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An *Scn1a* epilepsy mutation in *Scn8a* alters seizure susceptibility and behavior

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Abstract

Understanding the role of SCN8A in epilepsy and behavior is critical in light of recently identified human SCN8A epilepsy mutations. We have previously demonstrated that Scn8a^{med} and Scn8a^{med-jo} mice carrying mutations in the Scn8a gene display increased resistance to flurothyl and kainic acid-induced seizures; however, they also exhibit spontaneous absence seizures. To further investigate the relationship between altered SCN8A function and epilepsy, we introduced the SCNIA-R1648H mutation, identified in a family with generalized epilepsy with febrile seizures plus (GEFS+), into the corresponding position (R1627H) of the mouse Scn8a gene. Heterozygous R1627H mice exhibited increased resistance to some forms of pharmacologically and electrically induced seizures and the mutant Scn8a allele ameliorated the phenotype of Scn1a-R1648H mutants. Hippocampal slices from heterozygous R1627H mice displayed decreased bursting behavior compared to wild-type littermates. Paradoxically, at the homozygous level, R1627H mice did not display increased seizure resistance and were susceptible to audiogenic seizures. We furthermore observed increased hippocampal pyramidal cell excitability in heterozygous and homozygous Scn8a-R1627H mutants, and decreased interneuron excitability in heterozygous Scn8a-R1627H mutants. These results expand the phenotypes associated with disruption of the Scn8a gene and demonstrate that an Scn8a mutation can both confer seizure protection and increase seizure susceptibility.

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Keywords

Nav1.6; Nav1.1; Audiogenic seizure; voltage sensor; sodium channel; GEFS+; Dravet Syndrome; interneuron

INTRODUCTION

Voltage-gated sodium channels (VGSCs) are important regulators of neuronal excitability and are responsible for the initiation and propagation of action potentials in neurons. Given their fundamental role in neuronal communication, disruptions in VGSC function can lead to a host of pathophysiological conditions. Most notably, mutations in the four VGSCs that are primarily expressed in the CNS—*SCN1A* (Na_v1.1), *SCN2A* (Na_v1.2), *SCN3A* (Na_v1.3), and *SCN8A* (Na_v1.6)—are responsible for several types of idiopathic epilepsy. Specifically, *SCN1A* mutations lead to genetic epilepsy with febrile seizures plus (GEFS+) and Dravet syndrome (DS) (Claes et al., 2001; Escayg et al., 2000), *SCN2A* mutations cause benign familial neonatal-infantile seizures (Heron et al., 2002), mutations in *SCN3A* have been identified in patients with partial epilepsy (Estacion et al., 2010; Holland et al., 2008; Vanoye et al., 2014), and *SCN8A* mutations are responsible for some cases of epileptic encephalopathies (EIEE13) (Carvill et al., 2013; Vaher et al., 2013; Veeramah et al., 2012).

Based on the VGSC mutations that have been identified to date, it is clear that distinct seizure and behavioral outcomes can result from different mutations in the same VGSC gene, with the magnitude and direction of the observed phenotypes potentially reflecting specific biophysical changes caused by the mutation. For example, within the spectrum of *SCN1A* epilepsies, the most severe condition, DS, often results from null mutations in the *SCN1A* gene, while amino acid substitutions that change the biophysical properties of the channel can give rise to less severe forms of epilepsy such as GEFS+ (Claes et al., 2009; Escayg and Goldin, 2010; Lossin, 2009).

SCN8A mutations were recently identified in several patients with epileptic encephalopathies (de Kovel et al., 2014; Estacion et al., 2014; O'Brien and Meisler, 2013; Vaher et al., 2013; Veeramah et al., 2012). Functional analyses initially suggested that increased channel activity was the likely biophysical consequence of these mutations (Estacion et al., 2014; Veeramah et al., 2012). However, the recent identification of potential loss-of-function *SCN8A* mutations in this disorder indicates that epilepsy might result from a range of alterations in *SCN8A* activity (de Kovel et al., 2014). The relationship between *SCN8A* dysfunction and disease outcome is also complex as illustrated by the cognitive and motor deficits, but not epilepsy, that were previously observed in a family with a *SCN8A* loss-of-function truncation mutation (Trudeau et al., 2006).

The introduction of clinically relevant VGSC mutations into mouse models has proven critical to our understanding of the functional consequences of distinct VGSC mutations on seizure and behavioral phenotypes. Mice expressing human *SCN1A* mutations recapitulate many of the clinical features of DS and GEFS+ (Martin et al., 2010; Yu et al., 2006), and have led to the "loss of inhibition" model of DS (Yu et al., 2006). Similarly, mice with loss-of-function *Scn8a* mutations exhibit behavioral alterations that are consistent with some

psychiatric findings in patients (McKinney et al., 2008; Papale et al., 2010). Interestingly, we demonstrated that mutations that reduce the activity of *Scn8a* in the mouse confer resistance to induced seizures (Hawkins et al., 2011; Martin et al., 2007), but result in the generation of absence epilepsy (Papale et al., 2009). Recently, the *SCN8A* N1768D gain-of-function mutation, identified in a patient with epileptic encephalopathy, was introduced into the mouse *Scn8a* gene. Mice expressing this mutation exhibited spontaneous seizures, reduced lifespan, impaired motor coordination and deficits in social interaction (Wagnon et al., 2014).

To gain insight into the spectrum of seizure and behavioral outcomes that can arise when the same mutation is expressed in different VGSC genes, we generated a mouse mutant in which the well-characterized *SCN1A* mutation R1648H, first identified in a family with GEFS+ (Escayg et al., 2000), was knocked into the corresponding location in the mouse *Scn8a* gene. The mutant line (*Scn8a*-R1627H) was evaluated for seizure susceptibility, spontaneous seizure generation, behavioral deficits, and neuronal excitability.

MATERIAL AND METODS

Generation of Construct and Electrophysiology in Xenopus oocytes

For expanded methods refer to Supplemental Materials and Methods. We introduced the *Scn1a*-R1648H mutation plus two additional silent substitutions into exon 26 of the *Scn8a* cDNA plasmid for expression in *Xenopus laevis oocytes*. Sodium currents were recorded by two-electrode voltage-clamp.

Generation of mice expressing the Scn8a R1627H mutation

A targeting construct consisting of a 4.5-kb 5' arm of homology, the R1627H substitution, a neomycin cassette flanked by *FLP1* recombinase (Fft) sites, and a 4-kb 3' arm of homology, was electroporated into 129X1/SvJ-derived PAT-5 embryonic stem (ES) cells at the University of Michigan Transgenic Core. PCR analysis and Southern blotting were performed to identify correctly targeted ES cells.

Animal breeding and maintenance

All experimental procedures were performed in accordance with the guidelines of Emory University and the University of California, Irvine Institutional Animal Care and Use Committees. All mice were housed in a temperature and humidity controlled vivarium and provided drinking water and food *ad libitum*. R1627H targeted mice were crossed to mice carrying FLP recombinase to remove the neomycin cassette included in the targeting vector. Male mutants were then crossed to female C57BL6/J wild-type (WT) mice for three generations. Heterozygous males and females from this generation (N3) were crossed to generate WT, heterozygous (*Scn8a^{RH/+}*), and homozygous (*Scn8a^{RH/RH}*) mice (N3F1) for experiments. A cohort of N3F1 mice were monitored, weighed, and videotaped weekly from P10 to P120 in order to assess locomotor function and to make general assessments of health and survival. Male *Scn8a^{RH/+}* mice (N3 generation) were also crossed to female C3/HeJ mice for three generations to generate progeny for EEG analysis. To generate mice carrying both the *Scn8a*-R1627H and the *Scn1a*-R1648H mutations, $Scn8a^{RH/+}$ (N3) females were first crossed to $Scn1a^{RH/+}$ (N10) males to produce $Scn8a^{RH/+}/Scn1a^{RH/+}$ offspring. $Scn8a^{RH/+}/Scn1a^{RH/+}$ males were then crossed to N10 $Scn1a^{RH/+}$ females to produce WT, $Scn8a^{RH/+}$, $Scn1a^{RH/+}$, $Scn8a^{RH/+}$, $Scn1a^{RH/+}$, $Scn1a^{$

Genotyping of mutants

Genotyping of R1627H mutant mice was performed by PCR analysis of DNA extracted from tail biopsies. A PCR product spanning the R1627H position was amplified using the following primers: (RH F, AAG ACA GGT TAT CTG TGT AAA CTG; RH R, AAT CGG TTT TGT CTG CAA GAC TGG) to produce a 600bp (R1627H) or a 500bp (WT) product. Genotyping of *Scn1a* R1648H mutant mice was performed as previously reported (Martin et al., 2010).

