Title: Heterozygous expression of a *Kcnt1* gain-of-function variant has differential effects on SST- and

PV-expressing cortical GABAergic neurons.

Amy N. Shore[1,2], Keyong Li[1], Mona Safari[1,3], Alshaima'a M. Qunies[4,5], Brittany D. Spitznagel[6,7,8], C. David

Weaver[6,7,8], Kyle A. Emmitte[4], Wayne N. Frankel[9,10], and Matthew C. Weston[1,2,3,11*]

[1]Fralin Biomedical Research Institute at Virginia Tech Carilion, Center for Neurobiology Research,

Roanoke, VA, USA.

[2]Department of Neurological Sciences, University of Vermont, Burlington, VT, USA.

[3]Translational Biology, Medicine, and Health Graduate Program, Fralin Biomedical Research Institute at

Virginia Tech Carilion, Roanoke, VA, USA.

[4]Department of Pharmaceutical Sciences, UNT System College of Pharmacy, University of North Texas

Health Science Center, Fort Worth, TX, USA.

[5]School of Biomedical Sciences, University of North Texas Health Science Center, Fort Worth, TX, USA.

[6]Department of Pharmacology, Vanderbilt University, Nashville, TN, USA.

[7]Vanderbilt Institute of Chemical Biology, Vanderbilt University, Nashville, TN, USA.

[8]Department of Chemistry, Vanderbilt University, Nashville, TN, USA.

[9]Institute for Genomic Medicine, Columbia University, New York, NY, USA.

[10]Department of Neurology, Columbia University, New York, NY, USA.

[11]School of Neuroscience, Virginia Tech, Blacksburg, VA, USA.

*Lead contact and correspondence: mcweston7c@vt.edu

Conflict of interest statement: The authors declare no competing financial interests.

ABSTRACT

More than twenty recurrent missense gain-of-function (GOF) mutations have been identified in the

sodium-activated potassium (K_{Na}) channel gene *KCNT1* in patients with severe developmental and

epileptic encephalopathies (DEEs), most of which are resistant to current therapies. Defining the neuron

types most vulnerable to KCNT1 GOF will advance our understanding of disease mechanisms and

provide refined targets for precision therapy efforts. Here, we assessed the effects of heterozygous

expression of a *Kcnt1* GOF variant (Y777H) on K_{Na} currents and neuronal physiology among cortical

glutamatergic and GABAergic neurons in mice, including those expressing vasoactive intestinal

polypeptide (VIP), somatostatin (SST), and parvalbumin (PV), to identify and model the pathogenic

mechanisms of autosomal dominant *KCNT1* GOF variants in DEEs. Although the *Kcnt1*-Y777H variant had

no effects on glutamatergic or VIP neuron function, it increased subthreshold K_{Na} currents in both SST

and PV neurons but with opposite effects on neuronal output; SST neurons became hypoexcitable with a

higher rheobase current and lower action potential (AP) firing frequency, whereas PV neurons became

hyperexcitable with a lower rheobase current and higher AP firing frequency. Further neurophysiological

and computational modeling experiments showed that the differential effects of the Y777H variant on

SST and PV neurons are not likely due to inherent differences in these neuron types, but to an increased

persistent sodium current in PV, but not SST, neurons. The Y777H variant also increased excitatory input

onto, and chemical and electrical synaptic connectivity between, SST neurons. Together, these data

suggest differential pathogenic mechanisms, both direct and compensatory, contribute to disease

phenotypes, and provide a salient example of how a pathogenic ion channel variant can cause opposite

functional effects in closely related neuron subtypes due to interactions with other ionic conductances.

Keywords: epilepsy; DEE; ADNFLE; KCNT1 channel; Slack; K_{Na} current; potassium channel; GOF variant;

glutamatergic; GABAergic; SST; PV; VIP; electrophysiology

INTRODUCTION

Heterozygous missense variants have been identified in the sodium-gated potassium channel

gene *KCNT1* in more than 200 individuals exhibiting a wide spectrum of developmental and epileptic

encephalopathies (DEEs), with the majority being classified as either epilepsy of infancy with migrating

focal seizures (EIMFS) or autosomal dominant or sporadic sleep-related hypermotor epilepsy (ADSHE)

(Barcia et al., 2012; Bonardi et al., 2021; Heron et al., 2012). Each of these epilepsy syndromes result in

early-onset, frequent seizures that are largely pharmacoresistant and often accompanied by a range of

cognitive, psychiatric, and motor impairments. Thus, there is a critical need for a better understanding

of how heterozygous expression of these *KCNT1* variants in the developing brain alters neuronal

physiology and network behavior to lead to such devastating neurodevelopmental disorders.

KCNT1 encodes a tetrameric potassium channel that is widely expressed in both glutamatergic

and GABAergic neurons of the brain, particularly those of the cerebellum, striatum, thalamus,

hippocampus, and cortex (Berg et al., 2007; Bhattacharjee et al., 2002; Gertler et al., 2022; Rizzi et al.,

2016). Although its precise role in normal physiology is not well understood, at least in some neuronal

types, KCNT1 is activated by a persistent inward sodium leak (Na_P) current at rest, where it has a

proposed role in fine tuning neuronal excitability by countering the effects of the Na_P current across

subthreshold voltages (Budelli et al., 2009; Hage & Salkoff, 2012). Consistent with this role, loss-of-

function (LOF) studies using mouse models lacking KCNT1, and the associated sodium-activated

potassium (K_{Na}) current, have shown enhanced AP firing across multiple neuron types (Evely et al., 2017;

Liu et al., 2022; Lu et al., 2015; Martinez-Espinosa et al., 2015; Reijntjes et al., 2019; Zhang et al., 2022).

Characterizations of pathogenic DEE-associated *KCNT1* variants in heterologous cells found that nearly

all cause gain-of-function (GOF) effects on the channel, increasing the associated K_{Na} current (Hinckley et

al., 2023; Kim et al., 2014; McTague et al., 2018; Milligan et al., 2014; Tang et al., 2016). Based on LOF

studies, this would be expected to reduce neuronal excitability; however, it is difficult to predict the

effects of these GOF variants on AP generation in neurons, particularly among neuronal subtypes, *a*

priori.

To address this knowledge gap, we previously generated and characterized a mouse model

expressing a human ADNFLE-associated *KCNT1* GOF variant (Y796H, or Y777H in mice) (Shore et al.,

2020). Although heterozygous expression of the *KCNT1*-Y796H variant is sufficient to cause severe

childhood epilepsy in humans, we only observed rare behavioral seizures in heterozygous *Kcnt1*-Y777H

mice [see Discussion-(Shore et al., 2020)]; however, we identified hyperexcitable, hypersynchronous

cortical networks and frequent, early-onset seizures in homozygous *Kcnt1*-Y777H mice. As a potential

underlying mechanism of these network alterations, we demonstrated that homozygous *Kcnt1*-Y777H

expression increases subthreshold K_{Na} currents and reduces excitability in GABAergic neuron

populations, particularly in those classified as non-fast spiking (NFS), but it does not alter glutamatergic

neuron excitability. We further observed evidence of homeostatic compensation and network

remodeling downstream of K_{Na} current increases during development, including increased excitatory

input onto glutamatergic and NFS GABAergic neurons, and enhanced homotypic synaptic connectivity.

Although these findings provide a strong mechanistic basis for understanding how KCNT1 GOF disrupts

neuronal physiology and network behavior to lead to seizure disorders, key issues remain unresolved.

First, considering the heterozygous nature of *KCNT1* GOF variants in the overwhelming majority of

KCNT1-related DEE patients, it is crucial to determine whether heterozygous *Kcnt1*-Y777H expression

results in similar neuronal impairments and network alterations to those with homozygous expression.

Second, more recent studies, using both *in silico* modeling and additional construct-valid mouse models,

have similarly identified impairments in GABAergic neuron excitability downstream of KCNT1 GOF

(Gertler et al., 2022; Kuchenbuch et al., 2021; Wu et al., 2024), indicating impaired inhibition as a shared

pathogenic mechanism in *KCNT1*-related DEEs; however, precisely which GABAergic subtypes are most

impacted, and how, remains unknown.

Here, we assessed the effects of heterozygous *Kcnt1*-Y777H expression on K_{Na} currents and

neuronal physiology among cortical glutamatergic and GABAergic neurons, including those expressing

vasoactive intestinal polypeptide (VIP), somatostatin (SST), and parvalbumin (PV). Initial assessments of

cortical neuron populations with heterozygous Y777H expression showed strikingly similar effects on K_{Na}

currents and AP generation to those with homozygous expression, although not surprisingly, these

effects were lesser in magnitude. Across all cortical neuron types examined, the heterozygous *Kcnt1*-

Y777H variant caused a range of effects on neuronal excitability and action potential (AP) generation,

from no change (glutamatergic and VIP GABAergic) to decreased excitability (SST GABAergic) to

increased excitability (PV GABAergic). Neuron types that showed no change had K_{Na} currents that were

only significantly increased at suprathreshold voltages and exhibited a steeper voltage dependence of

activation. Interestingly, both SST and PV neurons showed similar increases in K_{Na} currents across

subthreshold voltages, however, only PV neurons had additional increases in the persistent Na^+ current,

which modeling experiments indicated was sufficient to overcome the effects of KCNT1 GOF and cause

an overall increase in AP generation. SST neurons also showed an increase in excitatory input, and in

homotypic electrical and chemical coupling. Taken together, these data provide further evidence of the

enhanced vulnerability of GABAergic neurons, particularly those expressing SST and PV, to KCNT1 GOF.

Moreover, these data show that heterozygous expression of a single *KCNT1* GOF variant can result in a

complex array of neuron type-dependent effects, both direct and indirect, each potential contributors to

the neural circuit pathology underlying *KCNT1*-related DEEs.

RESULTS

Heterozygous *Kcnt1*-Y777H expression alters the shape and frequency of APs in NFS GABAergic

neurons.

We previously identified frequent, early-onset seizures in mice with homozygous expression of

the *Kcnt1* GOF variant Y777H (hereafter referred to as YH-HOM), and as a potential underlying

pathological mechanism, we demonstrated that homozygous *Kcnt1*-Y777H expression drastically impairs

action potential (AP) shape and generation in non-fast spiking (NFS) GABAergic cortical neurons, with

lesser effects on fast spiking (FS) GABAergic, and no significant effects on glutamatergic, cortical neurons

(Shore et al., 2020). Considering that patients with *KCNT1*-associated epilepsy are predominantly

heterozygous for *KCNT1* GOF variants, it is crucial to determine whether heterozygous expression of

these variants, which likely leads to the formation of heteromeric channels consisting of WT and mutant

subunits, results in similar neuronal impairments. To assess neuron subtype-specific effects of

heterozygous KCNT1 GOF on membrane properties and AP firing, we isolated and cultured cortical

neurons from pups with heterozygous Kcnt1-Y777H expression (hereafter referred to as YH-HET), and

their WT littermates, at postnatal day 0 (P0). After infecting the cultured neurons with AAV-*CaMKII*-GFP

to facilitate glutamatergic neuron identification, we performed whole-cell, current-clamp analysis

between 13 and 17 days *in vitro* (DIV). Moreover, to compare to homozygous KCNT1 GOF effects

observed previously, the recorded neurons were classified as glutamatergic, FS GABAergic, or NFS

GABAergic, based on GFP expression, AP parameters, and evoked synaptic responses (see Methods).

Current-clamp recordings from YH-HET and WT glutamatergic neurons showed no significant

differences in any membrane or AP shape property measured (Fig. 1A$_1$ and Supplemental Table 1),

similar to observations from the homozygous KCNT1 GOF studies. Accordingly, AP firing frequencies

across increasing current steps in YH-HET glutamatergic neurons were not altered compared with those

of WT (Fig. 1B$_1$). For FS GABAergic neurons, we previously showed that homozygous Y777H expression

increases the rheobase—the minimal amount of current necessary to induce an AP—and reduces the AP

firing frequency. Heterozygous Y777H expression did not alter the rheobase, or any other passive or

active membrane property of FS GABAergic neurons (Fig. 1A$_2$ and Supplemental Table 1). Although YH-

HET FS GABAergic neurons appeared to fire fewer APs than their WT counterparts, particularly at higher

current steps, this effect was not significant (Fig. 1B$_2$). Lastly, we previously showed that homozygous

Y777H expression has the strongest effects in NFS GABAergic neurons, showing a decrease in input

resistance, accompanied by an increase in rheobase and a reduction in AP firing frequency. YH-HOM NFS

GABAergic neurons also have narrower AP half-widths and larger afterhyperpolarizations (AHPs) than

those of WT. Although heterozygous Y777H expression in NFS GABAergic neurons did not cause a

significant decrease in input resistance (WT: 171 ± 16; YH-HET: 146 ± 14, p=0.19), it did increase the

rheobase and reduce the AP firing frequency, particularly at lower current steps, relative to those of WT

(Fig. 1A$_3$ and 1B$_3$). YH-HET NFS GABAergic neurons also had narrower APs, larger AHPs, and more

depolarized AP thresholds than those of WT (Supplemental Table 1). Together, these data demonstrate

that the neurophysiological effects on cortical neurons with monoallelic expression of the *Kcnt1*-Y777H

variant, expressing channels with mutant and WT subunits, are similar to those with biallelic expression,

expressing only mutant subunits, with both causing the strongest impairments in NFS GABAergic

neurons.

Heterozygous *Kcnt1*-Y777H expression increases subthreshold K$_{Na}$ currents in NFS GABAergic neurons.

Previously, we showed that homozygous *Kcnt1*-Y777H expression in GABAergic cortical neurons

increases the K$_{Na}$ current across subthreshold voltages, an effect that is particularly evident in NFS

GABAergic neurons; conversely, in glutamatergic cortical neurons with homozygous expression of the

same variant, increases in K$_{Na}$ currents are only apparent at depolarized voltages (> +30 mV) (Shore et al.,

2020). To assess the effects of heterozygous Y777H expression on KCNT1 channel function, we

measured the associated K$_{Na}$ current in each cortical neuron subtype. We recorded K$_{Na}$ currents by

applying voltage steps to voltage-clamped neurons and comparing the delayed outward current before

and after the addition of the voltage-gated sodium channel inhibitor TTX (Fig. 2A$_{1-3}$). As reported

previously, there were K_{Na} currents in all three WT neuron subtypes, beginning around -10 mV and

increasing with depolarization (Fig. $2A_{1-3}$ and $2B_{1-3}$), whereas at more negative potentials, the TTX-

sensitive current was net inward (Fig. $2C_{1-3}$) due to the counteracting persistent Na^+ current.

In glutamatergic and NFS GABAergic YH-HET neurons, the overall K_{Na} current was increased

relative to those of WT, as measured by a significant effect of genotype using a linear model (Fig. $2B_{1,3}$).

Importantly, in each of these neuron subtypes, heterozygous Y777H expression increased K_{Na} currents

with distinct, voltage-dependent patterns that appeared strikingly similar to those reported with

homozygous expression. For instance, we previously showed that, across negative potentials, YH-HOM

and WT glutamatergic neuron K_{Na} currents are indistinguishable, whereas at more positive potentials (>

+30 mV), the Y777H variant causes significant increases in K_{Na} currents. Similar voltage-dependent

effects were observed in YH-HET glutamatergic neurons (Fig. $2B_1$ and $2C_1$), although pairwise

comparisons showed that K_{Na} current increases at positive potentials in the YH-HET glutamatergic

neurons were not significant. Conversely, we previously demonstrated broad increases in K_{Na} currents

across negative potentials, with significant increases from -60 to +10 mV, in YH-HOM NFS GABAergic

neurons compared with those of WT, with lesser effects across positive potentials. Similar voltage-

dependent increases were observed in YH-HET NFS GABAergic neurons, with pairwise comparisons

showing significant K_{Na} current increases at -70, -60, -50, and -20 mV (Fig. $2B_3$ and $2C_3$). For both

glutamatergic and NFS GABAergic neurons, the magnitudes of the K_{Na} current increases in YH-HET

neurons were intermediate to those of WT and YH-HOM neurons (Supplemental Fig. 2-1),

demonstrating a gene dose-dependent effect of the *Kcnt1*-Y777H variant on K_{Na} current increases and

validating a GOF effect of the heterozygous Y777H variant on channel function. Lastly, although previous

studies showed that homozygous Y777H expression increases K_{Na} current at several negative voltage

steps (-50, -40, and -10 mV) in FS GABAergic neurons, heterozygous expression of the same variant in FS

GABAergic neurons caused no significant differences in K_{Na} currents compared with those of WT (Fig. $2B_2$

and $2C_2$).

Heterozygous *Kcnt1*-Y777H expression does not alter synaptic connectivity or the excitation-inhibition

balance.

196   In addition to alterations in the intrinsic passive and active membrane properties of cortical

neurons with homozygous Y777H expression, we previously showed evidence of altered synaptic

connectivity and activity, leading to hypersynchronous, hyperexcitable YH-HOM networks; more

specifically, we found an increase in homotypic coupling between glutamatergic [excitatory–excitatory

(E–E)] and GABAergic [inhibitory–inhibitory (I–I)] neuron pairs, and an increase in the frequency of

spontaneous excitatory postsynaptic currents (sEPSCs), accompanied by an increase in the E/I ratio,

onto YH-HOM glutamatergic cortical neurons (Shore et al., 2020). First, to determine whether there are

similar changes in synaptic connectivity in YH-HET networks, we performed paired recordings of

glutamatergic (excitatory, E) and GABAergic (inhibitory, I) neurons and alternatively stimulated each

neuron at 0.1 Hz to test baseline connection probability and strength at the four possible motifs (E–E, I–

E, E–I, and I–I). Connection probability was not altered in YH-HET networks at any of the motifs tested

(Fig. $3A_{1-4}$, left bar graphs), and the amplitudes of the evoked postsynaptic currents (ePSCs) between

connected neurons were not different between genotypes for any of the four connection types (Fig. $3A_{1-}$

$_4$, right bar graphs), indicating grossly normal synaptic interactions among glutamatergic and GABAergic

neurons in YH-HET networks. Next, to assess potential alterations in synaptic activity, we recorded

spontaneous postsynaptic currents (sEPSCs and sIPSCs) onto voltage-clamped glutamatergic and

GABAergic neurons. Unlike the observation of an increase in sEPSC frequency onto YH-HOM

glutamatergic neurons, there was no difference in sEPSC, or sIPSC, frequency onto YH-HET glutamatergic

neurons (Fig. $3B_1$). Furthermore, although we previously found no alterations in sPSC frequency onto

YH-HOM GABAergic neurons, sEPSC and sIPSC frequencies were both slightly reduced onto YH-HET

GABAergic neurons (Fig. 3B$_2$). Finally, to assess the net effect of altered sPSC activity onto YH-HET

neurons, we calculated the E/I ratio, based on the relative frequency and size of the sPSCs, and found no

difference in the E/I ratio onto either YH-HET neuron type (Fig. 3B$_3$). Thus, although heterozygous Y777H

expression resulted in similar K$_{Na}$ current increases and neuronal physiology effects to those found with

homozygous Y777H expression, the broad effects on synaptic connectivity and activity found in YH-HOM

networks were absent in YH-HET networks, which likely results in the observed reduction in seizure

incidence in heterozygous, relative to homozygous, mice.

