24)	TS1 (n = 7)	TS2 (n = 2)	Other LQTS CRD (n = 3)	LQTS CRD (n = 12)	Non-LQTS CRD (n = 12)	Fisher's exact P-value
Any psychiatric dx	62.5	71.4	50	33.3	58.3	66.7	1
Any autistic symptoms	91.7	85.7	100	100	91.7	91.7	1
Autism spectrum disorder dx	50	57.1	50	33.3	50	50	1
Short attention span	54.2	57.1	50	66.7	58.3	50	1
Hyperactivity	20.8	28.6	0	0	16.7	25	1
ADHD dx	20.8	28.6	0	0	16.7	25	1
Aggression, self-injurious behavior	29.2	57.1	0	0	33.3	25	1
Specific phobias	25	28.6	0	33.3	25	25	1
Frequent worrying	16.7	28.6	0	33.3	25	8.3	0.59
Anxiety dx	16.7	28.6	0	0	16.7	16.7	1
Social anxiety, agoraphobia	12.5	14.3	0	33.3	16.7	8.3	1
Frequent sadness or depressed mood	4.2	14.3	0	0	8.3	0	1
Depression dx	4.2	14.3	0	0	8.3	0	1
Manic symptoms	4.2	0	0	0	0	8.3	1
Mania or bipolar dx	4.2	14.3	0	0	8.3	0	1
Schizophrenia, psychosis dx	4.2	14.3	0	0	8.3	0	1
Oppositional defiant disorder dx	4.2	14.3	0	0	8.3	0	1
Obsessive compulsive disorder dx	4.2	14.3	0	0	8.3	0	1

Percent of participants who reported experiencing psychiatric symptoms and who reported a formal diagnosis for all participants and genetic or cardiac subsets. Fisher's exact test two tailed P-value was calculated on the number of participants with and without cardiac arrhythmia. Bonferroni multiple comparison correction P <0.0029.

Abbreviations: ADHD: attention deficit hyperactivity disorder; CRD: *CACNA1C*-related disorder; dx: diagnosis (indicates report of a formal diagnosis); LQTS: long QT syndrome; TS1: Timothy syndrome type 1; TS2: Timothy syndrome type 2.