Protein extraction and Western blot analysis

Protein extraction and Western blot analysis was performed as previously described (Makinson et al., 2014). Detailed methods can be found in Supplemental Materials and Methods.

Animal behavior

All behavioral procedures were conducted between the hours of 10:00 AM and 3:00 PM to minimize possible circadian rhythm effects.

Locomotor activity was measured by photobeam breaks of individually housed mice (San Diego Instruments, La Jolla, CA, USA) over a 24-hour period. Ambulations were counted as consecutive beam breaks.

Rotarod performance was assessed by measuring the latency to fall from an accelerating rotating rod (Columbus Instruments, Rotamex-5 1.3). Each mouse was trained to walk on the rod three times per day for three days. Training trials were 5 minute duration at 5 r.p.m., with each training trial separated by 60 minutes. Mice that fell from the rod during the training trial were placed back on the rod. Following training, rod acceleration was increased in 0.2 r.p.m. increments every second starting at 0 r.p.m. and reaching a maximum of 50 r.p.m. Falls were detected by photobeam break.

Stride length was determined by measuring the distance between paw prints as mice walked across a 5 cm wide 60 cm long corridor. Mice were trained daily for 5 consecutive days to walk down the corridor. On training days, each mouse was placed at the beginning of the corridor and the home cage was placed on its side at the other end of the corridor. Once the mouse traversed the length of the corridor to the home cage, it was allowed to remain for at least 5 minutes before the next training trial. Three training trials were conducted on each day. Mouse forepaws were dipped in black ink so that a print would be left on white paper as the mouse traversed the corridor. Strides were measured from the tip of one paw print to the base of the next paw print on the same side. Stride length was measured in instances in

which at least 5 consecutive uninterrupted strides occurred. Average stride length was based on the analysis of at least 15 strides for each mouse.

Open field performance was measured by individually placing each mouse in an opaque Plexiglas box ($61 \text{ cm} \times 61 \text{ cm}$) for 5 minutes. The center of the open field was set 15 cm from the each side of the box. The latency to enter the center of the open field, the time spent in the center, and the number of entries into the center were measured.

The forced swim task was performed by placing each mouse in a beaker (diameter 30 cm) filled with water $27^{\circ}C \pm 1^{\circ}C$. The time spent struggling, defined as movements in which forelimbs broke the surface of the water, versus floating, defined as minimum movement required to remain afloat, were scored for the first 6 minutes.

The tail suspension task was performed by suspending each mouse from its tail at a height of 15 cm above a surface. Struggling was defined as forelimb and hindlimb movement or twisting movements. Mice were considered to be immobile if their forelimb and hindlimbs were not moving. Swinging was not scored as movement. Behavior was scored for the first 6 minutes.

The novel object recognition task was conducted as we previously described (Papale et al., 2010). Briefly, mice were acclimated to an open-field box for 10 minutes each day for five consecutive days. On the sixth day each animal was placed in the open-field for 5 minutes and then returned to the home cage for 2 minutes. Mice were then returned to the open field where they were given 5 minutes to explore three objects placed in three of the four corners of the box, at least 10 inches from the edges. Mice were returned to their home cage for either 5 or 20 minutes. Mice were then returned to the open-field for 5 minutes where one object had changed position (novel location), one object had been exchanged for a different object (novel object), and one object was the same (familiar object). The time spent exploring each object during the second exposure was recorded.

Video recordings of open field, forced swim, tail suspension, and novel object recognition tasks were analyzed using Any-Maze video tracking software (Stoelting Co. IL).

The elevated-plus maze consisted of four 30 cm arms (two open and two enclosed), suspended 76 inches above the floor. The test began with the mouse oriented towards an open arm. Each mouse was given 5 minutes to freely explore the apparatus during which time beam breaks were recorded to track movement and location. The percentage of time spent in the open arms was recorded and used as a measure of anxiety as mice naturally prefer the enclosed arms and anxiolytics have been shown to increase the time rodents spend in the open arm (Pellow et al., 1985). The percentage of time spent exploring the open arms was calculated by dividing the time spent in the open arms by the combined time spent in the open and closed arms. Total distance traveled in the apparatus was also recorded.

Sucrose preference was assessed by providing two food cups to each individually housed mouse. One cup contained between 1–2 g of standard rodent diet (Purina, Lab Diet 5001) the second cup contained 1–2 g of high sucrose diet (AIN-76A Rodent Tablets, TestDiet). The amounts of the high sucrose and standard diets consumed over a 1 hour period was

determined by subtracting the post-feeding weight of the food from the pre-feeding weight of the food. This procedure was repeated on 3 consecutive days. The reported sucrose preference values were determined on the third day.

The Y-maze task was performed by placing each mouse at the end of one arm of the Y-maze and allowing free exploration of the apparatus for 8 minutes. The Y-maze consists of three identical arms and a center zone. Spontaneous alternations were recorded.

Continuous video-electrocorticogram (EEG) monitoring

Video-EEG recordings to detect cortical seizure activity were collected and analyzed as we previously described (Dutton et al., 2012; Martin et al., 2007; Papale et al., 2009). EEG recordings from the inferior colliculus were collected as follows. Mice were anesthetized with isoflurane (2-chloro-2-(difluoromethoxy)-1,1,1-trifluoro-thane). Two sterile $0-80\times3/32$ sterile screw electrodes (Vintage Machine Supplies, Medina, OH) were implanted in the skull at the following coordinates from Bregma (anteroposterior (AP) 1.5 mm and mediolateral (ML) 1.2 mm; AP 1.0 mm and ML 1.2 mm). Stainless steel depth electrodes (diameter 0.25 mm, length 1.5 mm; Plastics One, Roanoke, VA) were implanted at the following coordinates (AP -3.5 mm, ML ± 1.5 mm, depth 1.2 mm from the surface of the skull). Two fine-wire electrodes were implanted in the neck muscle for electromyography (EMG) acquisition. Electrodes were covered with dental acrylic following implantation. After a minimum of five days for recovery from surgery, mice were placed in the recording chamber and connected to the EEG acquisition system via a flexible tether and commutator (Dragonfly inc.). Animals were allowed to freely move and were provided food and water ad libitum. EEG recordings were sampled at 200 Hz. Signals were digitized, amplified and processed by Stellate Harmonie EEG system (Natus Medical, Inc.). EEG traces were manually scored as previously described (Papale et al., 2013).

Seizure induction

Flurothyl seizure induction was performed as we previously described (Martin et al., 2007). Latencies to the first myoclonic jerk (MJ) and generalized tonic-clonic seizure (GTCS) were recorded. The MJ presents as a jerk of the upper body sometimes accompanied by tail-limb clonus. The GTCS was defined by complete loss of postural control and clonus of all limbs. Mice were also observed for the presence of hindlimb extension immediately following the GTCS.

Partial psychomotor seizures were evoked using the 6 Hz paradigm. Briefly, mice were administered a corneal analgesic 30 minutes prior to stimulation. Corneal electrical stimulation (6-Hz, 3 sec, 14 mA to 30 mA) was applied. Resulting seizures were scored according to the following modified Racine scale: 1 = staring, 2 = forelimb clonus, 3 = rearing and falling.