Heterozygous *Kcnt1*-Y777H expression differentially affects the intrinsic excitability of SST- and PV-

expressing GABAergic neurons.

Next, we sought to determine which GABAergic subtypes are most impacted by heterozygous

Kcnt1-Y777H expression. About 80–90% of cortical GABAergic neurons fall into three, largely non-

overlapping populations that can be identified by their expression of unique markers: the Ca^{2+}-binding

protein parvalbumin (PV; ~40%) and the neuropeptides somatostatin (SST; ~30%) and vasoactive

intestinal polypeptide (VIP; ~15%) (Rudy et al., 2011; Tremblay et al., 2016). The majority of PV-

expressing neurons have been characterized as FS, and VIP-expressing as NFS, whereas SST-expressing

neurons, although largely thought to show NFS firing properties, also contain a population that exhibits

a FS-like phenotype (Large et al., 2016; Ma et al., 2006). *KCNT1* is expressed at higher levels in both

human and mouse cortical GABAergic neurons expressing PV and SST, than in those expressing VIP

(Gertler et al., 2022; Shore et al., 2020). Based on the expression profile of KCNT1, and our findings that

homozygous and heterozygous expression of the *Kcnt1* GOF variant strongly reduced the excitability

GABAergic neurons with NFS firing properties, we hypothesized that the SST-expressing neurons are the

most vulnerable to the effects of KCNT1 GOF. To test this hypothesis, we crossed PV-, SST-, and VIP-Cre

mouse lines to the *Kcnt1*-Y777H mouse line and cultured neurons from the cortices of WT and YH-HET

littermate progeny (Fig. 4A and Supplemental Fig. 4-1A). We infected the cultured neurons with AAV-

CaMKII-GFP, to mark glutamatergic neurons, and AAV-*hSyn*-DIO-mCherry, to mark Cre recombinase-

expressing GABAergic neurons, and performed whole-cell, patch-clamp electrophysiology at DIV 13–17

on GFP$^-$/mCherry$^+$ neurons from each group.

First, we measured the intrinsic membrane properties and AP shape parameters of the three

GABAergic subtypes from the WT control groups to verify that they accurately reflected

electrophysiological behaviors of these subtypes from previous *ex vivo* recordings (Taniguchi et al.,

2011). As expected, the VIP neurons showed a relatively large input resistance, small rheobase, wide AP

half-width, small AHP, and low firing rate (Supplemental Fig. 4-1B and 4-1C, purple); in contrast, PV

neurons showed a small input resistance, large rheobase, narrow AP half-width, large AHP, and high

firing rate (Supplemental Fig. 4-1B and 4-1C, orange). For SST neurons, the values measured for input

resistance, rheobase, AP half-width, AHP, and firing rate were all intermediate to those of VIP and PV

neurons (Supplemental Fig. 4-1B and 4-1C, blue). Moreover, as reported previously, SST neurons

showed the most hyperpolarized AP threshold of the three groups. Together, these data demonstrate

the ability of cortical GABAergic neuron subtypes to retain their characteristic *ex vivo* passive and active

membrane properties *in vitro*.

Next, we assessed the effects of the Y777H variant on the passive and active membrane

properties of each GABAergic subtype. Current-clamp recordings from YH-HET VIP neurons revealed no

significant effect of the variant on any of the membrane or AP properties measured (Fig. 4B$_1$).

Conversely, YH-HET SST neurons showed a strong decrease in input resistance, with an accompanying

increase in rheobase current, relative to those of WT SST neurons (Fig. 4B$_2$), similar to the hypoexcitable

membrane phenotype observed in the YH-HET NFS neurons, without altering AP shape parameters. YH-

HET SST neurons also showed a reduction in the membrane time constant and an increase in membrane

capacitance, which was accompanied by an increase in soma size (Supplemental Table 4 and

Supplemental Fig. 4-2), compared with those of WT SST neurons. Unexpectedly, YH-HET PV neurons

exhibited a decrease in the rheobase current compared with that of WT (Fig. 4B$_3$), and an increase in AP

amplitude (Supplemental Table 4), suggesting the Y777H variant increases PV neuron excitability.

Next, we measured AP firing frequency with incremental, 500-ms current steps in each of the

three GABAergic subtypes. WT neurons showed AP firing frequencies similar to those reported

previously from *ex vivo* recordings of neurons from the three Cre lines (Taniguchi et al., 2011), with VIP

showing the lowest maximal firing frequency, PV the highest, and SST intermediate between VIP and PV

(Fig. 4C$_{1-3}$ and Supplemental Table 4). Consistent with a lack of any effects on membrane and AP shape

properties, the AP firing frequencies across increasing current steps were indistinguishable between WT

and YH-HET VIP neurons (Fig. 4C$_1$). As expected, based on their hypoexcitable membrane properties, the

YH-HET SST neurons fired fewer APs across all current steps relative to WT SST neurons (Fig. 4C$_2$). In

contrast to SST neurons, but consistent with their decreased rheobase current, YH-HET PV neurons fired

more APs across all current steps relative to their WT counterparts (Fig. 4C$_3$). Notably, although these

experiments were performed at room temperature (~22°C), and increasing temperatures to 35°C

altered multiple neuronal physiology parameters (Supplemental Fig. 4-3A), both WT and YH-HET PV

neurons responded similarly to increased temperature (Supplemental Fig. 4-3B), suggesting that the

observed phenotypes are not temperature dependent. Together, these data showed that, as

hypothesized, KCNT1 GOF indeed strongly reduces the excitability of SST neurons, but unexpectedly, it

also causes a hyperexcitable effect in PV neurons; thus, the same ion channel mutation can lead to

opposite effects on excitability in the two largest GABAergic neuron subtypes.

The Y777H variant increases KCNT1-mediated currents across subthreshold voltages in SST- and PV-

expressing GABAergic neurons.

Are the observed differential effects of KCNT1 GOF on GABAergic neuronal physiology due to

distinct patterns of K_{Na} current increases among GABAergic neuron subtypes? To answer this question,

we first recorded K_{Na} currents from GABAergic neurons cultured and labeled as described above, by

applying voltage steps to voltage-clamped neurons and comparing the delayed outward current before

and after the addition of TTX (Supplemental Fig. 5-1A$_{1-3}$). In all three subtypes, for both WT and YH-HET

neurons, we observed K_{Na} currents beginning around -10 mV and increasing with depolarization

(Supplemental Fig. 5-1B$_{1-3}$). For YH-HET VIP neurons, pairwise comparisons to assess voltage-dependent

differences showed that their K_{Na} currents were not different from those of WT at any voltage step

(Supplemental Fig. 5-1B$_1$ and 5-1C$_1$). On the other hand, YH-HET SST neurons showed significant

increases in K_{Na} currents compared with those of WT across multiple subthreshold voltage steps,

including -60, -50, and -40 mV (Supplemental Fig. 5-1B$_2$ and 5-1C$_2$), similar to those observed in NFS

GABAergic neurons with both heterozygous (Fig. 2) and homozygous expression of the Y777H variant

(Shore et al., 2020), indicating a direct current-to-phenotype relationship in YH-HET SST neurons.

Somewhat unexpectedly, pairwise comparisons showed that K_{Na} currents were not different between

YH-HET and WT PV neurons at any voltage step (Supplemental Fig. 5-1B$_3$ and 5-1C$_3$).

The lack of an increase in TTX-sensitive currents in PV neurons, despite alterations in rheobase,

AP amplitude, and AP firing frequency, suggests that either a TTX-insensitive Na^+ source activates KCNT1,

or that compensatory alterations in an opposing current mask an increase in K_{Na} and lead to the changes

in AP firing. In past studies, the lack of selective KCNT1-specific inhibitors necessitated the use of indirect

methods, such as TTX application or Na^+ replacement, to estimate the magnitude of K_{Na}. However, more

recently, we identified and validated a small-molecule, selective KCNT1 channel inhibitor termed

VU0606170, or VU170 (Spitznagel et al., 2020). To further validate the specificity of this small molecule,

we applied voltage steps to voltage-clamped neurons lacking KCNT1 and KCNT2 channels, isolated and

cultured from *Kcnt1*$^{-/-}$; *Kcnt2*$^{-/-}$ double knockout mice (Martinez-Espinosa et al., 2015), and we compared

the delayed outward current before and after the addition of either 10 µM VU170 or 0.5 µM TTX. This

analysis indicated a non-specific outward VU-sensitive current that started around -10 mV and peaked

at +50 mV (~1 nA peak current compared with ~3 nA in WT neurons), which was not present using TTX

subtraction (Supplemental Fig. 5-2A). Across negative potentials, there was an inward TTX-sensitive

current, which again is likely due to the persistent Na^+ current, whereas there was little to no outward

VU-sensitive current (Supplemental Fig. 5-2B), validating the utility of VU170 in isolating K_{Na} currents

across this voltage range.

To better assess KCNT1-mediated currents in GABAergic subpopulations and disentangle

potential confounds of using TTX, particularly across negative potentials, we applied the same voltage

step protocol as in the TTX subtraction experiments but applied 10 µM VU170 instead. In all three WT

GABAergic neuron subtypes, subtraction of the VU170 trace from the control trace revealed nA-sized

outward currents at depolarized potentials, and in YH-HET neurons, the overall VU170-sensitive current

was increased relative to those of WT, as measured by a significant effect of genotype using a linear

model (Fig. $5A_{1-3}$ and $5B_{1-3}$). Pairwise comparisons at each voltage step showed voltage-dependent

differences in current increases among the YH-HET neuron subtypes. For YH-HET VIP neurons, a

significant increase in the VU170 current was only observed at +50 mV (Fig. $5B_1$), whereas subthreshold

currents were indistinguishable from those of WT (Fig. $5C_1$). Previously, we similarly observed K_{Na}

current increases only at more positive potentials in YH-HOM glutamatergic neurons, and like YH-HET

VIP neurons, their membrane and AP properties were unaltered by expression of the YH variant (Shore

et al., 2020). Conversely, in YH-HET SST neurons, significant increases occurred at more negative voltage

steps, from -60 to -30 mV (Fig. $5C_2$). These changes were similar to those observed with TTX treatment,

but slightly larger, possibly due to the lack of the counteracting effect of the persistent Na^+ current,

which is also blocked by TTX. In contrast to the TTX results, YH-HET PV neurons showed an increase in

VU170-sensitive currents, with significant increases from -70 to -50 mV (Fig. $5C_3$). Taken together, these

data identify distinct patterns of K_{Na} current increases among GABAergic subtypes, and for VIP and SST

neurons, these current increases are consistent with the observed effects of KCNT1 GOF on neuronal

physiology. On the other hand, PV neurons showed subthreshold-specific K_{Na} current increases that

were highly similar to, and overlapping with, those of SST neurons; thus, differential K_{Na} current

increases alone likely do not account for the opposite effects of KCNT1 GOF on physiology observed in

these two neuron types.

Compartmental models of KCNT1 GOF in SST, but not PV, neurons are consistent with experimental

data.

The VU170-subtraction experiments showed that YH-HET SST and PV neurons have similar

subthreshold increases in KCNT1-mediated currents, even though they exhibited opposing effects on

neuron excitability and AP generation, and that YH-HET VIP neurons have suprathreshold increases in

KCNT1-mediated currents with no effects on neuron physiology. Because GABAergic subtypes exhibit

varying morphologies and express unique repertoires of ion channels, which are known to give rise to

their characteristic membrane and AP firing properties, we hypothesized that at least some of the

observed differential effects of KCNT1 GOF, in particular the opposite effects in SST and PV neurons, are

due to these inherent, neuron-type-dependent differences. To test this hypothesis, we simulated the

effect of KCNT1 GOF in compartmental models of these three cortical neuron types. We used a KCNT1

conductance with the Na^+ dependence constrained by prior studies in neurons (EC_{50} = 40 mM, slope 3.5;

see Materials and Methods) and the voltage dependence based on the activation curves from our own

experimental data of the VU170-sensitive current in each subtype (Fig. 6A). The kinetics of the model

current were set by measuring the onset time course of the VU170-sensitive current in outside-out

membrane patches of each neuron type (Fig. 6B and 6C). This conductance was inserted into

compartmental models of 10 neurons per subtype, representing a variety of morphological and

electrical properties of each subtype (Markram et al., 2015). Because the Y796H variant was previously

shown to increase the Na^+ sensitivity of the channel, we modeled the GOF effect by reducing the EC_{50} for

Na^+ activation to two levels (35 mM and 30 mM), which resulted in moderate K_{Na} current increases

above WT levels (Fig. 6D).

Consistent with our experimental data, introducing KCNT1 GOF at the lower level (EC_{50} = 35 mM)

into model VIP neurons did not alter their AP firing rate, input resistance, or rheobase, whereas at the

higher level (EC_{50} = 30 mM), there were small changes in input resistance (40 mM: 266.5 ± 28.2; 30 mM:

264.8 ± 28.0, p=0.012) and rheobase (40 mM: 78.7 ± 11.6; 30 mM: 80.4 ± 11.1, p=0.014), but these

changes were not sufficient to affect the AP firing rate (Fig. $6E_1$ and $6E_2$, Supplemental Table 6, and

Supplemental Fig. 6-2A). On the other hand, model SST neurons with KCNT1 GOF at both levels fired

fewer APs per current step as the GOF was increased, and showed significantly reduced input

resistances, higher rheobases, and lower AP firing rates (Fig. $6F_1$ and $6F_2$, Supplemental Table 6), results

that agree with our experimental observations. In contrast to experimental data, but similar to model

SST neurons, model PV neurons responded to KCNT1 GOF with decreased input resistance, increased

rheobase, and a reduced AP firing rate (Fig. $6G_1$ and $6G_2$ and Supplemental Table 6), although the

magnitude of these effects were smaller than those in model SST neurons (Supplemental Fig. 6-2A).

These data suggest that the increased excitability observed in YH-HET PV neurons is not simply due to an

intrinsic property of this GABAergic subtype, but instead results from an indirect mechanism (e.g. a

compensatory response like an increase in Na^+ channels) or depends on a feature of KCNT1 function not

captured by the model.

Lastly, we hypothesized that the lack of GOF effects on VIP and glutamatergic neuron physiology

is due to the altered kinetics of their KCNT1 channel activation curves, in particular their right shifted V_{50}

and decreased slope, relative to those of SST or PV neurons (Fig. 6A). To test this hypothesis, we

performed simulations in which the activation curve parameters (V_{50} and slope) measured in SST

neurons were inserted into VIP and glutamatergic neurons, with the Na$^+$ EC$_{50}$ set to 30 mM to simulate

KCNT1 GOF (Supplemental Fig. 6-1A$_1$ and 6-1B$_1$, blue traces). Inserting KCNT1 channels with SST kinetics

into VIP and glutamatergic neurons indeed enhanced the GOF effects on input resistance and rheobase,

and importantly, resulted in a significantly decreased AP firing rate relative to control neurons for both

neuron types (Supplemental Fig. 6-1A$_2$ and 6-1B$_2$, Supplemental Table 6, and Supplemental Fig. 6-2B).

This model-based evidence suggests that the relative sensitivity of SST and PV neurons, and resistance of

VIP and glutamatergic neurons, to KCNT1 GOF is due to the differences in voltage dependence of the K$_{Na}$

current in each subtype.

The persistent Na$^+$ current is increased by the *Kcnt1*-Y777H variant in PV, but not SST, neurons.

The modeling results suggest that a subthreshold increase in the K$_{Na}$ current alone is not

sufficient to account for the alterations in PV neuron excitability. Moreover, the finding of an increase in

the KCNT1-mediated current in YH-HET PV neurons when measured with VU170, but not TTX, isolation

suggests there may be a compensatory upregulation of an opposing current that masks the increase in

K$_{Na}$ and leads to the unexplained increase in AP firing. The persistent Na$^+$ current (I$_{NaP}$) is active on the

same time scale as K$_{Na}$ and was previously shown to provide the Na$^+$ source for K$_{Na}$ (Hage & Salkoff,

2012); therefore, we hypothesized that KCNT1 GOF causes an increase in I$_{NaP}$ in PV neurons that

enhances their excitability. To test this hypothesis, we measured I$_{NaP}$ in SST and PV neurons by applying

TTX during slow voltage ramp protocols (20 mV/s) designed to isolate I$_{NaP}$ from both the transient Na$^+$

current and the K$_{Na}$ current (Supplemental Fig. 7-1).

In SST neurons, I$_{NaP}$ did not differ between WT and YH-HET neurons, either in the shape of the

response (Fig. 7A and 7B), or the peak amplitude (Fig. 7B). In PV neurons, however, the peak amplitude

of I$_{NaP}$ was significantly increased by the Y777H variant (Fig. 7C and 7D). To examine whether this

increase in I$_{NaP}$ could account for the increase in AP firing observed in PV neurons, we again simulated

the effect of KCNT1 GOF (Na^+ EC_{50} = 35 mM) on PV neuron activity, but this time included an increase in

I_{NaP} in the compartmental models. Indeed, an increase in I_{NaP} conductance, similar to what was seen in

the experimental results (2-fold), was sufficient to increase the number of APs fired in response to

increasing current steps (Fig. 7E and 7F and Supplemental Table 6), even in the face of the higher level of

KCNT1 GOF (30 mM; Supplemental Table 6 and Supplemental Fig. 7-2). Also, like the experimental data,

the rheobase current was decreased (Fig. 7F and Supplemental Fig. 7-2) and the AP height was increased

(80.4 ± 0.7 vs. 82.1 ± 0.8 mV). These results suggest that the differential effects on AP firing in SST and

PV neurons can be accounted for by the absence or presence of a secondary increase in I_{NaP}.