Audiogenic seizures were examined following auditory stimulation using either key ringing or a high-intensity tone. For the key ringing method, each mouse was placed in a clear Plexiglas box and keys were shaken for 5 minutes approximately 25 cm above the mouse. Shaking a key bundle was found to produce a broad-spectrum acoustic stimulus (13–85

kHz) at an intensity ranging from 50–80 dB, as determined by a 1/4" Brüel & Kjaer microphone (Brüel & Kjaer, Denmark). For the high-intensity tone method, a wide range speaker (Model RT1.3, HiVi inc., Arcadia, CA) was used to produce a 12 kHz, 80 dB tone. Each mouse was exposed to this tone for 5 min. Audiogenic seizures were defined by wild-running behavior progressing to loss of posture and forelimb and hindlimb clonus, sometimes followed by tonic hindlimb extension.

Electrophysiology

Mice P16–28 were deeply anesthetized with halothane, rapidly decapitated, and their brains were removed. Horizontal hippocampal slices were cut 350 µm thick with a vibratome (VT1000S; Leica Systems, Germany) in ice-cold sucrose-containing artificial cerebrospinal fluid (sACSF) (in mM: 85 NaCl, 75 sucrose, 2.5 KCl, 25 glucose, 1.25 NaH₂PO₄, 4 MgCl₂, 0.5 CaCl₂, and 24 NaHCO₃). Slices were incubated in oxygenated normal ACSF (in mM: 126 NaCl, 2.5 KCl, 26 NaHCO₃, 2 CaCl₂, 2 MgCl₂, 1.25 NaH₂PO₄, and 10 glucose) for 30 minutes to 1 hour at 27°C. All solutions used in preparation and recording were oxygenated by bubbling 95% O₂–5% CO₂.

Current clamp recordings were obtained using a MultiClamp 700B amplifier (Molecular Devices, Union City, CA), digitized with a Digidata 1322A digitizer (Molecular Devices), and data were acquired and analyzed with pClamp 10.2 software (Molecular Devices). Signals were sampled at 25 kHz and filtered at 10 kHz. The pipette solution contained the following (in mM): 126 K-gluconate, 4 KCl, 10 HEPES, 4 Mg-ATP, 0.3 Tris-GTP, and 10 Phospho-creatine, pH 7.2. The bath solution contained the following (in mM): 126 NaCl, 2 CaCl₂, 2 Mg Cl₂, 26 NaHCO₃, and 10 glucose, pH 7.3. Whole-cell recordings were obtained with access resistance <25 M Ω , and cells were held at -70 mV for all experiments. Cells were visualized using infrared DIC illumination under 40x magnification. Hyperpolarizing current injections of -10 pA and -30 pA were used to calculate cellular impedance. Firing patterns were recorded in response to 2-second depolarizing current injections in 20-pA increments, starting at 10 pA, up to 350 pA.

Field recordings were performed as previously described (Makinson et al., 2014). Slices were held at 33°C and perfused (at 2 mL/min) with oxygenated ACSF containing reduced Mg⁺⁺, 0.5 mM. Recording pipettes (2–3 M Ω) were pulled from borosilicate glass with a P-87 Flaming-Brown puller (Sutter Instruments, Novato, CA). Electrophysiological recordings were performed using a MultiClamp 700B amplifier (Molecular Devices, Union City, CA) and a Digidata 1322A digitizer (Molecular Devices). Data were acquired and analyzed with pClamp 10.2 software (Molecular Devices). Pipettes were filled with 150 mM NaCl and positioned in the CA3 stratum pyramidal layer. ACSF with elevated potassium was prepared by supplementing standard ACSF with 3 M KCl to raise the potassium concentration to 8.5 mM. The experimental paradigm consisted of a control recording for 2 min in standard ACSF, followed by 15–25 min with 8.5 mM K⁺ ACSF perfusion, during which burst activity was recorded. This was followed by a 7-minute washout with standard ACSF. Population spikes were elicited in the CA3 pyramidal layer by stimulation of the mossy fiber tract with a tungsten wire. Population spikes were evoked in physiological 2.5 mM [K⁺]_O before and after exposing the slice to 8.5 mM [K⁺]_O. Slices in which population

spike amplitudes did not change more than 20% after exposing the slice to 8.5 mM $[K^+]_O$ were used for analysis.

Auditory Brainstem Nuclei Response

Mice were anesthetized with ketamine and xylazine (i.p., 100 and 10 mg/kg respectively) and placed in a heated (25°C) sound attenuating booth (Industrial Acoustics Company, Bronx, NY). Auditory brainstem responses (ABRs) were recorded using Tucker Davis Technologies BioSigRP© software running on a System 3 platform equipped with an RX5 Pentusa Base Station connected to subdermal electrodes via an RA4LI low impedance headstage (TDT, Alachua, FL, USA). Sets of tone pips at different frequencies were presented in a random order to each animal to avoid possible ordering effects.

A subdermal needle electrode was placed over the skull vertex as the active lead. The ground was placed ventral lateral to the left external pinna and the reference was placed ventral lateral to the right external pinna. The bioelectric signals were sampled at 25 kHz, bandpass filtered between 100 Hz – 3000 Hz, amplified 200,000 times and averaged over 500 consecutive responses, following a previously established ABR screening protocol in mice (Zheng et al., 1999). To determine the ABR threshold, we reduced the stimulus intensity in 10 dB steps and then 5 dB steps until the lowest intensity at which the dominant ABR wave was visible.

Auditory stimuli

Calibrated stimuli were generated using TDT SigGenRP© software and presented through BioSigRP© software via a TDT RX6 digital signal processor at a sample rate of 195 kS/s. Sounds were attenuated by a TDT PA5 programmable attenuator and played from an Infinity EMIT tweeter placed 90° to the right side of the animal. Absolute sound pressure levels (SPL) of sound stimuli were measured prior to ABR recording experiments using a calibrated ¼" Bruel and Kjaer (B&K, Denmark) Type 4139 microphone with a Type 2669 preamplifier. Pure tone pips of 3 ms duration with 1.5 ms cos2 rise/fall times were presented for 5, 8, 10, 12, 15, and 24 kHz at a rate of 21 per second.

Immunohistochemistry

Following auditory stimulation, mice were transcardially perfused with ice-cold 4% paraformaldehide (PFA). Brains were extracted and stored in 30% sucrose for 5 days before slicing. Coronal slices (45 µm) were cut using a cryostat (Leica, Germany). Every fifth slice was washed in phosphate buffered saline (PBS) and incubated overnight in primary rabbit anti-c-Fos antibody (Abcam, 1:5,000). Slices were washed in PBS and incubated in horseradish peroxidase conjugated goat anti-rabbit secondary antibody (GE, 1:5,000) for 30 minutes. Staining was performed using the Vectastain Elite ABC System (Vector). Images were collected and stereological analysis performed using the Microbrightfield stereology system (MBF Bioscience, Williston, VT).

Statistics

All behavioral tests and latencies to flurothyl-induced seizures were analyzed using a oneway analysis of variance (ANOVA) followed by Dunnett's or Tukey's *post hoc* tests. Binary

seizure outcomes were analyzed using the χ^2 test. Weight data was analyzed using a twoway analysis of variance, genotype X time. ABRs were analyzed by two-way ANOVA or by the Student's *t-test*. Electrophysiology results were analyzed by Student's *t-test* or ANOVA (one or two-way) followed by Dunn's or Holm-Sidak correction.

RESULTS

The R1627H mutation alters recovery from inactivation, use-dependent inactivation, and persistent current

The effects of a mutation may not be the same in two different VGSCs either because the mutation has different effects on the biophysical properties of the two channels, or because the channels have different physiological functions. To determine how the R1627H mutation altered the properties of Na_v1.6 channels, WT and mutant channels were first expressed in *Xenopus* oocytes and examined by two-electrode voltage clamping. The channels were expressed with the β 1 subunit because β 1 might differentially modulate WT and mutant channels, as we and others have previously observed for mutations that cause GEFS+ (Spampanato et al., 2004). Sample sodium current traces through WT *Scn8a* and R1627H channels are shown in Fig. 1A and B, respectively. The R1627H mutation altered a number of properties in the presence of β_1 .

Mutant channels demonstrated a significantly more depolarized $V_{1/2}$ of activation $(-15\pm1mV)$ compared to WT channels $(-20\pm2mV)$ (Student's *t-test*, P < 0.05, n = 5 per group, Fig. 1C). The mutation also led to faster recovery from inactivation (Fig. 1D) and decreased use-dependent inactivation (Fig. 1F), which correlates with faster recovery and increased persistent current (two-way ANOVA, P<0.05, Fig. 1E). The biophysical changes caused by R1627H in Na_v1.6 are predicted to have variable effects on neuronal excitability, with the positive shift in voltage-dependence leading to a decrease in excitability and the faster recovery from inactivation and increased persistent current leading to an increase in excitability.

Targeted knock-in of the R1627H mutation into the mouse Scn8a gene

To determine the effects of the R1627H mutation in neurons and on the animal, we generated a mouse model by knocking in the human *SCN1A* R1648H GEFS+ mutation into the corresponding position (R1627) of the mouse *Scn8a* gene. Western blotting was performed on whole brain membrane enriched samples from homozygous *Scn8a^{RH/RH}*, heterozygous *Scn8a^{RH/+}*, and WT littermates (n = 3 per genotype). Protein levels of Na_v1.1, Na_v1.2, Na_v1.3, and Na_v1.6 were found to be comparable between WT, *Scn8a^{RH/+}* and *Scn8a^{RH/RH}* littermates, indicating that knock-in of the R1627H mutation did not lead to altered VGSC protein levels (Supplemental Fig. 1).

Survival, weight, and video monitoring of the R1627H line

Mice were monitored and weighed weekly from P10 to P120 to examine locomotor function and to make general assessments of health and survival. R1627H mice were born in approximately the expected Mendelian ratio (1:2:1) (WT = 24, $Scn8a^{RH/+}$ = 36, $Scn8a^{RH/RH}$ = 28) and had a normal lifespan. However, homozygous $Scn8a^{RH/RH}$ mutants gained weight

more slowly so that by 14 weeks of age the average weights of male and female $Scn8a^{RH/RH}$ mice were 5.6% (males) and 9.5% (females) less than WT and $Scn8a^{RH/+}$ sex-matched littermates, respectively (two-way ANOVA, P < 0.05, n = 11–18). While WT and $Scn8a^{RH/+}$ mice appeared visibly normal, tremors and uncoordinated gait were observed in $Scn8a^{RH/RH}$ mutants by the third postnatal week and persisted throughout the life of the animal. The hind legs of $Scn8a^{RH/RH}$ mice were rotated outward, possibly to provide a wider base for postural support, and repetitive "high stepping" movements often preceded locomotion. These findings are consistent with recessive motor abnormalities that have been reported in other Scn8a mutant mouse lines (Kohrman et al., 1996; Sprunger et al., 1999).

Scn8a^{RH/RH} mutant mice have reduced motor function

Recessive motor dysfunction has been described in other mouse models of *Scn8a* dysfunction. Therefore, we tested whether the *Scn8a*-R1627H mutation also results in altered motor function, even though we did not observe any motor abnormalities in mice with the *Scn1a* R1648H mutation (Martin et al., 2010). Motor function was assessed by measuring 24 hr locomotor activity, rotarod performance, and stride length. Significantly reduced locomotor activity was observed in *Scn8a^{RH/RH}* mice compared to *Scn8a^{RH/+}* and WT littermates in the first 5 hours of the recording period and at the beginning of the dark cycle, when mice are most active (two-way ANOVA, genotype X time, P < 0.05, n = 9–12, Fig. 2A). Average latency to fall from the rotarod was not significantly different between WT and *Scn8a^{RH/RH}* mice when compared to *Scn8a^{RH/RH}* mutants (one-way ANOVA, P < 0.05, n = 7–18, Fig. 2B). Shorter stride lengths were also observed in *Scn8a^{RH/RH}* mice when compared to *Scn8a^{RH/RH}* and WT littermates (one-way ANOVA, P < 0.001, n = 6–10, Fig. 2C–D).

The R1627H mutation is not associated with alterations in anxiety, depressive-like behavior, or learning and memory

Scn8a mutations have been associated with alterations in anxiety in mice (McKinney et al., 2008; Sawyer et al., 2014) and psychiatric and cognitive deficits in humans (Trudeau et al., 2006; Wang et al., 2008; Wasserman et al., 2005). However, anxiety levels, as measured by the percentage of time spent in the open arm or the number of entries into the open arm of an elevated plus maze, were comparable between $Scn8a^{RH/+}$, $Scn8a^{RH/RH}$ and WT littermates (Table 1). Likewise, no significant differences were observed between the three genotypes in the time spent in the center zone or the number of entries into the open field (Table 1). In the tail suspension task, $Scn8a^{RH/RH}$ mice spent significantly more time struggling and less time immobile, when compared to $Scn8a^{RH/+}$ and WT littermates (one-way ANOVA, P < 0.05, n = 8–10); however, no differences in time spent struggling or floating were observed in the forced swim task (one-way ANOVA, P > 0.05, n = 10 per group, Table 1). Likewise, preference for sucrose was comparable between $Scn8a^{RH/RH}$, $Scn8a^{RH/RH}$ and WT littermates (one-way ANOVA, P > 0.05, n = 8–12, Table 1), providing no evidence for depressive-like behavior in the mutant mice.

We previously reported modest improvement in spatial memory in mice with the *Scn8a^{medjo}* mutation (Papale et al., 2010). However, mice carrying the R1627H mutation were found to perform comparably to WT littermates in the novel object recognition task (one-way

ANOVA, P > 0.05, n = 8-12, Table 1). Spatial learning was also comparable between the three genotypes according to performance in the Y-maze task (one-way ANOVA, P > 0.05, n = 10-12, Table 1).

Scn8a^{RH/RH} and Scn8a^{RH/+} mice do not exhibit spontaneous seizures

We previously observed spontaneous absence seizures in three *Scn8a* mouse lines (Papale et al., 2009). To determine if the R1627H mutation also leads to seizure generation, ten days of continuous video-EEG recordings were collected from WT, *Scn8a*^{*RH/+*}, and *Scn8a*^{*RH/RH*} mice. No seizures were detected in any mice (n = 4–6). Based on previous work, the C3H/HeJ genetic background is known to be more permissive to the generation of absence seizures (Beyer et al., 2008). Therefore, we crossed *Scn8a*^{*RH/+*} mice to the C3H/HeJ background for five generations and then repeated the EEG analysis. No absence seizures were observed over five days of continuous video-EEG recordings collected from *Scn8a*^{*RH/+*} mutants and WT littermates at this generation (n = 5–6).

Scn8a^{RH/+} but not Scn8a^{RH/RH} mice are resistant to 6 Hz- and flurothyl-induced seizures

Average latency to the flurothyl-induced myoclonic jerk (MJ) and the generalized tonicclonic seizure (GTCS) were significantly increased in *Scn8a^{RH/+}* mice when compared to *Scn8a^{RH/RH}* and WT littermates (one-way ANOVA, P < 0.05, Fig. 3A). However, the severity of the seizures, as assessed by the presence of hindlimb extension and mortality following the GTCS, was reduced in both *Scn8a^{RH/+}* and *Scn8a^{RH/RH}* mice when compared to WT littermates (χ^2 , P < 0.05, Fig. 3B). Similarly, *Scn8a^{RH/+}* but not *Scn8a^{RH/RH}* mice were found to be resistant to 6 Hz-induced psychomotor seizures when compared to WT littermates (Fig. 3C), and the severity of the observed 6 Hz-induced seizures was lower in both *Scn8a^{RH/+}* and *Scn8a^{RH/RH}* mice when compared to WT littermates at current intensities of 20, 22, and 24 mA (one-way ANOVA, P < 0.0001, Dunnett's *post hoc*, P < 0.001, Fig. 3D).