YH-HET SST neurons show increased chemical and electrical coupling and receive increased excitatory

input.

We previously showed that homozygous expression of the Y777H variant increased connections

between GABAergic neurons (Shore et al., 2020), but heterozygous expression did not (Fig. 3). Because

heterozygous Y777H variant expression caused the strongest excitability impairments in SST neurons,

we hypothesized that alterations in synaptic connectivity and activity may be more likely to occur in this

neuronal population. To test this hypothesis, we recorded from pairs of WT or YH-HET SST neurons as

described above to measure connection probability and strength. Indeed, similar to the increase

observed in I–I connections in YH-HOM networks, we found an increase in homotypic synaptic

connections of SST neurons in YH-HET networks (30/40 connections) relative to those in WT (15/40

connections; Fig. 8A). The amplitudes of the evoked postsynaptic currents (ePSCs) between connected

SST neurons were not significantly different between the YH-HET and WT groups (Fig. 8B).

There are two main types of synaptic coupling between neurons: chemical, which is mediated by

neurotransmitter release, and electrical, which is mediated by gap junctions. To test for alterations in

each type of synaptic coupling due to expression of the *Kcnt1* variant, we recorded from 20 pairs of WT

or YH-HET SST neurons in close proximity (< 100 μm apart). Under current-clamp conditions, we injected

a series of hyperpolarizing and depolarizing current steps into one neuron and assessed whether the

current induced a simultaneous voltage deflection in the paired, non-injected neuron (Fig.8C_1; 100-pA-

evoked response). We then evoked a train of APs in one neuron and assessed the AP-induced voltage

changes in the paired, non-injected neuron (Fig. 8C_1; AP-evoked response). SST pairs were considered to

be chemically coupled if the AP trains in one neuron induced corresponding IPSCs, without inducing

voltage deflections in response to the current steps, onto the paired neuron (Fig. 8C_1; top panels). SST

pairs were considered to be electrically coupled if the current steps injected into one neuron induced

simultaneous voltage deflections in the paired neuron (Fig. 8C_1; bottom panel, left). Electrically coupled

SST neurons also frequently showed AP-evoked spikelets onto paired neurons due to low-pass filtering

(Fig. 8C_1; bottom panel, right), which results in a greater attenuation of the high frequency portion

(spike), than the low frequency portion (AHP), of an action potential as it passes through an electrical

synapse.

Of the 20 WT SST neuron pairs tested, 12 were coupled, and the majority (9/12) of those were

one-way chemical connections (Fig. 8C_2). Of the 20 YH-HET SST neuron pairs tested, 17 were coupled,

and the majority (13/17) of those were two-way, or bidirectionally, connected (Fig. 8C_2). Of the 3

bidirectionally connected WT SST neuron pairs, 2 were chemical and 1 was electrical, whereas of the 13

bidirectionally connected YH-HET SST neuron pairs, 7 were chemical and 6 were electrical (Fig. 8C_2).

Thus, these data suggest that heterozygous Y777H expression increases reciprocal connectivity among

SST neurons, of both the chemical and electrical sort.

Finally, although we previously found no alterations in the sEPSC frequency onto the total

population of YH-HOM GABAergic neurons, there was an increase in the sEPSC frequency onto those

with an NFS phenotype (Shore et al., 2020), suggesting there may be a compensatory increase in

excitatory drive onto NFS YH-HOM GABAergic neurons to offset the effects of their decreased

membrane excitability. To assess whether there is a similar increase in sEPSC frequency onto SST-

expressing YH-HET neurons, which showed a similar decrease in membrane excitability to that of NFS

YH-HOM GABAergic neurons, we recorded sEPSCs onto voltage-clamped WT and YH-HET SST neurons.

Indeed, there was an increase in sEPSC frequency onto YH-HET SST neurons compared with that of WT

(Fig. 8D), suggesting a compensatory increase in excitatory drive onto the most impaired GABAergic

subtype.

Acute cortical slice recordings confirm the reduced intrinsic excitability of YH-HET SST neurons.

Of the three GABAergic neuron populations assessed, SST-expressing neurons showed the

strongest impairments, both direct and indirect, downstream of the heterozygous Y777H variant.

Importantly, the observed effects on YH-HET SST neuron physiology were consistent with those that

would be expected, based on YH-HET SST subthreshold K_{Na} current increases (Supplemental Fig. $5C_2$, and

Fig. $5C_2$), and predicted, based on SST KCNT1 GOF neuron modeling data (Fig. $6F_1$ and $6F_2$). Thus, to test

whether the most significant physiological alterations identified in cortical neuron cultures are also

present in an experimental preparation with a more organized network, we crossed YH HET mice with

homozygous GIN (GFP-expressing Inhibitory Neurons) mice, which express GFP in a subpopulation of

SST-expressing neurons of the hippocampus and cortex (Ma et al., 2006; Oliva et al., 2000), and

performed electrophysiological recordings in acute brain slices from WT and YH-HET progeny.

First, we assessed the effects of the heterozygous Y777H variant on the passive and active

membrane properties by performing whole-cell, current-clamp recordings of layer 2/3, GFP-expressing

neurons in motor and somatosensory cortical regions of slices prepared from P21–P30 mice (Fig. 9A).

Incremental current injections to elicit APs demonstrated a strong impairment in AP generation

downstream of the heterozygous Y777H variant, as the YH-HET GFP-expressing neurons fired fewer APs

across all current steps than the WT GFP-expressing neurons, similar to that observed in the YH-HET SST

cultured neurons. (Fig. 9B and 9C). Although YH-HET GFP-expressing neurons in slice did not show a

significant decrease in input resistance relative to those from WT, they did show a large increase in

rheobase current, without alterations in AP shape (Fig. 9D). Thus, we confirmed the main

electrophysiological finding from culture—the reduced excitability of YH-HET SST neurons—in acute

slices from YH-HET juvenile mice.

DISCUSSION

More than a decade ago, autosomal dominant mutations were identified in the sodium-gated

potassium channel *KCNT1* in multiple patients with severe, childhood epilepsy syndromes (Barcia et al.,

2012; Heron et al., 2012). *KCNT1*-related epilepsy patients suffer not only with frequent, early-onset

seizures, but also with cognitive impairments, movement disorders, and sometimes additional

behavioral and/or psychiatric problems (Bonardi et al., 2021). Unfortunately, after over 10 years of

research efforts, *KCNT1*-related epilepsies remain highly refractory to current anti-seizure medications

(Bonardi et al., 2021); thus, a better understanding of the pathological mechanisms underlying *KCNT1*-

related epilepsies—from mutated gene to altered K^+ current to altered neuronal physiology to altered

neural network to seizure—is imperative for advancing therapeutic strategies to improve seizure control

and enhance patient quality of life.

Heterozygous vs. homozygous *KCNT1* GOF variant effects

Despite the heterozygous nature of *KCNT1* GOF variants in the overwhelming majority of *KCNT1*-

related DEE patients, research efforts, including our own, have largely focused on homozygous GOF

variant effects on channel function and neuronal physiology (Gertler et al., 2019; Kim et al., 2014;

McTague et al., 2018; Mikati et al., 2015; Milligan et al., 2014; Quraishi et al., 2019; Shore et al., 2020;

Tang et al., 2016). Heterozygous *KCNT1* GOF variant expression likely results in the formation of

heteromeric KCNT1 channels, consisting of both WT and mutant subunits, whose characteristics and

kinetics may differ from those of homomeric KCNT1 channels; thus, it is critical to distinguish between

GOF variant effects in heteromeric and homomeric KCNT1 channels to identify the pathogenic

mechanisms of *KCNT1* GOF variants in DEEs. Recent work in heterologous cells has shown that KCNT1

GOF heteromeric channels, with WT and mutant subunits, exhibit K_{Na} current increases that lie between

those of WT and KCNT1 GOF homomeric channels (Cole et al., 2021; Dilena et al., 2018; Rizzo et al.,

2016). Furthermore, other recent studies have similarly shown that heterozygous *Kcnt1* GOF variant

expression in neurons causes effects on K_{Na} current and excitability that are intermediate to those of WT

and homozygous variant-expressing neurons (Gertler et al., 2022; Wu et al., 2024).

Here, we assessed the effects of heterozygous expression of *Kcnt1*-Y777H on K_{Na} currents,

neuronal physiology, and synaptic connectivity among cortical neuron subtypes in mice, and compared

the effects to those observed previously with homozygous expression of the same variant (Shore et al.,

2020). We found that the heterozygous YH variant caused similar effects on K_{Na} currents and neuronal

physiology to those caused by the homozygous YH variant, including subthreshold-specific increases in

K_{Na} current, altered AP shape, and impaired AP firing of NFS GABAergic neurons, but with magnitudes

intermediate to those of WT and YH-HOM neurons. At the network and whole-animal level, we

previously showed that homozygous expression of the YH variant in mice increases homotypic synaptic

connectivity (E–E and I–I) and the E/I ratio, resulting in hypersynchronous, hyperexcitable YH-HOM

cortical networks and frequent seizures. Here, we found no evidence of similar network alterations in

YH-HET cortical networks, which likely explains the infrequent seizures observed in the heterozygous YH

mouse model, and together with the YH-HOM data, indicate a strong relationship between network

abnormalities and seizure propensity downstream of the YH variant. Furthermore, these data suggest

that heterozygous and homozygous *KCNT1* variant expression cause similar, gene-dose-dependent GOF

effects at the current and neuron level, but not at the network or whole-animal level.

Neuron-type-dependent *KCNT1* GOF variant effects

Prior work has illustrated the importance of cell context in delineating the functional effects of ion channel variants on neuronal physiology. For instance, several studies using LOF channelopathy models of both sodium and potassium channels have identified excitability effects on one neuron type without effects on another (Hedrich et al., 2014; Soh et al., 2018; Tai et al., 2014; Yu et al., 2006). Other studies have shown that the same ion channel variant can even cause opposite effects on excitability in different neuron types (Makinson et al., 2016; Rush et al., 2006; Studtmann et al., 2022). Thus, understanding how KCNT GOF alters networks to lead to hyperexcitability and seizures requires a thorough evaluation of how the variant impacts each neuron type that participates in the network. With this goal in mind, we assessed the effects of heterozygous *Kcnt1* GOF on K_{Na} currents and AP generation among cortical glutamatergic neurons and all three major GABAergic neuron subtypes, including those expressing VIP, SST, and PV.

Using two methods to isolate and measure the KCNT1-mediated current, VU170- and TTX-subtraction, we observed voltage-dependent differences in current increases among the YH-HET neuron types. YH-HET glutamatergic and VIP neurons showed KCNT1-mediated current increases across suprathreshold voltages, whereas YH-HET SST and PV neurons had current increases across subthreshold voltages. These neuron-type dependent effects of KCNT1 GOF on K_{Na} current increases are similar to those observed previously in YH-HOM, glutamatergic and GABAergic neuron subpopulations using TTX (Shore et al., 2020). Moreover, these results are similar to those reported in a recent study using a different KCNT1 GOF mouse model (human R474H, mouse R455H), which showed K_{Na} current increases selectively across positive potentials in glutamatergic, but across positive and negative potentials in FS and NFS GABAergic, neurons (Wu et al., 2024). The mechanism underlying these neuron-type-dependent KCNT1 GOF effects on K_{Na} current increases is unknown, however, we have observed

differences even in WT KCNT1 channel activation curves between glutamatergic and VIP neurons

(steeper slopes of activation) and SST and PV neurons (shallower slopes of activation), suggesting there

are factors affecting KCNT1 channel kinetics that are inherent to each neuron type (Fig. 6A). Future

studies should investigate these potential factors including (1) differential expression of alternative

KCNT1 splice forms, some of which are known to have different activation kinetics, (2) differential

expression of other channels such as KCNT2, which can form heteromers with KCNT1 and alter its

biophysical properties, and (3) differential expression and/or localization of, and/or coupling to, sodium

channels that act as the source of KCNT1 channel activation (Chen et al., 2009; Hage & Salkoff, 2012;

Joiner et al., 1998).

Consistent with the differential KCNT1 GOF effects on K_{Na} currents, we observed a range of

effects on membrane properties and AP generation among the cortical neuron populations. We

observed no effects on YH-HET glutamatergic and VIP neurons, whereas YH-HET SST neurons showed

reduced AP firing, with decreased input resistance and increased rheobase, and YH-HET PV neurons

showed increased AP firing, with decreased rheobase. As a complementary approach, we performed *in

silico* electrophysiology using our measured neuron-type-specific KCNT1 channel activation kinetics, with

and without GOF, for each neuron type. For glutamatergic and VIP neurons, the steeper slopes of

activation limited the effects of GOF on intrinsic excitability, whereas for SST neurons, the shallower

slope of activation decreased the input resistance, increased the rheobase, and impaired AP generation

downstream of KCNT1 GOF. Each of these modeling results are similar to those observed using *in vitro*

electrophysiology, indicating direct K_{Na} current-to-KCNT1 GOF phenotype relationships in these neuron

types.

Conversely, for PV neurons, unlike the experimental observation of increased YH-HET PV neuron

excitability, *in silico* electrophysiology predicted a slight reduction in PV neuron excitability downstream

of KCNT1 GOF. In support of this *in silico* prediction, in a different KCNT1 GOF mouse model,

heterozygous expression of the human L456F variant indeed reduced AP firing of PV neurons, without

altering any other membrane properties or AP parameters (Gertler et al., 2022). The opposite effect of

KCNT1 GOF on PV excitability between these two studies may simply result from variant-specific

alterations in KCNT1 channel kinetics; however, both GOF variants showed a similar lack of effect on

membrane properties and AP generation of glutamatergic neurons. Alternatively, the opposite effects

could be due to the different experimental conditions used in the two studies, including (1) different

preps and mouse ages (acute slices from P42–P60 mice vs. DIV 13–17 cultured neurons from P0 pups)

and (2) different brain regions (hippocampal vs. cortical). The idea of brain region-dependent effects of

ion channel variants is not unprecedented, as a recent study showed that a potassium channel GOF

variant causes opposite effects on excitability in the same neuron type from different brain regions

(hippocampal CA1 vs. cortical layer 2/3 pyramidal neurons), and this opposite effect was suggested to

be due to the compensatory upregulation of sodium channels in one neuron type but not the other

(Varghese et al., 2023). Taken together, these findings illustrate the importance of interrogating not only

multiple neuron types but also multiple brain regions to fully unravel the complexity underlying KCNT1

GOF-related neurodevelopmental disorders.

Evidence of potential neuron and network compensatory responses to KCNT1 GOF

In a healthy brain, there are multiple homeostatic compensation mechanisms to maintain

neuronal and network activity at their proper physiological levels (Davis, 2013; Debanne et al., 2019;

Debanne & Russier, 2019; Marder & Goaillard, 2006; Turrigiano, 2011, 2012; Yang & Prescott, 2023),

whereas in a brain with epilepsy-causing genetic mutations or injury, it is thought that failure of

homeostatic compensation can result in hypersynchronous, hyperexcitable networks and, ultimately,

seizures (Issa et al., 2023; Lignani et al., 2020; Staley, 2015). As YH-HET mice have only been observed to

have infrequent seizures and YH-HET networks appear largely intact, it is plausible that there are

adaptive or homeostatic mechanisms downstream of the YH variant that limit KCNT1 GOF effects largely

to those observed at the current and neuron level. These regulatory mechanisms may include

alterations in ion currents and/or synaptic input to restore proper AP shape or neuronal excitability, or

alterations in chemical and/or electrical connectivity to stabilize neural networks.

Neuronal excitability is regulated by the inward and outward flow of opposing currents through

sodium and potassium channels; thus, it is not surprising that mutations affecting these channels can

lead to compensatory up- or down-regulation of other ion channels or currents to maintain excitability

and proper neuron function. In this study, our *in silico* electrophysiology predictions of YH-HET PV

neuron excitability were opposite to those of our *in vitro* electrophysiology data; thus, we hypothesized

that there may be a compensatory increase in an ion current that counteracts the increased K_{Na} current

selectively in YH-HET PV neurons. Accordingly, we identified a subthreshold increase in the persistent

Na^+ current, across a similar voltage range to the increased K_{Na} current, in YH-HET PV, but not SST,

neurons. Furthermore, *in silico* approaches indicated that this increased I_{NaP} was sufficient to overcome

the effects of KCNT1 GOF and cause an overall increase in AP generation in YH-HET PV neurons, which

aligned with our experimental data. Similar compensatory mechanisms have been identified in other

models such as the downregulation of multiple potassium channels in a sodium channel (*Scn2a*) LOF

mouse model of epilepsy, resulting in neuronal hyperexcitability (Zhang et al., 2021), or the upregulation

of Na^+ currents in response to increased potassium channel activity in a developing Xenopus neural

circuit, resulting in restored intrinsic excitability and network stability (Pratt & Aizenman, 2007). Notably,

another KCNT1 GOF model (*Kcnt1*-R455H) showed upregulation of a sodium channel ($Na_v1.6$),

accompanied by an increase in transient and persistent sodium currents, in *Kcnt1*-R455H cortical

neurons, but the effects of Nav1.6 upregulation were not investigated (Wu et al., 2024).

In addition to the KCNT1 GOF current-mediated impairments in SST neuron excitability, we

identified several potential compensatory YH-HET SST synaptic and network alterations. For instance,

there was an increased frequency of sEPSCs onto YH-HET SST neurons, suggestive of a compensatory

increase in excitatory input to counteract their reduced intrinsic excitability and increase their AP firing,

as has been observed previously in hypoexcitable SST neurons in another epilepsy model (Halabisky et

al., 2010). There was also an increase in chemical and electrical coupling between YH-HET SST neurons.