Scn8a mutant mice have reduced hippocampal bursting activity in the presence of high extracellular potassium

To examine the susceptibility of the CNS network to seizure-like activity, extracellular recordings were performed in the CA3 region of hippocampal slices from WT, *Scn8a*^{*RH*/+} and *Scn8a*^{*RH*/*RH*} mice in the presence of high extracellular potassium (Fig. 4A). We previously observed reduced hippocampal bursting in *Scn8a*^{*med*/+} mice that carry a loss of function *Scn8a* mutation using the high [K⁺] seizure model (Makinson et al., 2014) and reduced hippocampal network excitability is thought to be a key component of *Scn8a*-related seizure resistance (Blumenfeld et al., 2009; Makinson et al., 2014). We found that the percentage of slices exhibiting bursting was significantly lower in *Scn8a*^{*RH*/+} mice (1/8, 12.5%) compared to WT (5/5, 100%) and *Scn8a*^{*RH*/*RH*} mice (4/7, 57.1%) (Fig. 4B). The average latency to the onset of bursting was comparable between slices from WT mice (6.1 min) and *Scn8a*^{*RH*/*RH*} mutants (6.3 min). However slices from *Scn8a*^{*RH*/+} did not show bursting even upon prolonged (> 20 min) exposure to elevated potassium, except for 1 slice (1/8) that did show bursting with a latency of 20 min (Fig. 4C). The intra-burst spiking frequency was reduced in that one *Scn8a*^{*RH*/+} slice (32.0 Hz) and also in slices from

Scn8a^{RH/RH} mice (49.3 Hz) when compared to WT slices (61.5 Hz) (Fig. 4D). These results indicate that slices from heterozygous *Scn8a* R1627H mice are resistant to seizure-like activity compared to WT, and slices from homozygous mice are intermediate between heterozygous and WT in their susceptibility to seizure-like activity.

Scn8a mutant mice have altered hippocampal cell firing

Based on oocyte data (Fig. 1), the effects of the R1627H mutation on sodium channel recovery and persistent current would be predicted to increase neuronal activity, whereas the effects on the voltage-dependence of activation would be predicted to decrease activity. Therefore, we examined the effect of the mutation on network excitability by current clamp in hippocampal slices. Consistent with the predictions based on the biophysical properties of the channel in oocytes, the R1627H mutation altered action potential firing in CA3 pyramidal neurons in a complex manner. Scn8a^{RH/+} neurons fired more action potentials than WT and Scn8a^{RH/RH} neurons, and Scn8a^{RH/RH} neurons fired more action potentials than WT. These differences were statistically significant (two-way ANOVA, P < 0.05, n = 21–28, Fig. 5A–B). The action potential height measured from threshold was significantly decreased in $Scn8a^{RH/RH}$ neurons compared to WT and $Scn8a^{RH/+}$ (two-way ANOVA, P < 0.05, n = 21–28, Table 2), and the action potential amplitude half-width was increased (broadened) for both $Scn8a^{RH/RH}$ and $Scn8a^{RH+}$ compared to WT (two-way ANOVA, P < 0.05, n = 21–28, Table 2). The threshold for firing an action potential was slightly higher for the mutants compared to WT, but the differences were not significantly different (two-way ANOVA, P > 0.05, n = 21–28, Table 2). Overall, pyramidal neurons from $Scn8a^{RH/+}$ and *Scn8a^{RH/RH}* mutants demonstrated greater excitability compared to WT. Hippocampal interneurons recorded from sub-pyramidal layers of CA3 in Scn8a^{RH/+} mice fired fewer action potentials compared to WT (two-way ANOVA, P < 0.05, n = 6-8, Fig. 5D), while no significant difference was observed between interneuron firing of Scn8a^{RH/RH} and WT neurons. Furthermore, no differences were observed in interneuron action potential properties (two-way ANOVA, P > 0.05, n = 6-8, Table 2). The resting membrane potentials of pyramidal neurons and interneurons across all genotypes were approximately -70 mV and -65 mV, respectively. There were no significant differences in the cell impedance of pyramidal and inhibitory neurons.

The Scn8a-R1627H mutation increases the lifespan and seizure thresholds of Scn1a-R1648H GEFS+ mice

The R1627H mutation in *Scn8a* is orthologous to the *Scn1a* R1648H mutation that causes GEFS+ in humans. To determine if there is a functional interaction between the two channels with orthologous mutations, we crossed the two lines to generate offspring with the following genotypes: $Scn1a^{RH/+}$, $Scn1a^{RH/RH}$, $Scn8a^{RH/+}$, $Scn1a^{RH/+}/Scn8a^{RH/+}$, $Scn1a^{RH/+}/Scn8a^{RH/+}$, $Scn1a^{RH/RH}/Scn8a^{RH/+}$, and WT. $Scn1a^{RH/+}$ and $Scn1a^{RH/RH}$ mice experienced 18% (2/11) and 100% (11/11) mortality by 3 months of age, respectively. In contrast, 40% of $Scn1a^{RH/RH}/Scn8a^{RH/+}$, and $Scn1a^{RH/+}/Scn8a^{RH/+}$ mice (Fig. 6A). As previously observed in WT, $Scn8a^{RH/+}$, and $Scn1a^{RH/+}/Scn8a^{RH/+}$ mice (Fig. 6A). As previously observed, the average latency to the flurothyl-induced GTCS was lower in $Scn1a^{RH/+}$ mutants; however, latencies to the GTCS were comparable between $Scn1a^{RH/+}/Scn8a^{RH/+}$ mutants and WT littermates (Fig. 5B). In contrast, significantly lower latencies to the MJ

and GTCS were still observed in $Scn1a^{RH/RH}/Scn8a^{RH/+}$ mutants (one-way ANOVA, Dunnett's *post hoc*, *P < 0.05, ***P <0.001, n = 5–17, Fig. 6B). The increase in lifespan and seizure thresholds is consistent with our previous results examining the interactions between the $Scn8a^{medjo}$ and GEFS+ or DS Scn1a mutants (Hawkins et al., 2011; Martin et al., 2007).

Scn8a^{RH/RH} mice exhibit wild running behavior in response to high intensity sound stimuli

 $Scn8a^{RH/RH}$ mice were sometimes observed exhibiting wild-running in response to elevated sound stimuli. To further explore this behavior, $Scn8a^{RH/+}$, $Scn8a^{RH/RH}$, and WT littermates were exposed to tones (12 kHz, 80 dB) or broad-spectrum (13–85 kHz, 50–80 dB) acoustic stimuli. Approximately 36% (11/31) of $Scn8a^{RH/RH}$ mice exhibited wild-running behavior followed by loss of postural control and forelimb and hindlimb clonus in response to sound stimuli (χ^2 , *P < 0.05, Fig. 7A). In contrast, only one WT littermate (1/15, 7%) exhibited wild running behavior in response to the sound stimulus. Furthermore, this WT mouse progressed to tonic hindlimb extension, which was not observed in the $Scn8a^{RH/RH}$ mutants. No $Scn8a^{RH/+}$ mice (0/10) exhibited abnormal audiogenic responses.