Although SST neurons normally show a relatively low rate of homotypic chemical coupling, particularly

in the adult cortex, more than half of nearby SST neuron pairs are coupled electrically (Amitai et al.,

2002; Gibson et al., 1999, 2005; Hu & Agmon, 2015; Urban-Ciecko & Barth, 2016). This coupling is

thought to be important for fine tuning neural circuits by regulating such processes as synchronous AP

firing, synaptic integration, and network rhythmicity (Alcami & Pereda, 2019; Connors, 2017). The

observed increase in electrical coupling should not only contribute to the reduced intrinsic excitability

properties of individual YH-HET SST neurons, increasing their capacitance and decreasing their input

resistance, but may also act as a homeostatic mechanism to restore synchronicity and stability to YH-

HET SST networks (Lane et al., 2016; Parker et al., 2009).

Taken together, how might the YH-HET variant-induced alterations to SST and PV neurons, both

direct and adaptive, ultimately contribute to neural network behavior and seizure propensity in YH-HET

mice? The paucity of effects on synaptic activity and connectivity in YH-HET networks, relative to those

of YH-HOM, suggests that the current- and neuron-level effects of the YH-HET variant are below some

threshold required to induce broader network hyperexcitability and seizures. Moreover, as described

above, several of the proposed homeostatic mechanisms, such as increased excitatory input onto, and

connectivity between, SST neurons and increased I_{NaP} in PV neurons, should theoretically act to suppress

pathological network activity in YH HET mice. However, predicting the downstream effects of altered

SST and PV intrinsic excitability is not straightforward, particularly with the emerging appreciation of

both anti- and pro-epileptic roles for these neuron types in seizure initiation and maintenance (Dudok et

al., 2022; Magloire et al., 2019). Furthermore, studies using genetic mouse models of DEEs have

identified seemingly paradoxical, and sometimes ephemeral, roles of interneurons in disease

progression. For instance, in an *SCN8A* GOF DEE mouse model, increased I_{NaP} causes initial SST

hyperexcitability followed by depolarization block, and in parallel, chemogenetic activation of WT SST

neurons mimics these effects, leading to seizure generation (Wengert et al., 2021). Similarly, in a

KCNQ2/3 LOF DEE model, PV-specific *Kcnq2/3* deletion causes early PV hyperexcitability that is thought

to increase seizure susceptibility via homeostatic potentiation of excitatory transmission (Soh et al.,

2018). Lastly, in an *Scn1a*$^{+/-}$ mouse model of Dravet syndrome, PV neurons are transiently hypoexcitable

during development, whereas later they show an increased firing rate, both interictally and at seizure

onset, and a decrease in synchrony in the transition to seizure (Favero et al., 2018; Tran et al., 2020).

These studies illustrate the challenges in connecting altered interneuron intrinsic excitability with

network behavioral outcomes not only because of the contradictory nature of some of these

connections but also because of the often transient nature of interneuron phenotypes during disease

progression. Thus, future studies should aim to assess neuron-type-specific effects of the YH variant at

different developmental time points in both YH-HET mice, with rare seizures, and YH-HOM mice, with

frequent seizures, to better understand the contributions of interneuron dysfunction to network

pathology and seizure generation in KCNT1 GOF-related epilepsy.

Limitations and Conclusions

The *in vitro* and *in silico* nature of this study are limitations. Although the cultured VIP, SST, and

PV neurons showed characteristic electrophysiological properties, it is unknown how much overlap

there is between the *in vitro* and *in vivo* populations. For instance, it is possible that our culture

conditions select for more mature neurons of each type or preferentially support the survival of

subpopulations; thus, we may be assessing KCNT1 GOF effects on a small portion of a given subtype.

However, an advantage of the *in vitro* prep is that the compensatory alterations we observe are not

likely consequences of seizure activity, which has been shown to alter interneuron properties in epilepsy

models. Furthermore, the *in silico* neurons we used to model each neuron type were designed to

accurately reflect the morphological and electrophysiological properties of each subtype, but they may

lack more detailed features of ion channels, such as post-translational modifications and subcellular

localizations, that can have important functional effects. Our KCNT1 model conductance is also

hampered by an incomplete understanding of the relationship between Na^+ influx, membrane voltage,

and channel gating in neurons.

Moreover, our K^+ current measures using *Kcnt1$^{-/-}$*; *Kcnt2$^{-/-}$* double knockout neurons clearly

showed the presence of an outward VU170-sensitive current across positive potentials, indicating non-

specific activity of this small molecule, potentially to the ERG K^+ channel (Spitznagel et al., 2020). This

off-target effect of VU170 may contribute to the overall increase in outward K^+ currents seen in our

VU170- vs TTX-subtracted currents in all neuron types across positive potentials, and it limits the

interpretation of the VU170-sensitive K_{Na} current data within this range. Finally, although it is well

known that temperature changes can alter neuron physiology, the electrophysiological recordings in this

study were performed at room temperature rather than physiological temperature. In this study, we

further assessed the effect of temperature (22°C vs. 35°C) on rheobase, AP amplitude, and maximal AP

firing frequency among WT and YH-HET PV neurons and showed that both groups responded similarly.

Previously, we also showed that the effects of the homozygous YH variant on glutamatergic and NFS

GABAergic neuron excitability were highly similar between cultured recordings at room temperature

(~22°C) and slice recordings at 32°C. Taken together, these data suggest that the reported

neurophysiological phenotypes downstream of the YH variant are likely not temperature dependent.

Despite the potential limitations, this study provides three major findings that advance our

understanding of the relationship between ion channel function and disease. First, heterozygous *KCNT1*

expression causes changes in current flow and neuron excitability that are qualitatively similar to

homozygous expression, but of a lower magnitude; however, the synaptic activity and connectivity

changes are different, which likely leads to the discrepancy in seizure incidence. This finding suggests a

non-linear relationship between GOF, at the level of the ion channel current, and disease severity.

Second, the same *KCNT1* variant can produce opposite effects on neuron excitability in closely related

GABAergic neuron subtypes, and these opposite effects are likely due to compensation in one neuron

type (PV) that is absent in the other (SST). Previous studies have observed opposite effects on neuronal

excitability due to Na$^+$ channel variants, but the underlying mechanism, when explored, was proposed to

be intrinsic differences in ion channel expression, not compensation. Third, we observed increased

synaptic and gap junction connectivity among SST neurons, demonstrating that the effects of KCNT1

GOF extend to structural alterations as well. The effects of this increased connectivity, especially by gap

junctions, on neuronal excitability and network behavior offers an exciting avenue for future research.

Finally, it will be critical to determine whether these alterations downstream of KCNT1 GOF potentiate

or attenuate network pathology and seizure activity, thus providing a better understanding of disease

mechanisms and prompting novel therapeutic design.

MATERIALS AND METHODS

Mice

Mice were bred, and mouse procedures were conducted, in compliance with the National Institutes of

Health (NIH) Guidelines for the Care and Use of Laboratory Animals and were approved by the

Institutional Animal Care and Use Committee at the University of Vermont (animal protocol numbers:

16-001, 19-034, and X9-022). Mice were maintained in ventilated cages at controlled temperature (22–

23°C), humidity ~60%, and 12-h light: 12-h dark cycles (lights on at 7:00 AM, off 7:00 PM). Mice had

access to regular chow and water, *ad libitum*. For all experiments, male and female littermates were

used for each genotype. The ages of the mice for each experiment are indicated in the following

relevant sections.

Mouse lines and strains used for these studies include: *Kcnt1*^{Y777H} knockin in the C57BL/6NJ (B6NJ) strain

(formal gene and allele symbol: *Kcnt1*^{em1(Y777H)Frk}), *Sst*-IRES-Cre (*Sst*^{tm2.1(cre)Zjh}/J; Jackson Labs stock:

013044), *Vip*-IRES-Cre (*Vip*^{tm1(cre)Zjh}/J; Jackson Labs stock: 010908), *Pvalb*-IRES-Cre (B6.129P2-

Pvalb^{tm1(cre)Arbr}/J; Jackson Lab stock: 017320), C57Bl/6J (Jackson Labs stock: 000664), and GIN (FVB-

Tg(GadGFP)45704Swn/J; Jackson Lab stock: 003718) mice.

Kcnt1^{Y777H} mice were genotyped using PCR amplification primers (*Kcnt1* forward primer: 5'-

CTAGGGCTGCAAACACAACA-3'; *Kcnt1* reverse primer: 5'-TCAAGCAGCAACACGATAGG-3') with standard

thermocycler amplification conditions, and the annealing temperature set at 58°C. Following

amplification, a restriction cut was performed with the enzyme *Nla*III to distinguish mutant (127 and 44

bp products after cut) from wildtype (WT) alleles (171 bp product). Progeny of the *Kcnt1*^{Y777H} mice

crossed to each Cre mouse line were genotyped for the presence of a *Cre* transgene using a forward

primer (5'-TCGCGATTATCT TCTATA TCTTCAG-3') and reverse primer (5'-GCTCGACCAGTTTAGTTACCC-3'),

resulting in a 455 bp product.

Primary Astrocyte Feeder Layer Culture

Astrocyte feeder layers were generated to support the growth and maintenance of primary neurons.

Briefly, cortices were dissected from P0–P1 WT C57BL/6J mice (Jackson Labs stock: 000664) of either sex.

The cortices were incubated in 0.05% trypsin-EDTA (Gibco) for 15 min at 37°C in a Thermomixer

(Eppendorf) with gentle agitation (800 rpm). Then, the cortices were mechanically dissociated with a 1-

mL pipette tip, and the cells were plated into T-75 flasks containing filter-sterilized astrocyte media

[DMEM media supplemented with glutamine (Gibco), 10% fetal bovine serum (FBS, GE Healthcare), 1X

MITO+ Serum Extender (Corning), and 0.2X penicillin/streptomycin (Gibco)]. After the astrocytes

reached confluency, they were washed with PBS (Gibco) and incubated for 5 min in 0.05% trypsin-EDTA

at 37°C, washed, and then resuspended in astrocyte media. Astrocytes were added to 6-well plates

containing 25-mm coverslips precoated with coating mixture [0.7 mg/ml collagen I (Corning) and 0.1

mg/ml poly-D-lysine (Sigma) in 10 mM acetic acid].

Primary Cortical Neuron Culture

For the primary neuron culture, the dorsomedial cortices from P0–P1 WT and *Kcnt1*$^{m/+}$ mice of both

sexes were dissected in cold HBSS (Gibco). The tissue was then digested with papain (Worthington) for

60–75 min and treated with inactivating solution (Worthington) for 10 min, both while shaking at 800

rpm at 37°C in a Thermomixer. The cells were then mechanically dissociated and counted. The

dissociated cells were added at 200,000 cells/well to 6-well plates containing astrocyte-coated

coverslips in filter-sterilized NBA plus [Neurobasal-A medium (Gibco) supplemented with 1X Glutamax

(Gibco), 1X B27 (Invitrogen), and 0.2X penicillin/streptomycin (Gibco)]. After plating (12-24 h),

approximately 4×10^{10} genome copies (GC) of AAV8-*CaMKII*-GFP (UNC Vector Core) were added to each

well. For experiments to assess KCNT1 effects on GABAergic subtypes, approximately 4×10^{10} genome

copies (GC) of AAV9-*hSyn*-DIO-mCherry (addgene) were also added to each well to mark Cre-expressing

neurons. Every 3–4 days, 20–40% of the media was replaced with fresh NBA plus.

Electrophysiology

Whole-cell recordings were performed with patch-clamp amplifiers (MultiClamp 700B; Molecular

Devices) under the control of Clampex 10.3 or 10.5 (Molecular Devices, pClamp, RRID:SCR_011323).

Data were acquired at 20 kHz and low-pass filtered at 6 kHz. The series resistance was compensated at

70%, and only cells with series resistances maintained at less than 15 MΩ were analyzed. Patch

electrodes were pulled from 1.5-mm o.d. thin-walled glass capillaries (Sutter Instruments) in five stages

on a micropipette puller (model P-97; Sutter Instruments). Internal solution contained the following: 136

mM K-gluconate, 17.8 mM HEPES, 1 mM EGTA, 0.6 mM $MgCl_2$, 4 mM ATP, 0.3 mM GTP, 12 mM creatine

phosphate, and 50 U/ml phosphocreatine kinase. Alternatively, internal solution contained: 136 mm KCl,

17.8 mm HEPES, 1 mm EGTA, 0.6 mm MgCl2, 4 mm ATP, 0.3 mm GTP, 12 mm creatine phosphate, and

50 U/ml phosphocreatine kinase. The pipette resistance was between 2 and 4 MΩ. Standard

extracellular solution contained the following (in mM): 140 NaCl, 2.4 KCl, 10 HEPES, 10 glucose, 4 $MgCl_2$,

and 2 $CaCl_2$ (pH 7.3, 305 mOsm). All experiments were performed at room temperature (22–23°C), with

the exception of the current-clamp experiments to compare the effects of temperature on neuronal

physiology (Supplemental Figure 4-3). For these recordings, all current-clamp parameters were

measured first at room temperature, then the temperature of the external solution was heated to 35°C

using a ThermoClamp Inline Heater (ThermoClamp-3; AutoMate Scientific), and the measurements were

repeated. Whole-cell recordings were performed on cortical neurons from control and mutant groups in

parallel on the same day (day 13–17 *in vitro*). All experiments were performed by two independent

investigators blinded to the genotypes.

For current-clamp experiments, the intrinsic electrophysiological properties of neurons were tested by

injecting 500-ms square current pulses incrementing in 20 pA steps, starting at −100 pA. Resting

membrane potential (V_m) was calculated from a 50 ms average before current injection. The membrane

time constant (τ) was calculated from an exponential fit of current stimulus offset. Input resistance (R_{In})

was calculated from the steady state of the voltage responses to the hyperpolarizing current steps.

Membrane capacitance was calculated by dividing the time constant by the input resistance. Action

potentials (APs) were evoked with 0.5 s, 20 pA depolarizing current steps. Rheobase was defined as the

minimum current required to evoke an AP during the 500 ms of sustained somatic current injections. AP

threshold was defined as the membrane potential at the inflection point of the rising phase of the AP.

AP amplitude was defined as the difference in membrane potential between the AP peak and threshold,

and the afterhyperpolarization (AHP) was the difference between the AP threshold and the lowest V_m

value within 50 ms. The AP half-width was defined as the width of the AP at half-maximal amplitude. To

obtain the neuron's maximum firing frequency, depolarizing currents in 20-pA steps were injected until

the number of APs per stimulus reached a plateau phase. The membrane potential values were not

corrected for the liquid junction potential. GABAergic neurons were classified as fast spiking (FS) if their

maximum mean firing rate reached above 60 Hz and their AP half-widths increased by less than 25%

during 1 s of sustained firing (Avermann et al., 2012; Casale et al., 2015). All others were considered

non-fast spiking (NFS).

For voltage-clamp experiments to measure synaptic currents, neurons were held at -70 mV, except for

evoked IPSC measurements, for which neurons were held at 0 mV. AP-evoked EPSCs were triggered by a

2 ms somatic depolarization to 0 mV. The shape of the evoked response, the reversal potential and the

effect of receptor antagonists [10 µM NBQX (Tocris Bioscience) or 20 µM bicuculline (BIC, Tocris

Bioscience)] were analyzed to verify the glutamatergic or GABAergic identities of the currents. Neurons

were stimulated at 0.1 Hz in standard external solution to measure basal-evoked synaptic responses.

Electrophysiology data were analyzed offline with AxoGraph X software (AxoGraph Scientific,

RRID:SCR_014284). Spontaneous synaptic potentials were recorded in control solution with either NBQX

or bicuculline to isolate EPSCs or IPSCs, respectively. Data were filtered at 1 kHz and analyzed using

template-based miniature event detection algorithms implemented in the AxoGraph X. The threshold

for detection was set at three times the baseline SD from a template of 0.5 ms rise time and 3 ms decay.

The E/I ratio was calculated as the product of the sEPSC frequency and charge over the sum of the sEPSC

frequency and charge and the product of the sIPSC frequency and charge.

K_{Na} and KCNT1 Current Measurements

For voltage-clamp experiments to measure the sodium-activated K^+ current or KCNT1-mediated current,

neurons were held at -70 mV and given 1 s voltage pulses in 10 mV steps over a range of -80 to +50 mV.

Recordings were obtained for each cell in standard extracellular solution or extracellular solution

containing 0.5 μM tetrodotoxin (TTX) or 10 μM VU170. TTX or VU170 was applied directly on the

recorded neuron with a custom-built fast flow perfusion system capable of complete solution exchange

in less than 1 s to minimize the time interval between control and drug recordings and current rundown.

Current traces from the TTX or VU170 solution were subtracted from the current traces obtained from

the standard solution. The difference current over the 100 ms at the end of the voltage pulse was

considered the steady state K_{Na} current. The same protocol was used for outside-out patch experiments.

I_{NaP} Current Measurements

For voltage-clamp experiments to measure the persistent Na^+ current, neurons were held at -80 mV and

given a 5 s voltage ramp at 20 mV/s. The ramp was repeated every 10 s. Recordings were obtained for

each cell in standard extracellular solution or extracellular solution containing 0.5 μM TTX. TTX was

applied directly on the recorded neuron with a custom-built fast flow perfusion system capable of

complete solution exchange in less than 1 s to minimize the time interval between control and TTX

recordings. The first current trace after TTX application was subtracted from the averaged current traces

obtained from the standard solution. At this time point, the effect of TTX on the Na^+-activated K^+ current

is minimal, while the persistent Na^+ current is maximally inhibited. This allowed us to isolate the

persistent Na^+ current from K_{Na} and obtain a more accurate estimate of changes in I_{NaP}.

Neuron Modeling

Model neurons (~10 per class) representing excitatory (Pyramidal Cell, PC), SST (Martinotti Cell, MC), PV

(Large Basket Cell, LBC), and (Bipolar Cell, BP) VIP neurons were downloaded from the EPFL/Blue Brain

project (https://bbp.epfl.ch/nmc-portal/downloads.html), and implemented in the NEURON

environment. Because no KCNT1-like conductance was present in the original models, a Hodgkin-Huxley

KCNT1 model conductance was created with both Na^+- and voltage-dependent gates and inserted into

the dendritic, somatic, and axonal compartments of model neurons (0.03 S/cm^2). The Na^+ dependence

was modeled after Bischoff et al. (EC_{50} = 40 mM, slope 3.5), and the voltage dependence was modeled

after our own experimental data (Fig. 5), corrected to account for the calculated liquid junction potential

in the experimental data. To verify that this approach reproduced experimental data, voltage step

protocols were run in NEURON with 10 mM and 0 mM internal Na^+. "K_{Na}" currents were obtained by

subtraction, and the amplitude and kinetics of the resulting currents were verified to be similar to

experimental values. To model the effects of KCNT1 GOF, the EC_{50} for Na^+ was decreased to either 35 or

30 mM, because the Y796H variant was previously shown to increase the Na^+ sensitivity of the channel.