To investigate whether the observed abnormal behaviors may have been the result of audiogenic seizures, we recorded EEG activity in the inferior colliculus, which has been shown to be an important site for the initiation and maintenance of audiogenic seizures (Browning, 1986; Kesner, 1966; Willott and Lu, 1980). EEG activity in Scn8a^{RH/RH} mice was unaltered by exposure to the tone prior to the onset of the behavioral response (Fig. 7B). However at the onset of the behavioral response, we observed high amplitude - greater than twice baseline - EEG signals in the inferior colliculus of *Scn8a^{RH/RH}* mice (Fig. 7B). High amplitude EEG activity was not observed in cortical EEG recordings during the soundinduced behavioral response (Fig. 7B). To provide an independent measure of neuronal activity following sound exposure, we performed immunohistochemistry to compare c-Fos expression in the cortex, hippocampus, superior colliculus and inferior colliculus of responding and non-responding Scn8a^{RH/RH} mice as well as Scn8a^{RH/+} and WT controls (Fig. 7C-E). In agreement with the EEG results, low levels of c-Fos expression was observed in the cortex and hippocampus of all animals (Fig. 7D). High levels of c-Fos expression were observed in the superior and inferior colliculus of responding Scn8a^{RH/RH} mice but not non-responding $Scn8a^{RH/RH}$, WT, or $Scn8a^{RH/+}$ mice (Fig. 7D).

Scn8a^{RH/RH} mice have reduced auditory brainstem responses (ABRs)

To test the possibility that the sound-induced wild running behavior in $Scn8a^{RH/RH}$ mice may have been the result of increased hearing ability, ABRs to sound stimuli were recorded (Fig. 8). WT, $Scn8a^{RH/+}$ and $Scn8a^{RH/RH}$ mice were exposed to 5–24 kHz tones from 110–20 dB SPL while measuring ABR responses. $Scn8a^{RH/RH}$ mice were found to have higher thresholds between 12–15 kHz (two-way ANOVA, genotype X frequency, Dunnett's *post hoc*, *P < 0.05, Fig. 8A,C). However, no differences in ABR thresholds were observed between $Scn8a^{RH/RH}$ mice that were not susceptible to sound-induced wild-running behavior (non-responders) and $Scn8a^{RH/RH}$ mice that were found to be susceptible (responders) (Student's *t-test*, P > 0.05, Fig. 8B).

DISCUSSION

We constructed a mouse line in which the *SCN1A* GEFS+ mutation R1648H was introduced into the corresponding position in the mouse *Scn8a* gene (R1627H). The biophysical properties of the two sodium channels were altered by the mutation in similar but not identical ways. In oocytes, the R1627H mutation altered some properties of Na_v1.6 function that would be predicted to increase neuronal excitability, including faster recovery from inactivation, reduced use dependence and increased persistent current. These effects are consistent with those previously observed for R1648H (Lossin et al., 2002; Spampanato et al., 2001).

Action potential firing of pyramidal neurons was increased in heterozygous and homozygous R1627H mice compared to WT. These results contrast with our previous investigation of the Scn1a-R1648H mutation in which we observed no significant changes in excitability of pyramidal neurons from mutant compared to WT mice (Martin et al., 2010). Because the Scn1a-R1648H mutation also markedly decreased action potential firing in inhibitory bipolar neurons (Martin et al., 2010) and Scn8a expression has been detected at the axon initial segments of parvalbumin positive interneurons (Lorincz and Nusser, 2008), we also recorded from CA3 interneurons of *Scn8a*-R1627H animals. The only difference we observed in interneuron firing was a decrease in the number of action potentials from heterozygous mutants compared to WT mice. To assess the net effect of the Scn8a-R1627H mutation on network excitability, we examined seizure-like activity (bursting) in response to elevated extracellular potassium. Slices from the Scn8a^{RH/+} mice exhibited less epileptiform bursting activity (Fig. 4), consistent with increased seizure resistance. Interestingly, no change in the latency to burst onset was observed between Scn8a^{RH/RH} compared to WT slices, and more slices from Scn8a^{RH/RH} mice displayed bursting behavior compared to Scn8a^{RH/+} littermates, indicating a partial loss of seizure resistance at the homozygous level (Fig. 4). Similarly Scn8a^{RH/+} but not Scn8a^{RH/RH} animals were found to be resistant to flurothyl and electrically induced seizures (Fig. 3). It is not clear if the observed increase in CA3 pyramidal neuron firing and decrease in interneuron firing contributes directly to seizure resistance or if it results from compensatory responses to reduced network excitability in the heterozygous mice.

Consistent with findings from some other *Scn8a* mouse mutants, at the heterozygous level, the R1627H mutation increased resistance to induced seizures and compensated for the increase in seizure susceptibility resulting from the orthologous GEFS+ R1648H mutation in the *Scn1a* gene (Hawkins et al., 2011; Makinson et al., 2014; Martin et al., 2007).

A novel feature of the R1627H mice was their abnormal behavioral response to sound stimuli, which has not been previously reported in *Scn8a* mutants. Three observations suggest that the sound-induced wild running and forelimb-hindlimb clonic events represented audiogenic seizures. First, the behavioral presentation of these events, consisting of loss of posture and forelimb and hindlimb clonus, was similar to flurothyl-induced brainstem seizure behaviors (Samoriski et al., 1998). Second, following exposure to auditory stimuli, *Scn8a*^{*RH/RH*} mice that exhibited the abnormal behavior expressed high levels of c-Fos expression in the brainstem, including the inferior colliculus, a region responsible for

the initiation and maintenance of audiogenic seizures (Browning, 1986; Kesner, 1966; Willott and Lu, 1980). In contrast, c-Fos expression was not increased in the midbrain and brainstem of non-responding $Scn8a^{RH/RH}$, $Scn8a^{RH/+}$, or WT littermates and comparatively little c-Fos expression was observed outside of brainstem regions in $Scn8a^{RH/RH}$ responders following the auditory stimulus. c-Fos expression in brainstem nuclei following exposure to sound stimuli is a hallmark of audiogenic seizures (Klein et al., 2004; Kwon and Pierson, 1997; Le Gal La Salle and Naquet, 1990; Snyder-Keller and Pierson, 1992). Third, we observed increased amplitude of EEG signals in the inferior colliculus during the abnormal behavior in $Scn8a^{RH/RH}$ mice, indicating hypersynchronous neuronal activity in a major audiogenic seizure region. $Scn8a^{RH/RH}$ mice were also found to have higher ABR thresholds. This was similarly observed in the Scn8a mutant 'Cloth-ears', lending further support to the prediction that Scn8a dysfunction may contribute to neural hearing loss (Mackenzie et al., 2009).

Homozygous *Scn8a^{RH/RH}* mutants exhibited motor impairments and reduced locomotor activity, consistent with previously described *Scn8a* rodent models (Dickie, 1965; Hamann et al., 2003; Kearney et al., 2002; Kohrman et al., 1996). However, in contrast to previously described *Scn8a* mutants, R1627H mutants did not display alterations in measures of anxiety, or learning and memory (McKinney et al., 2008; Papale et al., 2010).

Unexpectedly, R1627H mutants did not exhibit absence seizures, which were observed in mice expressing *Scn8a^{8J}*, *Scn8a^{med}*, and *Scn8a^{medjo}* alleles (Papale et al., 2009). Across measurements of seizure susceptibility, the R1627H mutants contrast strikingly to the severe convulsive seizures and premature lethality observed in mice expressing the epileptic encephalopathy *SCN8A* N1768D mutation (Wagnon et al., 2014). Since electrophysiological analysis of neurons from *Scn8a* N1768D mutants has not yet been reported, a more direct comparison of the effect of each mutation on neuronal excitability cannot be performed. However, the striking phenotypic differences between these two mutants highlight the importance of studying mouse lines expressing different *SCN8A* mutations.