These changes resulted in moderate increases in the current above WT levels (Fig. 6B), consistent with

the experimental data (Fig. 4 and Fig. 5). Modeling data was generated by running current clamp

protocols in NEURON for each of the model neurons and analyzed in the same way as experimental data.

For modeling the increase in I_{NaP}, the control I_{NaP} conductance was set to 0.001 times the conductance of

the transient Na^+ current in each compartment. This produced I_{NaP} currents approximately 1% of the

peak of the transient Na^+ current. KCNT1 GOF was modeled by decreasing the Na^+ EC_{50} to 30 mM and

the I_{NaP} increase was modeled by increasing its conductance 2-fold, in accordance with experimental

data.

Immunostaining, Imaging, and Quantification

For live imaging of GABAergic subtypes *in vitro*, cortical neurons isolated and cultured from VIP-Cre, SST-

Cre, and PV-Cre pups, were infected with AAV-*CamKII*-GFP and AAV-*hSyn*-DIO-mCherry viruses at DIV 1

to mark glutamatergic and Cre-expressing, GABAergic neurons, respectively. Images were captured from

live cultures at DIV 14 (VIP-Cre and SST-Cre) or DIV 16 (PV-Cre) using an inverted Olympus IX73

epifluorescent microscope with an Olympus PlanFluor 10× objective lens and an Andor Zyla sCMOS 4.2

camera controlled by µManager 2.0-β software (Edelstein et al., 2014). The same acquisition parameters

were used for each image and each group.

For soma size measurements, DIV 16 cultured SST-Cre neurons on coverslips were washed three times

with prewarmed PBS, fixed with 4% paraformaldehyde in PBS for 30 min at room temperature, and then

washed again three times with PBS. The fixed neurons were then incubated in blocking buffer (10%

normal goat serum and 0.1% Triton-100 in 1X PBS) for 1 h at room temperature, followed by incubation

with a Neu-N primary antibody (Synaptic Systems 266-004) diluted in blocking buffer (1:500) overnight

at 4°C. The next day, the neurons were washed three times with PBS for 10–15 min, incubated with goat

anti-guinea pig, Alexa Fluor 647 secondary antibody (Invitrogen A21450) in blocking buffer (1:1000) for 1

h at room temperature, and washed again three times with PBS for 10–15 min. The coverslips were then

flipped and mounted on glass slides using Fluoromount G with DAPI. Images (1024 × 1024 pixels) of Neu-

N to mark all neurons, *CamKII*-GFP to mark glutamatergic neurons, and *hSyn*-DIO-mCherry to mark Cre-

expressing neurons, were obtained using a DeltaVision Restoration Microscopy System (Applied

Precision/GE Life Sciences) with an inverted Olympus IX70 microscope with a 20 × oil objective,

SoftWoRx software, and a CoolSNAP-HQ charge-coupled device digital camera (Photometrics). Image

exposure times and settings were kept the same between groups in a culture and were optimized to

ensure that there were no saturated pixels. To analyze soma size, regions of interest (ROIs) were drawn

around cell bodies using the mCherry channel with Fiji software (Schindelin et al., 2012), and then the

cell body areas were measured for each mCherry[+] neuron imaged.

Mouse cortex slice preparation

Cortical slices were prepared from P21–P30 WT and *Kcnt1*-Y777H heterozygous pups, which were the

progeny of *Kcnt1*-Y777H heterozygous mice crossed to homozygous GIN mice (JAX stock #003718),

which express GFP in a subset of SST-positive neurons (Ma et al., 2006; Oliva et al., 2000), following a

modified protocol (Ting et al., 2014). Briefly, mice were deeply anesthetized with isoflurane and

transcardially perfused with 25–30 mL of ice-cold NMDG aCSF (in mM: 93 NMDG, 2.5 KCl, 1.2 NaH_2PO_4,

30 $NaHCO_3$, 20 HEPES, 25 D-glucose, 5 Na-ascorbate, 2 thiourea, 3 Na-pyruvate, 12 N-acetyl-L-cysteine,

10 $MgSO_4$, 0.5 $CaCl_2$, with pH adjusted to 7.3–7.4 with 10 N HCl) that was saturated with carbogen (95%

O_2, 5% CO_2 (pH 7.3–7.4). The mice were rapidly decapitated, and the heads were submerged in ice-cold

NMDG aCSF bubbled with carbogen. After extraction from the skull, the cortex was blocked and 250-µm

coronal sections were cut with a vibrating microslicer (Leica VT100S, Leica Biosystems) while submerged

in the same solution. Slices were first moved to NMDG aCSF at 32–34°C for a short-term recovery (≤ 15

min) before being transferred into HEPES-based holding aCSF (in mM: 92 NaCl, 2.5 KCl, 1.2 NaH_2PO_4, 30

$NaHCO_3$, 20 HEPES, 25 D-glucose, 5 Na-ascorbate, 2 thiourea, 3 Na-pyruvate, 12 N-acetyl-L-cysteine, 2

$MgSO_4$, 2 $CaCl_2$, bubbled with carbogen, with pH adjusted to 7.3–7.4 with KOH or HCl as necessary). The

slices were held in this solution until use.

Ex vivo slice electrophysiology

For recording, the cortical slices were placed in a chamber mounted on a fluorescence microscope (Zeiss

Axio Examiner.D1) and perfused with normal Ringer's solution containing (in mM): 130 NaCl, 3 KCl, 2

$MgCl_2$, 2 $CaCl_2$, 1.25 NaH_2PO_4, 26 $NaHCO_3$, and 10 D-glucose, bubbled with carbogen. Patch electrodes

were pulled from 1.5-mm o.d. thin-walled glass capillaries (Garner Glass Company) in two stages on a

micropipette puller (model P-1000; Sutter Instruments). Electrophysiological recordings were obtained

from GFP-labeled neurons in layer 2/3 of either motor or somatosensory cortex using pClamp, an

MultiClamp 700B amplifier, and a Digidata 1440A analog-to-digital converter (Molecular Devices). Data

were acquired at 20 kHz and low-pass filtered at 10 kHz. Internal solution contained the following (in

mM): 17.5 KCl, 112.5 K-gluconate, 1.5 NaCl, 5 Na_2-phosphocreatine, 1 $MgCl_2$, 10 HEPES, 0.2 EGTA, 3 Mg-

ATP, 0.3 GTP-Tris (pH 7.2, with KOH). The pipette resistance was between 3 and 5 MΩ. All experiments

were performed at room temperature (22–23°C).

The intrinsic electrophysiological properties of neurons were tested using current-clamp recording

under whole-cell configuration by injecting 1 s square current pulses in 5-pA steps, starting at −30 pA. All

recordings were saved in the computer for off-line analysis using Clampfit 11.2 or Axograph X software.

The membrane time constant (τ) was calculated from an exponential fit of current stimulus offset. Input

resistance (R_{in}) was calculated from the steady state of the voltage responses to the hyperpolarizing

current steps. Membrane capacitance was calculated by dividing the time constant by the input

resistance. Action potential (AP) frequency was counted on each depolarizing current step and plotted

as a function of injection current. Rheobase was defined as the minimum current required to evoke an

AP during the 1 s of sustained somatic current injections. AP threshold was defined as the membrane

potential at the inflection point of the rising phase of the AP, and the afterhyperpolarization (AHP) was

the difference between the AP threshold and the lowest V_m value within 50 ms. The membrane

potential values were not corrected for the liquid junction potential.

Data Presentation and Statistical Analysis

Prism 10 (GraphPad Prism, RRID:SCR_002798) was used to create all graphs. To test for statistical

significance for all whole cell electrophysiology experiments, we used generalized linear mixed models

(GLMM) in SPSS [28.0 Chicago, III (IBM, RRID:SCR_002865], which allows for within-subject correlations

and the specification of the most appropriate distribution for the data. Because neurons and animals

from the same culture or animal are not independent measurements, culture or litter was used as the

subject variable, and animals, neurons, and voltage steps were considered within-subject measurements.

All data distributions were assessed with the Shapiro-Wilk test. Datasets that were significantly different

from the normal distribution ($p < 0.05$) were fit with models using the gamma distribution and a log link

function, except for synaptic connection probability, which was fit with the binomial distribution and

probit link. Normal datasets were fit with models using a linear distribution and identity link. We used

the model-based estimator for the covariance matrix and goodness of fit was determined using the

corrected quasi likelihood under independence model criterion and by the visual assessment of

residuals. All values reported in the text, figures, and tables are estimated marginal means +/- standard

error, except where noted.

Acknowledgements

This work was supported by NIH/NINDS grants NS087095 (M.C.W.), NS110945 (M.C.W.), NS031348

(W.N.F.), R33NS109521 (C.D.W. and K.A.E.), and OD020351 (The Jackson Laboratory Center for Precision

Genetics). The authors would like to acknowledge the support of the Microscopy Imaging Center (RRID#

SCR_018821) and the Cellular & Molecular Core at the University of Vermont.

Author Contributions: Conceptualization, A.N.S., W.N.F., M.C.W.; Methodology, A.M.Q., B.D.S., K.A.E.,

C.D.W., Formal Analysis, A.N.S., M.C.W. K.L., M.S.; Investigation, A.N.S., M.C.W., K.L., M.S.; Writing-

Original Draft, A.N.S, M.C.W., Revising and Editing, A.N.S., M.C.W.; Supervision, K.A.E., C.D.W., W.N.F.,

M.C.W.; Funding Acquisition, W.N.F., M.C.W.

FIGURE LEGENDS

Figure 1. Heterozygous *Kcnt1*-Y777H expression alters AP shape and generation in NFS GABAergic

neurons. (A_1-A_3) On the left, representative responses to step currents are shown for glutamatergic, and

FS and NFS GABAergic, WT (black) and YH-HET (colors) neurons (top to bottom), illustrating the input

resistance (in response to a depolarizing step) and the rheobase (the first trace with an AP in response

to a hyperpolarizing step) for each neuron type. On the right, bar graphs show quantification and mean

± SEM of the membrane properties and AP parameters for each neuron type for WT (grey) and YH-HET

(colors) groups, with individual neuron measurements overlaid in scatter plots. The p-values are shown

on each graph where $p < 0.05$. **(B_1-B_3)** Representative traces are shown at low, medium, and high

current steps for glutamatergic, and FS and NFS GABAergic, WT (black) and YH-HET (colors) neurons (left

to right). The line graphs below show the number of APs (mean ± SEM) per current injection step in WT

(black) and YH-HET (colors) neurons. Statistical significance was tested using Generalized Linear Mixed

Models, and p-values are shown on each graph where $p < 0.05$.

Figure 2. Heterozygous *Kcnt1*-Y777H expression increases subthreshold K_{Na} currents in NFS GABAergic

neurons. (A_1-A_3) Representative traces in control (top), 0.5 μM TTX (middle), and the difference current

(bottom) calculated by subtracting the membrane current response to voltage steps (-80 to +50 mV)

from a holding potential of -70 mV in TTX from the response in control external solution in glutamatergic,

and FS and NFS GABAergic, WT (black) and YH-HET (colors) neurons. To include all of the representative

traces in the figures, and prevent overlap of the traces, the large inward sodium currents were removed

from each set of traces using the masking tool in Adobe Illustrator. **(B_1-B_3)** Summary data shows the K_{Na}

current (mean ± SEM) for each voltage step in glutamatergic, and FS and NFS GABAergic, WT (black and

gray) and YH-HET (colors) neurons. The p-values are shown on each graph where $p < 0.05$, and the n

values are the number of neurons recorded for each group. **(C$_1$-C$_3$)** Plots of the K$_{Na}$ current (mean ± SEM)

for each voltage step from -80 to 0 mV in WT (black and gray) and YH-HET (colors) neurons to illustrate

the values that are too small to be seen on the graphs in B$_1$-B$_3$. The shaded red area in C$_3$ indicates the

subthreshold voltage range with significantly higher K$_{Na}$ currents (red voltages along x-axis indicate steps

where $p < 0.05$) in YH-HET relative to WT neurons. The p-values are shown on each graph where $p < 0.05$,

and the n values are the number of neurons recorded for each group. Statistical significance for I-V plots

was tested using Generalized Linear Mixed Models with genotype and voltage step as fixed effects

followed by pairwise comparisons at each level.

Figure 3. Heterozygous *Kcnt1*-Y777H expression does not alter synaptic connectivity or the excitation-

inhibition balance. (A$_1$-A$_4$) Evoked postsynaptic currents (PSCs) were recorded from neuron pairs

[glutamatergic (excitatory, E) and GABAergic (inhibitory, I)] by stimulating the neuron type indicated on

the left and recording the response in the neuron type indicated on the right (WT, gray; YH-HET E,

green; YH-HET I, red). Bar graphs below each recorded neuron pair schematic show summary data

(mean ± SEM) of the connection probability (left graph; numbers on bars represent connected pair

number/recorded pair number) and peak evoked PSC amplitude (right graph; dots represent individual

evoked responses) between each motif. **(B$_1$)** Bar graphs with overlaid individual neuron measurements

and mean ± SEM show the spontaneous EPSC (sEPSC) or IPSC (sIPSC) frequency onto E neurons (WT,

gray; YH-HET, green). **(B$_2$)** Bar graphs with overlaid individual neuron measurements and mean ± SEM

show the sEPSC or sIPSC frequency onto I neurons (WT, gray; YH-HET, red). **(B$_3$)** Scatter plots show

individual E/I ratio measurements onto E neurons (WT, gray; YH-HET, green) and I neurons (WT, gray;

YH-HET, red). The p-values are shown on each graph where $p < 0.05$.

1003

Figure 4. Heterozygous *Kcnt1*-Y777H expression differentially affects the intrinsic excitability of SST-

and PV-expressing GABAergic neurons. (A) A schematic diagram illustrates the strategy for generating

fluorescently labeled GABAergic subtype-specific neurons. YH-HET mice were crossed to VIP-, SST-, or

PV-Cre mice, and the resulting P0 WT and YH-HET littermate pups were used to isolate and culture

cortical neurons. At DIV 1, neurons were infected with AAV-*CamKII*-GFP to label glutamatergic neurons,

and AAV-*hSyn*-DIO-mCherry to label Cre-expressing neurons. At DIV 13–17, whole-cell, patch-clamp

electrophysiology was performed on mCherry$^+$/GFP$^-$ neurons. **(B$_1$-B$_3$)** On the left, representative

responses to step currents are shown for VIP-, SST-, and PV-expressing WT (black) and YH-HET (colors)

neurons (top to bottom). For each neuron type, the superimposed dark traces illustrate the input

resistance (in response to a depolarizing step) and the rheobase (the first trace with an AP in response

to a hyperpolarizing step), and the light trace shows the first step current response to induce repetitive

AP firing across the step. Above the superimposed traces, the first AP of each rheobase trace is shown

(same vertical scale, increased horizontal scale). On the right, bar graphs show quantification of the

membrane properties and AP parameters for each neuron type (VIP, SST, and PV, top to bottom) for WT

(grey) and YH-HET (colors) groups, with individual neuron measurements overlaid in scatter plots. The p-

values are shown above each graph where $p < 0.05$. **(C$_1$-C$_3$)** For VIP-, SST-, and PV-expressing neurons

(left to right), representative traces are shown at low, medium, and high current steps, and the line

graphs below show the number of APs (mean ± SEM) per current injection step in WT (black) and YH-

HET (colors) neurons. Statistical significance was tested using Generalized Linear Mixed Models, and p-

values are shown above each graph where $p < 0.05$.

1024

Figure 5. The *Kcnt1*-Y777H variant increases KCNT1-mediated currents across subthreshold voltages in

SST- and PV-expressing GABAergic neurons. (A$_1$-A$_3$) Representative traces in control (top), 10 µM

VU170 (middle), and the difference current (bottom) calculated by subtracting the membrane current

response to voltage steps (-80 to +50 mV) from a holding potential of -70 mV in VU170 from the

response in control external solution in VIP-, SST-, and PV-expressing, WT (black) and YH-HET (colors)

neurons. To include all of the representative traces in the figures, and prevent overlap of the traces, the

large inward sodium currents were removed from each set of traces using the masking tool in Adobe

Illustrator. **(B_1-B_3)** Summary data shows the KCNT1 current (mean ± SEM) for each voltage step in VIP-,

SST-, and PV-expressing, WT (black and gray) and YH-HET (colors) neurons. The p-values are shown on

each graph where $p < 0.05$, and the n values are the number of neurons recorded for each group. **(C_1-C_3)**

Plots of the K_{Na} current (mean ± SEM) for each voltage step from -80 to 0 mV in WT (black and gray) and

YH-HET (colors) neurons to illustrate the values that are too small to be seen on the graphs in B_1-B_3. The

shaded red areas in C_2 and C_3 indicate the subthreshold voltage range with significantly higher K_{Na}

currents ($p < 0.05$) in YH-HET relative to WT neurons. Statistical significance for I-V plots was tested

using Generalized Linear Mixed Models with genotype and voltage step as fixed effects followed by

pairwise comparisons at each level.

1041

Figure 6. Compartmental models of KCNT1 GOF in SST, but not PV, neurons are consistent with

experimental data. (A) Activation curves for VIP, SST, and PV neurons used to model the KCNT1 channel

in each neuron type. **(B)** Representative traces of the response of an outside-out membrane patch

pulled from an SST neuron in control external solution (top) and after application of 10 μm VU170

(bottom). **(C)** Representative trace of the VU170-sensitive current in an outside-out patch pulled from an

SST neuron with a single exponential fit (red curve) overlaid and summary data showing the time

constant (mean ± SEM) obtained from these fits. **(D)** Simulated traces showing the KCNT1-mediated

current in model neurons at three levels of Na^+-sensitivity (Con-40 mM, GOF-35 mM, and GOF-30 mM).