CONCLUSIONS

The increased seizure resistance conferred by the *Scn8a* R1627H mutation at the heterozygous level raises the possibility that seizure protection may be achieved by targeting specific biophysical properties of *Scn8a* function without elevating the risk of spontaneous seizure generation or negative behavioral outcomes. However, we report that the seizure resistance of the R1627H mutation was lost at the homozygous level and that these mice additionally were susceptible to audiogenic seizures, which coincided with increased pyramidal cell excitability but normal interneuron excitability. Together these observations indicate that *Scn8a* is an important regulator of excitatory as well as inhibitory cell excitability. This study expands upon the phenotypes associated with disruption of the *Scn8a* gene and demonstrates that an *Scn8a* mutation can both confer seizure protection and increase seizure susceptibility.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Highlights

- The R1627H mutation provides seizure resistance without causing absence epilepsy.
- *Scn8a*-dependent seizure phenotypes are gene-dose dependent.
- The R1627H mutation alters excitatory and inhibitory neuronal excitability in the hippocampus.
- The Scn8a-R1627H mutation increases susceptibility to audiogenic seizures



FIGURE 1.

Characterization of the biophysical properties of R1627H channels in *Xenopus* oocytes. A and B. Sample two-electrode voltage clamp recordings of currents through wild-type Na_v1.6 (A) and R1627H channels (B) with co-expression of β 1 during depolarizations between -50 and +50 mV in 10 mV increments. C. The R1627H mutation leads to a depolarizing shift in the V_{1/2} of voltage-dependent activation (Student's *t* test, P < 0.05). D. R1627H channels have accelerated recovery from inactivation (two-way ANOVA, Holm-Sidak correction, P < 0.05). E. At holding potentials from -30 to +40 mV, R1627H channels exhibit increased persistent current (two-way ANOVA, Holm-Sidak correction, P < 0.05). F. The R1627H mutation decreased use-dependent inactivation at 30 Hz (two-way ANOVA, Holm-Sidak correction, P < 0.05).



FIGURE 2.

Recessive motor impairment in R1627H mice. A. Locomotor activity of *Scn8a*-R1627H mice in a new cage was measured over a 24-hour period. A significant deficit in locomotor activity was detected between *Scn8a*^{*RH*/*RH*} mice and WT littermates (two-way ANOVA, genotype X time, P < 0.05). No significant differences were detected between *Scn8a*^{*RH*/*RH*} and WT littermates (P > 0.05). Inset graph shows reduced locomotor activity of *Scn8a*^{*RH*/*RH*} mice during the initial period in a novel environment (P < 0.05). White and black bars under the X-axis represent the light-dark cycle. n = 9–12. B. Rotarod performance was reduced in *Scn8a*^{*RH*/*RH*} (red) but not WT (black) or *Scn8a*^{*RH*/*+*} (blue) mice (one-way ANOVA, P < 0.001, Dunnett's *post hoc*, P < 0.05, n = 7–18). C. Representative example of paw prints showing reduced stride length in *Scn8a*^{*RH*/*RH*} mice. Black lines show relative stride length. D. Stride lengths were found to be significantly reduced in *Scn8a*^{*RH*/*RH*} but not *Scn8a*^{*RH*/*+*} mice when compared to WT littermates (one-way ANOVA, P < 0.001; Dunnett's *post hoc*, P < 0.05. n = 6–10.



FIGURE 3.

Effects of the R1627H mutation on flurothyl- and 6 Hz-induced seizure susceptibility. A. $Scn8a^{RH/+}$ but not $Scn8a^{RH/RH}$ mice have longer latencies to flurothyl-induced seizures when compared to WT littermates (one-way ANOVA, MJ and GTCS, Tukey *post hoc*, * P < 0.05, n = 11–12). B. Seizure severity, as measured by hindlimb extension (HE) and death, were reduced in $Scn8a^{RH/+}$ and $Scn8a^{RH/RH}$ mice (χ^2 , *P < 0.05). C. $Scn8a^{RH/+}$ mice are more resistant to 6 Hz seizures compared to $Scn8a^{RH/RH}$ and WT littermates. D. Both $Scn8a^{RH/+}$ and $Scn8a^{RH/RH}$ mice experience less severe seizures in response to stimulus intensities of 20, 22, and 24 mA compared to WT littermates (one-way ANOVA, P < 0.0001, Dunnett's *post hoc*, P < 0.001, n = 8–10).



FIGURE 4.

Effects of the R1627H mutation on hippocampal bursting. A. Example of bursting observed during field recording from WT CA3 when bath solution is switched from regular ACSF to high K⁺ ACSF. Bursting stops soon after high K⁺ ACSF is replaced with regular ACSF. B,The percentage of slices that exhibited bursting in the presence of high K⁺ was WT (5/5, 100%), *Scn8a^{RH/+}* (1/8, 12.5%), and *Scn8a^{RH/RH}* (4/7, 57.1%, *P < 0.05, Fisher Exact Test). C. The latency to the occurrence of bursting activity in the presence of high potassium (K⁺) was increased in slices from *Scn8a^{RH/+}* but not WT or *Scn8a^{RH/RH}* littermates. D. The frequency of intra-burst high amplitude spiking activity was reduced in *Scn8a^{RH/+}* compared to WT and *Scn8a^{RH/RH}* littermates. (B and C, data are average ± SEM, *P < 0.05, one-way ANOVA, Holm-Sidak *post-hoc* test).



FIGURE 5.

Firing properties of CA3 neurons from R1627H mice. A. Sample traces from CA3 pyramidal neurons of WT, $Scn8a^{RH/+}$ and $Scn8a^{RH/RH}$ mice at depolarizing current injections of 30 and 190 pA from a resting potential of -70mV. B. Average number of action potentials (AP#) plotted against current injection (pA) (Error bars represent SEM, n = number of cells ; *P < 0.05, two-way ANOVA, Holm-Sidak correction). C. Sample traces from CA3 interneurons of WT, $Scn8a^{RH/+}$ and $Scn8a^{RH/RH}$ mice at depolarizing current injections of 30 and 190 pA from a resting potential of -70 mV. B. Average number of action potentials (AP#) plotted against current injection (pA). Error bars represent SEM, n = number of cells, *P < 0.05 two-way ANOVA, Holm-Sidak correction.



FIGURE 6.

The *Scn8a*-R1627H mutation increases lifespan and seizure resistance in a mouse model of GEFS+ (*Scn1a*-R1648H). A. No mortality was observed over the first three months of postnatal development in WT, *Scn8a*^{RH/+}, and *Scn1a*^{RH/+}/*Scn8a*^{RH/+} mice. In contrast, *Scn1a*^{RH/+} and *Scn1a*^{RH/RH} mice experienced 18% (2/11) and 100% (11/11) mortality, respectively. *Scn1a*^{RH/RH}/*Scn8a*^{RH/+} mice exhibited 40% (4/10) survival. B. *Scn1a*^{RH/+} mutants (green) have lower latencies to flurothyl-induced GTCS, while *Scn8a*^{RH/+} mutants (orange) exhibit elevated latencies when compared to WT littermates. Normal latencies were restored in *Scn1a*^{RH/+}/*Scn8a*^{RH/+} mice (blue). *Scn1a*^{RH/RH}/*Scn8a*^{RH/+} mice (grey) have reduced latencies to the MJ and GTCS. One-way ANOVA, Dunnett's *post hoc*, *P < 0.05, ***P < 0.001. Error bars represent SEM, n = 5–17.



FIGURE 7.