(E_1, F_1, and G_1) Simulated traces from model VIP (E_1; purple), SST (F_1; blue), and PV (G_1; yellow) neurons

in response to 500-ms current injections at three levels of Na$^+$-sensitivity [control (Con), and two levels

of KCNT1 GOF]. Representative traces are shown at increasing current steps from bottom to top for

each level. **(E$_2$, F$_2$, and G$_2$)** Summary data from 10 model VIP (E$_2$; purple), SST (F$_2$; blue), and PV (G$_2$;

yellow) neurons showing (from left to right) the number of APs at different current steps (F-I plot), the

input resistance, and rheobase. The bar graphs (mean ± SEM) are overlaid with scatter plots of individual

neuron measurements. Statistical significance was tested using a Repeated Measures ANOVA. Asterisks

indicate where $p < 0.05$ (*), $p \leq 0.01$ (**), and $p \leq 0.001$ (***).

1058

Figure 7. The persistent Na$^+$ current is increased by the *Kcnt1*-Y777H variant in PV, but not SST,

neurons. (A) Representative traces of WT (left) and YH-HET (right) SST neurons in response to slow

voltage ramp in control (dark gray) and 7 seconds after application of 0.5 nM TTX (light gray) with the

difference current (I$_{NaP}$) plotted on the same scale. **(B)** I$_{NaP}$ I-V curves of WT (gray) and YH-HET (blue) SST

neurons constructed from averaging the ramp-evoked difference current (mean ± SEM) at 5-mV

intervals (left), and a bar graph with overlaid scatter plot showing the peak negative value for each

neuron (right). **(C)** Representative traces of WT (left) and YH-HET (right) PV neurons in response to slow

voltage ramp in control (dark gray) and 7 seconds after application of 0.5 nM TTX (light gray) with the

difference current (I$_{NaP}$) plotted on the same scale. The upper traces and difference currents are plotted

on different scales. **(D)** I$_{NaP}$ I-V curves of WT (gray) and YH-HET (orange) PV neurons constructed from

averaging the ramp-evoked difference current (mean ± SEM) at 5-mV intervals (left), and a bar graph

with overlaid scatter plot showing the peak negative value for each neuron (right). **(E)** Simulated traces

from a model PV neuron in response to 500-ms current injections with control levels of KCNT1 (black

traces) and KCNT1 GOF (Na$^+$ EC$_{50}$ = 35 mM) with a 2-fold increase in I$_{NaP}$ (orange traces). Representative

traces are shown at increasing current steps from bottom to top for each level. **(F)** F-I plot shows the

increase in AP firing (left), and a bar graph with overlaid scatter plot of individual neuron values shows

the decrease in rheobase associated with modeling KCNT1 GOF (Na^+ EC_{50} = 35 mM) together with the

increase in I_{NaP}. For modeling data, statistical significance was tested using a Repeated Measures ANOVA.

Asterisks indicate where $p < 0.05$ (*), $p \leq 0.01$ (**), and $p \leq 0.001$ (***).

1078

Figure 8. YH-HET SST neurons show increased chemical and electrical coupling and receive increased

excitatory input. (A and B) Evoked postsynaptic currents (PSCs) were recorded from SST neuron pairs by

stimulating each neuron at 0.1 Hz and recording the response onto the partner. **(A)** A bar graph shows

the connection probability (mean + SEM) between SST neurons (WT, gray; YH-HET, blue; numbers on

bars represent connected pair number/recorded pair number). **(B)** On the left, example traces of evoked

IPSCs [individual IPSCs (light) overlaid by averaged IPSCs (dark)], and on the right, a graph of connection

strength between SST neurons (WT, gray; YH-HET, blue; individual measurements and summary violin

plots). **(C$_1$)** A schematic illustrates representative responses (presynaptic, black; postsynaptic, red)

between chemically coupled (upper two panels) and electrically coupled (lower two panels) SST neurons

following 100-pA (left two panels) and AP (right two panels) stimulation. **(C$_2$)** Summary data of four

possible connection motifs (not coupled, chemical: one-way, chemical: two-way, and electrical) tested

among WT (gray box) and YH-HET (blue box) SST neuron pairs (20 pairs/group). **(D)** On the left, example

traces of spontaneous EPSCs (sEPSCs) recorded onto SST neurons (WT, black; YH-HET, blue). On the right,

a graph shows individual neuron measurements and summary violin plots of the sEPSC frequency onto

SST neurons (WT, gray; YH-HET, blue). All significant p-values are displayed at the top of each graph.

1094

Figure 9. Acute cortical slice recordings confirm the reduced intrinsic excitability of YH-HET SST

neurons. (A) An inverted monochrome image illustrates the recording scheme from GFP-expressing

neurons in acute slices from P21–P30 GIN mice that express GFP in a subset of SST-positive neurons. **(B)**

Representative responses to current steps are shown for WT (black) and YH-HET (blue) to three,

equivalent 1 s current steps. **(C)** Plot of the action potential frequency (mean ± SEM) per current

injection step shows that YH-HET GFP-expressing neurons (blue) fire fewer action potentials than those

of WT (grey) across a range of current inputs. **(D)** Bar graphs show quantification of the membrane

properties and AP parameters for WT (grey) and YH-HET (colors) GFP-expressing neuron groups, with

individual neuron measurements overlaid in scatter plots.

1104

Supplemental Figure 2-1. The magnitudes of the K_{Na} current increases in YH-HET glutamatergic and

NFS GABAergic neurons are intermediate to those of WT and YH-HOM neurons. Bar graphs (top three)

illustrate differences in K_{Na} current (mean ± SEM) between YH-HET [dark color; WT littermates (black)]

and YH-HOM [light color; WT littermates (gray)] neurons for voltage steps where significant increases

were previously observed (red asterisks where $p < 0.05$) in glutamatergic, and FS and NFS GABAergic,

YH-HOM neurons. Line graphs (bottom three) illustrate differences in K_{Na} current (mean ± SEM) between

YH-HET [dark color; WT littermates (black)] and YH-HOM [light color; WT littermates (gray)] neurons

across negative voltage steps (-80 to -20 mV; red asterisks where $p < 0.05$ in YH-HOM compared with

WT) in glutamatergic, and FS and NFS GABAergic, neurons.

1114

Supplemental Figure 4-1. Cre recombinase-expressing, cortical GABAergic neuron subpopulations

retain characteristic electrophysiological features in culture. Cortical neurons isolated and cultured

from P0 pups (*Kcnt1*^{+/+}) from VIP-, SST-, or PV-Cre mouse lines were infected with AAV-*CamKII*-GFP to

label glutamatergic neurons, and AAV-*hSyn*-DIO-mCherry to label Cre-expressing neurons, at DIV 1. **(A)**

Images show GFP^{+} glutamatergic neurons and mCherry[+] VIP-, SST-, or PV-expressing neurons (left to

right) at DIV 14 (VIP and SST) or DIV 16 (PV). The scale bars represent 50 μm. **(B)** Representative

responses to step currents are shown for VIP (purple), SST (blue), and PV (orange) neurons (left to right).

For each neuron type, the superimposed dark traces illustrate the input resistance (in response to a

depolarizing step) and the rheobase (the first trace with an AP in response to a hyperpolarizing step),

and the light trace shows the first step current response to induce repetitive AP firing across the step.

(C) On the left, the first AP of each rheobase trace is shown for VIP (purple), SST (blue), and PV (orange)

neurons. On the right, bar graphs show quantification of the membrane properties and AP parameters

for each neuron type (VIP, SST, and PV, left to right), with individual neuron measurements overlaid in

scatter plots.

1129

Supplemental Figure 4-2. YH-HET SST neurons have increased soma size compared with those of WT.

Cortical neurons were isolated and cultured from P0 pup progeny (WT and YH-HET) of YH-HET mice

crossed to SST-Cre mice. Neurons were infected with AAV-*CamKII*-GFP to label glutamatergic neurons,

and AAV-*hSyn*-DIO-mCherry to label Cre-expressing neurons, at DIV 1, and immunostained using anti-

Neu-N antibodies at DIV 16. **(A)** Top two images show mCherry$^+$ SST-expressing neurons as a

subpopulation of Neu-N$^+$ cortical neurons, and bottom two images show GFP$^-$, mCherry$^+$ SST-expressing

neurons intermingled with GFP$^+$ glutamatergic neurons. The scale bars represent 50 μm. **(B)** A graph

shows individual neuron measurements and summary violin plots of the mCherry$^+$ neuron soma size

(WT, gray; YH-HET, blue) with the p-value displayed at the top.

1139

Supplemental Figure 4-3. YH-HET and WT PV neurons show similar neurophysiological responses to

increased temperature. YH-HET mice were crossed to PV-Cre mice, and the resulting P0 WT and YH-HET

littermate pups were used to isolate and culture cortical neurons. At DIV 1, neurons were infected with

AAV-*CamKII*-GFP to label glutamatergic neurons, and AAV-*hSyn*-DIO-mCherry to label Cre-expressing

neurons. At DIV 16 and 17, whole-cell, patch-clamp electrophysiology was performed on glutamatergic

(mCherry$^-$/GFP$^+$), GABAergic (mCherry$^-$/GFP$^-$), or PV (mCherry$^+$/GFP$^-$) neurons at room temperature

(~22°C). Then, the external solution was warmed to 35°C, and the measurements were repeated. **(A)**

Before-after plots of glutamatergic and GABAergic WT neurons (n=16) show alterations in all parameters

measured with increased temperature, excluding the resting membrane potential (V_{rest}) and membrane

capacitance (C_m). **(B)** Before-after plots show highly similar responses to temperature increases between

WT (n=11) and YH-HET (n=10) PV neurons for rheobase, AP amp, and maximal firing frequency (MFR).

Empty circles represent individual neurons, and filled-in circles represent the averages for each group.

Lines connect the before and after measurements for each neuron.

1153

Supplemental Figure 5-1. The *Kcnt1*-Y777H variant increases K_{Na} currents measured with TTX

subtraction across subthreshold voltages in SST-expressing GABAergic neurons. (A_1-A_3) Representative

traces in control (top), 0.5 µM TTX (middle), and the difference current (bottom) calculated by

subtracting the membrane current response to voltage steps (-80 to +50 mV) from a holding potential of

-70 mV in TTX from the response in control external solution in VIP-, SST-, and PV-expressing, WT (black)

and YH-HET (colors) neurons. To include all of the representative traces in the figures, and prevent

overlap of the traces, the large transient inward sodium currents were removed from each set of traces

using the masking tool in Adobe Illustrator. **(B_1-B_3)** Summary data shows the K_{Na} current (mean ± SEM)

for each voltage step in VIP-, SST-, and PV-expressing, WT (black and gray) and YH-HET (colors) neurons.

The p-values are shown on each graph where $p < 0.05$, and the n values are the number of neurons

recorded for each group. **(C_1-C_3)** Plots of the K_{Na} current (mean ± SEM) for each voltage step from -80 to

0 mV in WT (black and gray) and YH-HET (colors) neurons to illustrate the values that are too small to be

seen on the graphs in B_1-B_3. The shaded red area in C_2 indicates the subthreshold voltage range with

significantly higher K_{Na} currents in YH-HET relative to WT neurons. Statistical significance for I-V plots

was tested using Generalized Linear Mixed Models with genotype and voltage step as fixed effects

followed by pairwise comparisons at each level.

1170

Supplemental Figure 5-2. Isolation of K_{Na} currents from neurons lacking KCNT1 and KCNT2 channels

illustrates the specificity of VU170- vs. TTX-subtraction methods. Voltage steps were applied to

voltage-clamped neurons isolated and cultured from $Kcnt1^{-/-}$; $Kcnt2^{-/-}$ double knockout mice to compare

the delayed outward currents before and after the addition of either 10 μM VU170 or 0.5 μM TTX. **(A)** A

summary plot shows the VU-sensitive (red) and TTX-sensitive (grey) difference currents (mean ± SEM)

for each voltage step, which were calculated by subtracting the membrane current response to voltage

steps (-80 to +50 mV) from a holding potential of -70 mV in VU170 or TTX from the response in control

external solution. **(B)** A summary plot of the VU-sensitive (red) and TTX-sensitive (grey) difference

currents (mean ± SEM) for each voltage step from -80 to 0 mV to illustrate the values that are too small

to be seen on the graph in (A).

1181

Supplemental Figure 6-1. Model VIP and glutamatergic neurons with insertion of SST-like, KCNT1 GOF

kinetics show reduced excitability. (A_1) Simulated traces from model VIP neurons, using VIP KCNT1

activation curve parameters (purple) or SST KCNT activation curve parameters (blue), in response to

500-ms current injections at two levels of Na^+-sensitivity (Con-40 mM and GOF-30 mM). Representative

traces are shown at increasing current steps from bottom to top for each level. **(A_2)** Summary data from

10 model VIP neurons, with VIP KCNT1 parameters (VIP-Con and VIP-GOF) or SST KCNT1 parameters

(SST-GOF) showing (from left to right) the number of APs at different current steps (F-I plot), the input

resistance, and rheobase. The bar graphs (mean ± SEM) are overlaid with scatter plots of individual

neuron measurements. **(B_1)** Simulated traces from model glutamatergic neurons, using glutamatergic

KCNT1 activation curve parameters (green) or SST KCNT activation curve parameters (blue), in response

to 500-ms current injections at two levels of Na^+-sensitivity (Con-40 mM and GOF-30 mM).

Representative traces are shown at increasing current steps from bottom to top for each level. **(B_2)**

Summary data from 10 model glutamatergic neurons, with glutamatergic KCNT1 parameters (Glut-Con

and Glut-GOF) or SST KCNT1 parameters (SST-GOF) showing (from left to right) the number of APs at

different current steps (F-I plot), the input resistance, and rheobase. The bar graphs (mean ± SEM) are

overlaid with scatter plots of individual neuron measurements. Statistical significance was tested using a

Repeated Measures ANOVA. Asterisks indicate where $p < 0.05$ (*), $p \leq 0.01$ (**), and $p \leq 0.001$ (***).

1199

Supplemental Figure 6-2. Model neurons show differential responses to KCNT1 GOF. (A) Bar graphs

(mean ± SEM) with overlaid dot plots show the effects of two levels of KCNT1 GOF (Na^+ EC_{50} = 35 or 30

mM) relative to those of control (EC_{50} = 40 mM) on input resistance, rheobase, and AP firing rate at 1.5 ×

rheobase for GABAergic subtypes (10 model neurons/group). **(B)** Bar graphs (mean ± SEM) with overlaid

dot plots show the effects KCNT1 GOF (Na^+ EC_{50} = 30 mM) on VIP GABAergic or glutamatergic model

neurons (10/group) with their respective KCNT1 activation curve parameters (VIP-GOF and Glut-GOF) or

SST KCNT1 parameters (SST-GOF) relative to those of control (EC_{50} = 40 mM) on input resistance,

rheobase, and AP firing rate at 1.5 × rheobase.

1208

Supplemental Figure 7-1. The rate of K_{Na} current block after TTX application is slow relative to I_{NaP}

block in GABAergic neurons. Plots show the time course over which I_{NaP} and K_{Na} currents are blocked

following 0.5 µM TTX application. Neurons of the labeled type in each panel were voltage clamped and

subjected to a slow voltage ramp protocol (20 mV/s over 5 s). TTX (0.5 µM) was applied at time 0 and

K_{Na} currents and I_{NaP} were measured on each successive sweep. The K_{Na} current was measured as the

normalized difference between each trace and the mean of the pre-TTX current between +20 and +30

mV. I_{NaP} was measured as the normalized difference between each trace and the mean of the pre-TTX

current at the 10 mV bracketing the local peak inward current. Once the minimum value was reached

for I_{NaP}, all subsequent values were considered zero.

1218

1219 **Supplemental Figure 7-2. Model PV neurons show differential responses to KCNT1 GOF with increased**

1220 **I_{NaP}.** Bar graphs (mean ± SEM) with overlaid dot plots show the effects of two levels of KCNT1 GOF (Na^+

1221 EC_{50} = 35 or 30 mM), with or without additional I_{NaP}, relative to those of control (EC_{50} = 40 mM) on input

1222 resistance, rheobase, and AP firing rate at 1.5 × rheobase for PV neurons (10/group).

1223

1224

1225

1226 **Figure 1- Supplemental Table. Electrophysiological parameters of current clamp recordings from**
1227 **neuronal cultures.**

1228

1229

	Glutamatergic			FS GABAergic			NFS GABAergic		
	WT	YH-HET		WT	YH-HET		WT	YH-HET	
	N = 9	N = 9		N = 9	N = 9		N = 9	N = 9	
	n = 27	n = 30	p-value	n = 27	n = 29	p-value	n = 26	n = 22	p-value
V_{rest} (mV)	-57.9±1.6	-57.0±1.7	0.61	-53.8±1.8	-52.8±1.8	0.60	-56.3±1.8	-54.4±1.9	0.34
R_{in} (MΩ)	177±17	182±16	0.80	95±9	105±10	0.40	171±16	146±14	0.19
Tau (ms)	32.9±3.8	39.3±4.3	0.14	17.3±2.0	16.7±1.9	0.75	24.9±2.9	22.2±2.7	0.37
C_m (pF)	198±15	215±16	0.23	183±14	163±12	0.15	143±11	164±13	0.09
Rheo (pA)	171±19	148±16	0.25	400±45	415±47	0.78	196±22	297±36	**0.005**
AP_{thresh} (mV)	-28.2±1.5	28.3±1.5	0.93	-25.7±1.6	-25.8±1.6	0.95	-28.2±1.6	-24.4±1.7	**0.012**
AP_{amp} (mV)	77.8±2.0	77.5±1.5	0.51	70.9±1.8	67.8±1.7	0.15	71.5±1.8	68.7±1.9	0.28
AP_{hw} (ms)	2.19±0.12	2.20±0.12	0.95	0.89±0.05	0.91±0.05	0.85	1.36±0.08	1.11±0.07	**0.01**
AHP (mV)	11.8±0.6	11.8±0.6	0.96	27.5±1.3	26.9±1.4	0.78	19.3±1.0	22.6±1.2	**0.034**
AP_{mfr} (Hz)	22.4±2.3	22.4±2.4	0.99	68.8±6.2	71.4±6.6	0.53	35.0±3.8	31.8±3.5	0.25

For an explanation of the parameters, see Methods. R_{in} = input resistance, Tau = membrane time
constant, Rheo = rheobase current, C_m = membrane capacitance, AP = action potential, thresh =
threshold, amp = amplitude, hw = half width, AHP = afterhyperpolarization, and mfr = maximum firing
rate. Values shown are estimated marginal means ± the standard error as determined by implementing
a Generalized Linear Mixed Model. P values less than 0.05 are in red, bold type. For each subgroup, N
values are the number of mouse pups, and n values are the number of neurons.

1236

1237

1238

1239

1240

Figure 4- Supplemental Table. Electrophysiological parameters of current clamp recordings from

neuronal cultures of three major GABAergic subtypes.

	SST$^+$ GABAergic			VIP$^+$ GABAergic			PV$^+$ GABAergic		
	WT	YH-HET		WT	YH-HET		WT	YH-HET	
	N = 4	N = 5		N = 3	N = 4		N = 3	N = 4	
	n = 45	n = 46	p-value	n = 50	n = 48	p-value	n = 23	n = 25	p-value
R_{in} (MΩ)	133±18	95±13	**0.001**	243±10	249±11	0.645	67±4	66±3	0.850
Tau (ms)	16.9±1.1	13.5±0.8	**0.012**	35.1±2.4	33.1±2.3	0.384	6.7±0.5	6.0±0.4	0.302
C_m (pF)	135±13	152±15	**0.032**	148±6	134±6	0.055	103±5	90±4	0.050
Rheo (pA)	232±41	319±56	**0.001**	149±12	157±13	0.344	546±33	441±24	**0.031**
AP_{thresh} (mV)	-36.4±0.8	-35.2±0.8	0.113	-30.9±0.4	-31.0±0.4	0.850	-34.2±1.8	-35.5±1.9	0.308
AP_{amp} (mV)	79.9±1.5	80.2±1.5	0.850	75.6±0.9	75.5±0.9	0.943	77.6±1.0	82.7±1.0	**0.002**
AP_{hw} (ms)	0.91±0.02	0.86±0.02	0.142	1.09±0.04	1.09±0.04	0.916	0.73±0.03	0.69±0.02	0.180
AHP (mV)	23.0±0.7	23.5±0.7	0.554	17.2±0.9	17.4±0.9	0.860	26.6±0.6	25.3±0.5	0.150
AP_{mfr} (Hz)	57.6±5.1	55.7±4.9	0.432	35.5±2.8	35.9±2.8	0.857	69.4±4.9	79.0±5.3	0.068

For an explanation of the parameters, see Methods. R_{in} = input resistance, Tau = membrane time

constant, Rheo = rheobase current, C_m = membrane capacitance, AP = action potential, thresh =

threshold, amp = amplitude, hw = half width, AHP = afterhyperpolarization, and mfr = maximum firing

rate. Values shown are estimated marginal means ± the standard error as determined by implementing

a Generalized Linear Mixed Model. P values less than 0.05 are in red, bold type. For each subgroup, N

values are the number of mouse pups, and n values are the number of neurons.

1264

1265

1266 **Figure 6- Supplemental Table. Electrophysiological parameters using compartmental model neurons.**

Model neuron	Input Resistance (MΩ)	p-value vs 40 mM	Rheobase (pA)	p-value vs 40 mM	AP rate at 1.5x Rheobase (Hz)	p-value vs 40 mM
VIP-40 mM	266.5 ± 28.2		78.7 ± 3.7		7.0 ± 0.7	
VIP-35 mM	266.2 ± 28.1	0.193	79.1 ± 3.6	0.177	7.0 ± 0.7	0.999
VIP-30 mM	264.8 ± 28.0	0.012	80.4 ± 3.5	0.014	6.9 ± 0.7	0.527
VIP-30 mM + SST	260.2 ± 27.8	<0.001	86.3 ± 3.8	<0.001	6.5 ± 0.8	0.011
SST-40 mM	210.3 ± 23.0		127.5 ± 8.5		17.3 ± 2.1	
SST-35 mM	199.9 ± 22.1	0.009	156.3 ± 11.7	<0.001	11.9 ± 2.0	0.025
SST-30 mM	187.1 ± 21.2	0.001	180.0 ± 12.3	<0.001	7.1 ± 1.9	0.003
PV-40 mM	105.8 ± 7.7		258.0 ± 7.5		18.3 ± 1.3	
PV-35 mM	105.5 ± 7.7	0.140	264.5 ± 7.9	0.004	16.8 ± 1.4	0.033
PV-30 mM	103.3 ± 7.5	<0.001	273.0 ± 9.5	0.004	12.4 ± 2.0	0.004
PV-35 mM + I_{NaP}	104.6 ± 7.5	<0.001	238.5 6.3	<0.001	21.1 ± 1.3	<0.001
PV-30 mM + I_{NaP}	103.4 ± 7.4	0.009	250.5 ± 6.9	0.002	20.1 ± 1.5	0.012
Glut-40 mM	147.1 ± 12.0		117.0 ± 10.4		2.46 ± 0.3	
Glut-30 mM	146.4 ± 12.1	0.017	121.5 ± 10.0	<0.001	2.36 ± 0.4	0.593
Glut-30 mM + SST	140.4 ± 11.9	0.001	146.0 ± 9.7	<0.001	1.54 ± 0.3	0.012

1267

1268

1269

1270 For an explanation of the parameters, see Methods. AP = action potential. For each subgroup, 10 model
1271 neurons were used and simulations were repeated with the EC_{50} (in mM) for Na^+ for the KCNT1
1272 conductance set at the indicated levels. "+ SST" refers to the activation curve parameters (V_{50} and slope)
1273 for the KCNT1 conductance in that neuron type being replaced with those measured in SST neurons.
1274 Values shown are means ± the standard error. The statistical test is a repeated measures ANOVA.

1275

1276

1277

1278

1279

1280

1281

REFERENCES

Alcami, P., & Pereda, A. E. (2019). Beyond plasticity: the dynamic impact of electrical synapses on neural circuits. *Nat Rev Neurosci*, *20*(5), 253-271. https://doi.org/10.1038/s41583-019-0133-5

Amitai, Y., Gibson, J. R., Beierlein, M., Patrick, S. L., Ho, A. M., Connors, B. W., & Golomb, D. (2002). The spatial dimensions of electrically coupled networks of interneurons in the neocortex. *J Neurosci*, *22*(10), 4142-4152. https://doi.org/10.1523/JNEUROSCI.22-10-04142.2002

Avermann, M., Tomm, C., Mateo, C., Gerstner, W., & Petersen, C. C. (2012). Microcircuits of excitatory and inhibitory neurons in layer 2/3 of mouse barrel cortex. *J Neurophysiol*, *107*(11), 3116-3134. https://doi.org/10.1152/jn.00917.2011

Barcia, G., Fleming, M. R., Deligniere, A., Gazula, V. R., Brown, M. R., Langouet, M., Chen, H., Kronengold, J., Abhyankar, A., Cilio, R., Nitschke, P., Kaminska, A., Boddaert, N., Casanova, J. L., Desguerre, I., Munnich, A., Dulac, O., Kaczmarek, L. K., Colleaux, L., & Nabbout, R. (2012). De novo gain-of-function KCNT1 channel mutations cause malignant migrating partial seizures of infancy. *Nat Genet*, *44*(11), 1255-1259. https://doi.org/10.1038/ng.2441

Berg, A. P., Sen, N., & Bayliss, D. A. (2007). TrpC3/C7 and Slo2.1 are molecular targets for metabotropic glutamate receptor signaling in rat striatal cholinergic interneurons. *J Neurosci*, *27*(33), 8845-8856. https://doi.org/10.1523/JNEUROSCI.0551-07.2007

Bhattacharjee, A., Gan, L., & Kaczmarek, L. K. (2002). Localization of the Slack potassium channel in the rat central nervous system. *J Comp Neurol*, *454*(3), 241-254. https://doi.org/10.1002/cne.10439

Bonardi, C. M., Heyne, H. O., Fiannacca, M., Fitzgerald, M. P., Gardella, E., Gunning, B., Olofsson, K., Lesca, G., Verbeek, N., Stamberger, H., Striano, P., Zara, F., Mancardi, M. M., Nava, C., Syrbe, S., Buono, S., Baulac, S., Coppola, A., Weckhuysen, S., . . . Rubboli, G. (2021). KCNT1-related epilepsies and epileptic encephalopathies: phenotypic and mutational spectrum. *Brain*, *144*(12), 3635-3650. https://doi.org/10.1093/brain/awab219

Budelli, G., Hage, T. A., Wei, A., Rojas, P., Jong, Y. J., O'Malley, K., & Salkoff, L. (2009). Na+-activated K+ channels express a large delayed outward current in neurons during normal physiology. *Nat Neurosci*, *12*(6), 745-750. https://doi.org/10.1038/nn.2313

Casale, A. E., Foust, A. J., Bal, T., & McCormick, D. A. (2015). Cortical Interneuron Subtypes Vary in Their Axonal Action Potential Properties. *J Neurosci*, *35*(47), 15555-15567. https://doi.org/10.1523/JNEUROSCI.1467-13.2015

Chen, H., Kronengold, J., Yan, Y., Gazula, V. R., Brown, M. R., Ma, L., Ferreira, G., Yang, Y., Bhattacharjee, A., Sigworth, F. J., Salkoff, L., & Kaczmarek, L. K. (2009). The N-terminal domain of Slack determines the formation and trafficking of Slick/Slack heteromeric sodium-activated potassium channels. *J Neurosci*, *29*(17), 5654-5665. https://doi.org/10.1523/JNEUROSCI.5978-08.2009

Cole, B. A., Pilati, N., & Lippiat, J. D. (2021). Epilepsy-causing KCNT1 variants increase K_{Na}1.1 channel activity by disrupting the activation gate. *bioRxiv*, 2021.2009.2016.460601. https://doi.org/10.1101/2021.09.16.460601

Connors, B. W. (2017). Synchrony and so much more: Diverse roles for electrical synapses in neural circuits. *Dev Neurobiol*, *77*(5), 610-624. https://doi.org/10.1002/dneu.22493

Davis, G. W. (2013). Homeostatic signaling and the stabilization of neural function. *Neuron*, *80*(3), 718-728. https://doi.org/10.1016/j.neuron.2013.09.044

Debanne, D., Inglebert, Y., & Russier, M. (2019). Plasticity of intrinsic neuronal excitability. *Curr Opin Neurobiol*, *54*, 73-82. https://doi.org/10.1016/j.conb.2018.09.001

Debanne, D., & Russier, M. (2019). The contribution of ion channels in input-output plasticity. *Neurobiol Learn Mem*, *166*, 107095. https://doi.org/10.1016/j.nlm.2019.107095

Dilena, R., DiFrancesco, J. C., Soldovieri, M. V., Giacobbe, A., Ambrosino, P., Mosca, I., Galli, M. A., Guez, S., Fumagalli, M., Miceli, F., Cattaneo, D., Darra, F., Gennaro, E., Zara, F., Striano, P., Castellotti, B., Gellera, C., Varesio, C., Veggiotti, P., & Taglialatela, M. (2018). Early Treatment with Quinidine in 2 Patients with Epilepsy of Infancy with Migrating Focal Seizures (EIMFS) Due to Gain-of-Function KCNT1 Mutations: Functional Studies, Clinical Responses, and Critical Issues for Personalized Therapy. *Neurotherapeutics*, *15*(4), 1112-1126. https://doi.org/10.1007/s13311-018-0657-9

Dudok, B., Klein, P. M., & Soltesz, I. (2022). Toward Understanding the Diverse Roles of Perisomatic Interneurons in Epilepsy. *Epilepsy Curr*, *22*(1), 54-60. https://doi.org/10.1177/15357597211053687

Edelstein, A. D., Tsuchida, M. A., Amodaj, N., Pinkard, H., Vale, R. D., & Stuurman, N. (2014). Advanced methods of microscope control using muManager software. *J Biol Methods*, *1*(2). https://doi.org/10.14440/jbm.2014.36

Evely, K. M., Pryce, K. D., Bausch, A. E., Lukowski, R., Ruth, P., Haj-Dahmane, S., & Bhattacharjee, A. (2017). Slack K(Na) Channels Influence Dorsal Horn Synapses and Nociceptive Behavior. *Mol Pain*, *13*, 1744806917714342. https://doi.org/10.1177/1744806917714342

Favero, M., Sotuyo, N. P., Lopez, E., Kearney, J. A., & Goldberg, E. M. (2018). A Transient Developmental Window of Fast-Spiking Interneuron Dysfunction in a Mouse Model of Dravet Syndrome. *J Neurosci*, *38*(36), 7912-7927. https://doi.org/10.1523/JNEUROSCI.0193-18.2018

Gertler, T. S., Cherian, S., DeKeyser, J. M., Kearney, J. A., & George, A. L., Jr. (2022). K(Na)1.1 gain-of-function preferentially dampens excitability of murine parvalbumin-positive interneurons. *Neurobiol Dis*, *168*, 105713. https://doi.org/10.1016/j.nbd.2022.105713

Gertler, T. S., Thompson, C. H., Vanoye, C. G., Millichap, J. J., & George, A. L., Jr. (2019). Functional consequences of a KCNT1 variant associated with status dystonicus and early-onset infantile encephalopathy. *Ann Clin Transl Neurol*, *6*(9), 1606-1615. https://doi.org/10.1002/acn3.50847

Gibson, J. R., Beierlein, M., & Connors, B. W. (1999). Two networks of electrically coupled inhibitory neurons in neocortex. *Nature*, *402*(6757), 75-79. https://doi.org/10.1038/47035

Gibson, J. R., Beierlein, M., & Connors, B. W. (2005). Functional properties of electrical synapses between inhibitory interneurons of neocortical layer 4. *J Neurophysiol*, *93*(1), 467-480. https://doi.org/10.1152/jn.00520.2004

Hage, T. A., & Salkoff, L. (2012). Sodium-activated potassium channels are functionally coupled to persistent sodium currents. *J Neurosci*, *32*(8), 2714-2721. https://doi.org/10.1523/JNEUROSCI.5088-11.2012

Halabisky, B., Parada, I., Buckmaster, P. S., & Prince, D. A. (2010). Excitatory input onto hilar somatostatin interneurons is increased in a chronic model of epilepsy. *J Neurophysiol*, *104*(4), 2214-2223. https://doi.org/10.1152/jn.00147.2010

Hedrich, U. B., Liautard, C., Kirschenbaum, D., Pofahl, M., Lavigne, J., Liu, Y., Theiss, S., Slotta, J., Escayg, A., Dihne, M., Beck, H., Mantegazza, M., & Lerche, H. (2014). Impaired action potential initiation in GABAergic interneurons causes hyperexcitable networks in an epileptic mouse model carrying a human Na(V)1.1 mutation. *J Neurosci*, *34*(45), 14874-14889. https://doi.org/10.1523/JNEUROSCI.0721-14.2014

Heron, S. E., Smith, K. R., Bahlo, M., Nobili, L., Kahana, E., Licchetta, L., Oliver, K. L., Mazarib, A., Afawi, Z., Korczyn, A., Plazzi, G., Petrou, S., Berkovic, S. F., Scheffer, I. E., & Dibbens, L. M. (2012). Missense mutations in the sodium-gated potassium channel gene KCNT1 cause severe autosomal dominant nocturnal frontal lobe epilepsy. *Nat Genet*, *44*(11), 1188-1190. https://doi.org/10.1038/ng.2440

Hinckley, C. A., Zhu, Z., Chu, J. H., Gubbels, C., Danker, T., Cherry, J. J., Whelan, C. D., Engle, S. J., & Nguyen, V. (2023). Functional evaluation of epilepsy associated KCNT1 variants in multiple

cellular systems reveals a predominant gain of function impact on channel properties. *Epilepsia*. https://doi.org/10.1111/epi.17648

Hu, H., & Agmon, A. (2015). Properties of precise firing synchrony between synaptically coupled cortical interneurons depend on their mode of coupling. *J Neurophysiol*, *114*(1), 624-637. https://doi.org/10.1152/jn.00304.2015

Issa, N. P., Nunn, K. C., Wu, S., Haider, H. A., & Tao, J. X. (2023). Putative roles for homeostatic plasticity in epileptogenesis. *Epilepsia*, *64*(3), 539-552. https://doi.org/10.1111/epi.17500

Joiner, W. J., Tang, M. D., Wang, L. Y., Dworetzky, S. I., Boissard, C. G., Gan, L., Gribkoff, V. K., & Kaczmarek, L. K. (1998). Formation of intermediate-conductance calcium-activated potassium channels by interaction of Slack and Slo subunits. *Nat Neurosci*, *1*(6), 462-469. https://doi.org/10.1038/2176

Kim, G. E., Kronengold, J., Barcia, G., Quraishi, I. H., Martin, H. C., Blair, E., Taylor, J. C., Dulac, O., Colleaux, L., Nabbout, R., & Kaczmarek, L. K. (2014). Human slack potassium channel mutations increase positive cooperativity between individual channels. *Cell Rep*, *9*(5), 1661-1672. https://doi.org/10.1016/j.celrep.2014.11.015

Kuchenbuch, M., Nabbout, R., Yochum, M., Sauleau, P., Modolo, J., Wendling, F., & Benquet, P. (2021). In silico model reveals the key role of GABA in KCNT1-epilepsy in infancy with migrating focal seizures. *Epilepsia*, *62*(3), 683-697. https://doi.org/10.1111/epi.16834

Lane, B. J., Samarth, P., Ransdell, J. L., Nair, S. S., & Schulz, D. J. (2016). Synergistic plasticity of intrinsic conductance and electrical coupling restores synchrony in an intact motor network. *Elife*, *5*. https://doi.org/10.7554/eLife.16879

Large, A. M., Kunz, N. A., Mielo, S. L., & Oswald, A. M. (2016). Inhibition by Somatostatin Interneurons in Olfactory Cortex. *Front Neural Circuits*, *10*, 62. https://doi.org/10.3389/fncir.2016.00062

Lignani, G., Baldelli, P., & Marra, V. (2020). Homeostatic Plasticity in Epilepsy. *Front Cell Neurosci*, *14*, 197. https://doi.org/10.3389/fncel.2020.00197

Liu, Y., Zhang, F. F., Song, Y., Wang, R., Zhang, Q., Shen, Z. S., Zhang, F. F., Zhong, D. Y., Wang, X. H., Guo, Q., Tang, Q. Y., & Zhang, Z. (2022). The Slack Channel Deletion Causes Mechanical Pain Hypersensitivity in Mice. *Front Mol Neurosci*, *15*, 811441. https://doi.org/10.3389/fnmol.2022.811441

Lu, R., Bausch, A. E., Kallenborn-Gerhardt, W., Stoetzer, C., Debruin, N., Ruth, P., Geisslinger, G., Leffler, A., Lukowski, R., & Schmidtko, A. (2015). Slack channels expressed in sensory neurons control neuropathic pain in mice. *J Neurosci*, *35*(3), 1125-1135. https://doi.org/10.1523/JNEUROSCI.2423-14.2015

Ma, Y., Hu, H., Berrebi, A. S., Mathers, P. H., & Agmon, A. (2006). Distinct subtypes of somatostatin-containing neocortical interneurons revealed in transgenic mice. *J Neurosci*, *26*(19), 5069-5082. https://doi.org/10.1523/JNEUROSCI.0661-06.2006

Magloire, V., Mercier, M. S., Kullmann, D. M., & Pavlov, I. (2019). GABAergic Interneurons in Seizures: Investigating Causality With Optogenetics. *Neuroscientist*, *25*(4), 344-358. https://doi.org/10.1177/1073858418805002

Makinson, C. D., Dutt, K., Lin, F., Papale, L. A., Shankar, A., Barela, A. J., Liu, R., Goldin, A. L., & Escayg, A. (2016). An Scn1a epilepsy mutation in Scn8a alters seizure susceptibility and behavior. *Exp Neurol*, *275 Pt 1*(0 1), 46-58. https://doi.org/10.1016/j.expneurol.2015.09.008

Marder, E., & Goaillard, J. M. (2006). Variability, compensation and homeostasis in neuron and network function. *Nat Rev Neurosci*, *7*(7), 563-574. https://doi.org/10.1038/nrn1949

Markram, H., Muller, E., Ramaswamy, S., Reimann, M. W., Abdellah, M., Sanchez, C. A., Ailamaki, A., Alonso-Nanclares, L., Antille, N., Arsever, S., Kahou, G. A., Berger, T. K., Bilgili, A., Buncic, N., Chalimourda, A., Chindemi, G., Courcol, J. D., Delalondre, F., Delattre, V., . . . Schurmann, F.