Brainstem activity in *Scn8a^{RH/RH}* mice following sound-induced wild-running behavior. A. *Scn8a^{RH/RH}*, *Scn8a^{RH/+}*, and WT mice were exposed to 5-minute duration tones (12 kHz, 80 dB) or broad-spectrum (13–85 kHz, 50–80 dB) acoustic stimuli. Aproximately 7% (1/15) of WT, 0% (0/10) of *Scn8a^{RH/+}*, and 36% (11/31) of *Scn8a^{RH/RH}* mice exhibited wild running behavior during sound stimulus presentation (χ^2 , *P < 0.05). B. Representative example of inferior colliculus (IC) and cortical EEG activity in an *Scn8a^{RH/RH}* mouse before (baseline) and during abnormal behavoir. C. Representative image of c-Fos expression in the IC of a responding and non-responding *Scn8a^{RH/RH}* mouse 2 hours after exposure to sound stimulus. D. Increased c-Fos immunoreactive cells were observed in the IC of responding compared to non-responding *Scn8a^{RH/RH}* mice 2 hours following exposure to acoustic stimuli (Mann-Whitney U test, *P < 0.05, Error bars represent SEM, n = 3–4). E. Regional expression of c-Fos 2 hours after exposure to sound stimulus. Relative c-Fos expression: (+) low; (++) light; (+++) moderate; (++++) high. High levels of c-fos expression were observed in the superior and inferior colliculus of *Scn8a^{RH/RH}* responding animals but not in

non-responding $Scn8a^{RH/RH}$, $Scn8a^{RH/+}$ or WT littermates. Low to light levels of c-Fos was observed in the hippocampus and cortex of all animals.



FIGURE 8.

 $Scn8a^{RH/RH}$ mice have reduced auditory brainstem nuclei responses (ABRs). A. Auditory brainstem nuclei responses (ABR) were measured in response to 5–24 kHz tones in WT, $Scn8a^{RH/+}$ and $Scn8a^{RH/RH}$ mice. $Scn8a^{RH/RH}$ mice were found to have reduced auditory responses between 12–15 kHz (two-way ANOVA, Dunnett's *post hoc*, *P < 0.05, n = 8–10). B. No differences in ABR thresholds at 12 kHz were observed between $Scn8a^{RH/RH}$ mice that did not display wild-running behavior (non-responders) and $Scn8a^{RH/RH}$ mice that did exhibit abnormal behavior (responders) (Student's t test, P > 0.05, n = 4–8). C. Representative examples of ABR responses in WT and R1627H mutant mice. Black dot indicates the response threshold. Error bars represent SEM.

Summary of sur	vival and behavioral a	assessment of	the R1627H line.							
Category	Test	# of Animals	Measurement	Genotype	Average	s.e.m.	d	Post hoc Test	Comparison	Significance
				WT	0.0	ı			,	,
Survival	Survival	WI = 24 RH/+ = 36	4 mo % mortality	RH/+	0.0					
		KH/KH = 28		RH/RH	0.0	-				
				WT	25.1	3.8			,	
			% time in open arm	RH/+	26.9	2.6	0.84			ı
				RH/RH	28.6	5.5		•		1
		10 TT 11		WT	13.3	2.5				ı
	Elevated Plus Maze	WI = 10 RH/+ = 12	Open arm entries	RH/+	12.1	1.6	0.73		,	
		KH/KH = 10		RH/RH	10.8	2.4			,	ı
		-		WT	106.2	8.1			,	ı
			Distance traveled (cm)	RH/+	115.9	7.3	0.64		,	ı
				RH/RH	104.6	12.6		•		ı
Anxiety				WT	5.3	5.5			,	
			Time in center (sec)	RH/+	7.9	8.5	0.05		,	ı
				RH/RH	1.1	2.0			,	ı
				WT	2.8	2.1			,	ı
	Open Field	WI = 8 RH/+ = 12	Number of entries into the center	RH/+	3.6	2.5	p < 0.01	Dunnett's Test	WT vs. RH/+	p > 0.05
		KH/KH = 10		RH/RH	0.6	0.7			WT vs. RH/RH	p > 0.05
				ΤW	67.3	86.4				
			Latency to first center entry (sec)	RH/+	132.0	93.9	0.23	ı		
				RH/RH	65.6	108.6				
				ΤW	90.1	10.3				
			Time swimning (sec)	RH/+	83.6	11.2	0.57			
	Econod Conjun	WT = 10		RH/RH	99.5	8.6				ı
Depression-like	LUICED SWIII	RH/RH = 10 RH/RH = 10		WT	122.2	22.3				ı
			Time floating (sec)	RH/+	121.5	24.5	0.2	ı	1	I
				RH/RH	71.2	15.8			ı	

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TABLE 1

Category	Test	# of Animals	Measurement	Genotype	Average	s.e.m.	a	Post hoc Test	Comparison	Significance
				TW	170.7	8.4				1
			Time struggling (sec)	RH/+	171.8	10.6	p < 0.01	Dunnett's Test	WT vs. RH/+	p > 0.05
	To Provide American	WT = 10		RH/RH	247.7	7.2			WT vs. RH/RH	p < 0.05
	I all Suspension	RH/H = 10 RH/RH = 10		ΜT	229.5	5.0				
			Time immobile (sec)	RH/+	220.7	12.9	p < 0.05	Dunnett's Test	WT vs. RH/+	p > 0.05
				RH/RH	164.1	8.5			WT vs. RH/RH	p < 0.05
		$\mathbf{WT} = 11$		WT	96.9	2.9			1	ı
	Sucrose Preference	WI = 11 $RH/+ = 10$	Sucrose preference (sucrose g/ total g) *100	RH/+	97.0	1.9	0.64		-	
		KH/KH = 11	ò	RH/RH	89.5	10.7				
				ΜT	59.7	4.9			,	
			Novel object preference	RH/+	62.1	6.8	0.77			
	Novel Object	WT = 8	-	RH/RH	55.1	10.9			,	
	kecognuon (20 mm exposure)	RH/t = 10 RH/RH = 12		ΜT	56.2	4.9				
			Novel position preference	RH/+	62.6	6.8	0.41		,	
				RH/RH	64.8	10.9			-	
				ΜT	69.69	10.3				
			Novel object preference	RH/+	58.5	8.8	0.923		,	
Tomino	Novel Object	WT = 8		RH/RH	66.4	7.8			-	
Leaning/Meniory	exposure)	RH/RH = 12		WT	61.9	10.8			I	I
			Novel position preference	RH/+	60.6	5.9	0.136		I	
				RH/RH	70.9	13.0			-	
				ΤW	59.3	2.2			-	
			% correct rotations	RH/+	56.8	1.4	0.031	Dunnett's Test	WT vs. RH/+	p > 0.05
	V Marro	WT = 10		RH/RH	64.2	6.4			WT vs. RH/RH	p > 0.05
	I -IVIAZE	RH/RH = 10 RH/RH = 10		ΜT	331.4	19.3			-	ı
			Distance traveled (cm)	RH/+	352.3	18.7	0.021	Dunnett's Test	WT vs. RH/+	p > 0.05
				RH/RH	264.3	27.11			WT vs. RH/RH	p > 0.05
Mice carrying the R10	627H mutation and WT litter	rmates were evalu	ated for survival and were subjected	d to a battery o	of behavioral	tests in or	der to evalu	aate anxiety, depres	sive-like, and learni	ng and

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Intrinsic properties of hippocampal neurons from R1627H mice.

	AP	properties of pyram	iidal cells		AF	properties of inter	neurons	
	Amplitude (mV)	Threshold (mV)	Half-width (ms)	N	Amplitude (mV)	Threshold (mV)	Half-width (ms)	Z
ΨT	87 ± 2	-45 ± 1	2 ± 0.1	21	80 ± 5	-49 ± 2	2 ± 0.2	8
RH/+	88 ± 2	-46 ± 1	$2.3 \pm 0.1^*$	28	81 ± 4	-43 ± 2	2.5 ± 0.1	8
RH/RH	$78 \pm 3^{*}$	-46 ± 2	$2.7\pm0.1^*$	23	80 ± 4	-45 ± 3	2.6 ± 0.4	9

Summary of intrinsic membrane properties of hippocampal pyramidal cells and interneurons obtained by current clamp recording (one-way ANOVA, Holm-Sidak test).

 $^{*}_{P < 0.05.}$

Values represent average \pm SEM, n = number of cells,

Makinson et al.