(2015). Reconstruction and Simulation of Neocortical Microcircuitry. *Cell*, *163*(2), 456-492.
https://doi.org/10.1016/j.cell.2015.09.029

Martinez-Espinosa, P. L., Wu, J., Yang, C., Gonzalez-Perez, V., Zhou, H., Liang, H., Xia, X. M., & Lingle, C. J.
(2015). Knockout of Slo2.2 enhances itch, abolishes KNa current, and increases action potential
firing frequency in DRG neurons. *Elife*, *4*. https://doi.org/10.7554/eLife.10013

McTague, A., Nair, U., Malhotra, S., Meyer, E., Trump, N., Gazina, E. V., Papandreou, A., Ngoh, A.,
Ackermann, S., Ambegaonkar, G., Appleton, R., Desurkar, A., Eltze, C., Kneen, R., Kumar, A. V.,
Lascelles, K., Montgomery, T., Ramesh, V., Samanta, R., . . . Kurian, M. A. (2018). Clinical and
molecular characterization of KCNT1-related severe early-onset epilepsy. *Neurology*, *90*(1), e55-
e66. https://doi.org/10.1212/WNL.0000000000004762

Mikati, M. A., Jiang, Y. H., Carboni, M., Shashi, V., Petrovski, S., Spillmann, R., Milligan, C. J., Li, M., Grefe,
A., McConkie, A., Berkovic, S., Scheffer, I., Mullen, S., Bonner, M., Petrou, S., & Goldstein, D.
(2015). Quinidine in the treatment of KCNT1-positive epilepsies. *Ann Neurol*, *78*(6), 995-999.
https://doi.org/10.1002/ana.24520

Milligan, C. J., Li, M., Gazina, E. V., Heron, S. E., Nair, U., Trager, C., Reid, C. A., Venkat, A., Younkin, D. P.,
Dlugos, D. J., Petrovski, S., Goldstein, D. B., Dibbens, L. M., Scheffer, I. E., Berkovic, S. F., &
Petrou, S. (2014). KCNT1 gain of function in 2 epilepsy phenotypes is reversed by quinidine. *Ann
Neurol*, *75*(4), 581-590. https://doi.org/10.1002/ana.24128

Oliva, A. A., Jr., Jiang, M., Lam, T., Smith, K. L., & Swann, J. W. (2000). Novel hippocampal interneuronal
subtypes identified using transgenic mice that express green fluorescent protein in GABAergic
interneurons. *J Neurosci*, *20*(9), 3354-3368. https://doi.org/10.1523/JNEUROSCI.20-09-
03354.2000

Parker, P. R., Cruikshank, S. J., & Connors, B. W. (2009). Stability of electrical coupling despite massive
developmental changes of intrinsic neuronal physiology. *J Neurosci*, *29*(31), 9761-9770.
https://doi.org/10.1523/JNEUROSCI.4568-08.2009

Pratt, K. G., & Aizenman, C. D. (2007). Homeostatic regulation of intrinsic excitability and synaptic
transmission in a developing visual circuit. *J Neurosci*, *27*(31), 8268-8277.
https://doi.org/10.1523/JNEUROSCI.1738-07.2007

Quraishi, I. H., Stern, S., Mangan, K. P., Zhang, Y., Ali, S. R., Mercier, M. R., Marchetto, M. C., McLachlan,
M. J., Jones, E. M., Gage, F. H., & Kaczmarek, L. K. (2019). An Epilepsy-Associated KCNT1
Mutation Enhances Excitability of Human iPSC-Derived Neurons by Increasing Slack KNa
Currents. *J Neurosci*, *39*(37), 7438-7449. https://doi.org/10.1523/JNEUROSCI.1628-18.2019

Reijntjes, D. O. J., Lee, J. H., Park, S., Schubert, N. M. A., van Tuinen, M., Vijayakumar, S., Jones, T. A.,
Jones, S. M., Gratton, M. A., Xia, X. M., Yamoah, E. N., & Pyott, S. J. (2019). Sodium-activated
potassium channels shape peripheral auditory function and activity of the primary auditory
neurons in mice. *Sci Rep*, *9*(1), 2573. https://doi.org/10.1038/s41598-019-39119-z

Rizzi, S., Knaus, H. G., & Schwarzer, C. (2016). Differential distribution of the sodium-activated potassium
channels slick and slack in mouse brain. *J Comp Neurol*, *524*(10), 2093-2116.
https://doi.org/10.1002/cne.23934

Rizzo, F., Ambrosino, P., Guacci, A., Chetta, M., Marchese, G., Rocco, T., Soldovieri, M. V., Manocchio, L.,
Mosca, I., Casara, G., Vecchi, M., Taglialatela, M., Coppola, G., & Weisz, A. (2016).
Characterization of two de novoKCNT1 mutations in children with malignant migrating partial
seizures in infancy. *Mol Cell Neurosci*, *72*, 54-63. https://doi.org/10.1016/j.mcn.2016.01.004

Rudy, B., Fishell, G., Lee, S., & Hjerling-Leffler, J. (2011). Three groups of interneurons account for nearly
100% of neocortical GABAergic neurons. *Dev Neurobiol*, *71*(1), 45-61.
https://doi.org/10.1002/dneu.20853

Rush, A. M., Dib-Hajj, S. D., Liu, S., Cummins, T. R., Black, J. A., & Waxman, S. G. (2006). A single sodium
channel mutation produces hyper- or hypoexcitability in different types of neurons. *Proc Natl*
Acad Sci U S A, *103*(21), 8245-8250. https://doi.org/10.1073/pnas.0602813103
Schindelin, J., Arganda-Carreras, I., Frise, E., Kaynig, V., Longair, M., Pietzsch, T., Preibisch, S., Rueden, C.,
Saalfeld, S., Schmid, B., Tinevez, J. Y., White, D. J., Hartenstein, V., Eliceiri, K., Tomancak, P., &
Cardona, A. (2012). Fiji: an open-source platform for biological-image analysis. *Nat Methods*,
9(7), 676-682. https://doi.org/10.1038/nmeth.2019
Shore, A. N., Colombo, S., Tobin, W. F., Petri, S., Cullen, E. R., Dominguez, S., Bostick, C. D., Beaumont, M.
A., Williams, D., Khodagholy, D., Yang, M., Lutz, C. M., Peng, Y., Gelinas, J. N., Goldstein, D. B.,
Boland, M. J., Frankel, W. N., & Weston, M. C. (2020). Reduced GABAergic neuron excitability,
altered synaptic connectivity, and seizures in a KCNT1 gain-of-function mouse model of
childhood epilepsy. *Cell Rep*.
Soh, H., Park, S., Ryan, K., Springer, K., Maheshwari, A., & Tzingounis, A. V. (2018). Deletion of KCNQ2/3
potassium channels from PV+ interneurons leads to homeostatic potentiation of excitatory
transmission. *Elife*, *7*. https://doi.org/10.7554/eLife.38617
Spitznagel, B. D., Mishra, N. M., Qunies, A. M., Prael, F. J., 3rd, Du, Y., Kozek, K. A., Lazarenko, R. M.,
Denton, J. S., Emmitte, K. A., & Weaver, C. D. (2020). VU0606170, a Selective Slack Channels
Inhibitor, Decreases Calcium Oscillations in Cultured Cortical Neurons. *ACS Chem Neurosci*,
11(21), 3658-3671. https://doi.org/10.1021/acschemneuro.0c00583
Staley, K. (2015). Molecular mechanisms of epilepsy. *Nat Neurosci*, *18*(3), 367-372.
https://doi.org/10.1038/nn.3947
Studtmann, C., Ladislav, M., Topolski, M. A., Safari, M., & Swanger, S. A. (2022). Na(V)1.1
haploinsufficiency impairs glutamatergic and GABAergic neuron function in the thalamus.
Neurobiol Dis, *167*, 105672. https://doi.org/10.1016/j.nbd.2022.105672
Tai, C., Abe, Y., Westenbroek, R. E., Scheuer, T., & Catterall, W. A. (2014). Impaired excitability of
somatostatin- and parvalbumin-expressing cortical interneurons in a mouse model of Dravet
syndrome. *Proc Natl Acad Sci U S A*, *111*(30), E3139-3148.
https://doi.org/10.1073/pnas.1411131111
Tang, Q. Y., Zhang, F. F., Xu, J., Wang, R., Chen, J., Logothetis, D. E., & Zhang, Z. (2016). Epilepsy-Related
Slack Channel Mutants Lead to Channel Over-Activity by Two Different Mechanisms. *Cell Rep*,
14(1), 129-139. https://doi.org/10.1016/j.celrep.2015.12.019
Taniguchi, H., He, M., Wu, P., Kim, S., Paik, R., Sugino, K., Kvitsiani, D., Fu, Y., Lu, J., Lin, Y., Miyoshi, G.,
Shima, Y., Fishell, G., Nelson, S. B., & Huang, Z. J. (2011). A resource of Cre driver lines for
genetic targeting of GABAergic neurons in cerebral cortex. *Neuron*, *71*(6), 995-1013.
https://doi.org/10.1016/j.neuron.2011.07.026
Ting, J. T., Daigle, T. L., Chen, Q., & Feng, G. (2014). Acute brain slice methods for adult and aging
animals: application of targeted patch clamp analysis and optogenetics. *Methods Mol Biol*, *1183*,
221-242. https://doi.org/10.1007/978-1-4939-1096-0_14
Tran, C. H., Vaiana, M., Nakuci, J., Somarowthu, A., Goff, K. M., Goldstein, N., Murthy, P., Muldoon, S. F.,
& Goldberg, E. M. (2020). Interneuron Desynchronization Precedes Seizures in a Mouse Model
of Dravet Syndrome. *J Neurosci*, *40*(13), 2764-2775. https://doi.org/10.1523/JNEUROSCI.2370-
19.2020
Tremblay, R., Lee, S., & Rudy, B. (2016). GABAergic Interneurons in the Neocortex: From Cellular
Properties to Circuits. *Neuron*, *91*(2), 260-292. https://doi.org/10.1016/j.neuron.2016.06.033
Turrigiano, G. (2011). Too many cooks? Intrinsic and synaptic homeostatic mechanisms in cortical circuit
refinement. *Annu Rev Neurosci*, *34*, 89-103. https://doi.org/10.1146/annurev-neuro-060909-
153238

Turrigiano, G. (2012). Homeostatic synaptic plasticity: local and global mechanisms for stabilizing
neuronal function. *Cold Spring Harb Perspect Biol*, *4*(1), a005736.
https://doi.org/10.1101/cshperspect.a005736
Urban-Ciecko, J., & Barth, A. L. (2016). Somatostatin-expressing neurons in cortical networks. *Nat Rev*
Neurosci, *17*(7), 401-409. https://doi.org/10.1038/nrn.2016.53
Varghese, N., Moscoso, B., Chavez, A., Springer, K., Ortiz, E., Soh, H., Santaniello, S., Maheshwari, A., &
Tzingounis, A. V. (2023). KCNQ2/3 Gain-of-Function Variants and Cell Excitability: Differential
Effects in CA1 versus L2/3 Pyramidal Neurons. *J Neurosci*, *43*(38), 6479-6494.
https://doi.org/10.1523/JNEUROSCI.0980-23.2023
Wengert, E. R., Miralles, R. M., Wedgwood, K. C. A., Wagley, P. K., Strohm, S. M., Panchal, P. S., Idrissi, A.
M., Wenker, I. C., Thompson, J. A., Gaykema, R. P., & Patel, M. K. (2021). Somatostatin-Positive
Interneurons Contribute to Seizures in SCN8A Epileptic Encephalopathy. *J Neurosci*, *41*(44),
9257-9273. https://doi.org/10.1523/JNEUROSCI.0718-21.2021
Wu, J., Quraishi, I. H., Zhang, Y., Bromwich, M., & Kaczmarek, L. K. (2024). Disease-causing Slack
potassium channel mutations produce opposite effects on excitability of excitatory and
inhibitory neurons. *Cell Rep*, *43*(3), 113904. https://doi.org/10.1016/j.celrep.2024.113904
Yang, J., & Prescott, S. A. (2023). Homeostatic regulation of neuronal function: importance of
degeneracy and pleiotropy. *Front Cell Neurosci*, *17*, 1184563.
https://doi.org/10.3389/fncel.2023.1184563
Yu, F. H., Mantegazza, M., Westenbroek, R. E., Robbins, C. A., Kalume, F., Burton, K. A., Spain, W. J.,
McKnight, G. S., Scheuer, T., & Catterall, W. A. (2006). Reduced sodium current in GABAergic
interneurons in a mouse model of severe myoclonic epilepsy in infancy. *Nat Neurosci*, *9*(9),
1142-1149. https://doi.org/10.1038/nn1754
Zhang, J., Chen, X., Eaton, M., Wu, J., Ma, Z., Lai, S., Park, A., Ahmad, T. S., Que, Z., Lee, J. H., Xiao, T., Li,
Y., Wang, Y., Olivero-Acosta, M. I., Schaber, J. A., Jayant, K., Yuan, C., Huang, Z., Lanman, N.
A., . . . Yang, Y. (2021). Severe deficiency of the voltage-gated sodium channel Na(V)1.2 elevates
neuronal excitability in adult mice. *Cell Rep*, *36*(5), 109495.
https://doi.org/10.1016/j.celrep.2021.109495
Zhang, Q., Gao, S. H., Shen, Z. S., Wang, Y., Hu, S. W., Duan, G. B., Liu, Y., Zhong, D. Y., Liu, J., Sun, M. H.,
Zhang, X., Cao, T. Y., Cao, J. L., Tang, Q. Y., & Zhang, Z. (2022). The Slack Channel Regulates
Anxiety-Like Behaviors via Basolateral Amygdala Glutamatergic Projections to Ventral
Hippocampus. *J Neurosci*, *42*(14), 3049-3064. https://doi.org/10.1523/JNEUROSCI.2027-21.2022

1547

Figure 1

Figure 2

Figure 3

Paired neuron recordings of evoked PSCs onto excitatory (E: WT●, YH-HET●) and inhbitory (I: WT●, YH-HET●) neurons

A₁.

A₂.

A₃.

A₄.

Spontaneous PSCs onto excitatory (E: WT●, YH-HET●) and inhbitory (I: WT●, YH-HET●) neurons

B₁.

B₂.

B₃.

Figure 4

Figure 5

Figure 6

Figure 7

Figure 8

Figure 9

GFP-expressing GIN neurons (WT, YH-HET)

A.

B.

25 mV
200 ms

C.

WT (25)
YH-HET (25)

p<0.001

Current (pA)

APs/second

D.

p=0.082

Input Resistance (MΩ)

p<0.001

Rheobase (pA)

AP threshold (mV)

AHP (mV)

Supplemental Figure 2-1

Supplemental Figure 4-1

Supplemental Figure 4-2

A.

DIV 16 cortical SST neurons

WT

YH-HET

Neu-N *hSyn*-DIO-mCherry

Neu-N *hSyn*-DIO-mCherry

CamKII-GFP *hSyn*-DIO-mCherry

CamKII-GFP *hSyn*-DIO-mCherry

B.

Supplemental Figure 4-3

A.

B.

Supplemental Figure 5-1

Supplemental Figure 5-2

A.

VU-sensitive
TTX-sensitive

difference current (nA)

Voltage (mV)

B.

VU-sensitive
TTX-sensitive

difference current (nA)

Voltage (mV)

Supplemental Figure 6-1

Supplemental Figure 6-2

A.

Input resistance

Rheobase

AP firing rate

B.

Input resistance

Rheobase

AP firing rate

Supplemental Figure 7-1

Supplemental Figure 7-2