Circuit-Specific Early Impairment of Proprioceptive Sensory Neurons in the SOD1^{G93A} Mouse Model for ALS

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Amyotrophic lateral sclerosis (ALS) is a neurodegenerative disease in which motor neurons degenerate, resulting in muscle atrophy, paralysis, and fatality. Studies using mouse models of ALS indicate a protracted period of disease development with progressive motor neuron pathology, evident as early as embryonic and postnatal stages. Key missing information includes concomitant alterations in the sensorimotor circuit essential for normal development and function of the neuromuscular system. Leveraging unique brainstem circuitry, we show *in vitro* evidence for reflex circuit-specific postnatal abnormalities in the jaw proprioceptive sensory neurons in the well-studied SOD1^{G93A} mouse. These include impaired and arrhythmic action potential burst discharge associated with a deficit in Nav1.6 Na⁺ channels. However, the mechanoreceptive and nociceptive trigeminal ganglion neurons and the visual sensory retinal ganglion neurons were resistant to excitability changes in age-matched SOD1^{G93A} mice. Computational modeling of the observed disruption in sensory patterns predicted asynchronous self-sustained motor neuron discharge suggestive of imminent reflexive defects, such as muscle fasciculations in ALS. These results demonstrate a novel reflex circuit-specific proprioceptive sensory abnormality in ALS.

Key words: amyotrophic lateral sclerosis; dynamic clamp; motor neurons; persistent sodium; proprioceptive neurons; SOD1

Significance Statement

Neurodegenerative diseases have prolonged periods of disease development and progression. Identifying early markers of vulnerability can therefore help devise better diagnostic and treatment strategies. In this study, we examined postnatal abnormalities in the electrical excitability of muscle spindle afferent proprioceptive neurons in the well-studied SOD1^{G93A} mouse model for neurodegenerative motor neuron disease, amyotrophic lateral sclerosis. Our findings suggest that these proprioceptive sensory neurons are exclusively afflicted early in the disease process relative to sensory neurons of other modalities. Moreover, they presented Nav1.6 Na⁺ channel deficiency, which contributed to arrhythmic burst discharge. Such sensory arrhythmia could initiate reflexive defects, such as muscle fasciculations in amyotrophic lateral sclerosis, as suggested by our computational model.

Introduction

Amyotrophic lateral sclerosis (ALS) is a neurodegenerative disease in which motor neurons throughout the brain and spinal

cord progressively degenerate. In this devastating disease, motor neuron (MN) degeneration leads to muscle paralysis and atrophy, and death ensues 3–5 years following clinical disease onset (Cleveland and Rothstein, 2001; Bruijn et al., 2004). Use of transgenic mouse models of ALS has provided key insights into presymptomatic mechanisms of disease development. These include central mechanisms encompassing glutamatergic excitotoxicity at MN synaptic terminals, diminished energy supply resulting from metabolite deficiency, and dysfunctional RNA metabolism, protein homeostasis, and aggregation (Bruijn et al., 2004; Taylor

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et al., 2016). Additional non–cell-autonomous contributors, such as dysfunctional astrocytes (Yamanaka et al., 2008), microglia (Boillée et al., 2006b), and oligodendrocytes (Lee et al., 2012), provide evidence that ALS disease mechanisms are not limited to MNs. Peripheral neural dysfunction, such as impairment in axonal transport (Williamson and Cleveland, 1999; Puls et al., 2003), and synaptic pruning/failure at the neuromuscular junctions (Pun et al., 2006; Casas et al., 2016), also plays a crucial role in initiating muscle paralysis. Importantly, these diverse aspects of early disease development collectively support the idea that additional neuronal pathways, such as early sensory input changes, are worth exploring as these may provide opportunities for identifying novel early biomarker and therapeutic targets.

Vulnerability to neurodegeneration is not the same for all MNs and within motor pools in ALS (Pun et al., 2006; Kanning et al., 2010). Both in normal aging and in ALS, motor unit physiology is a crucial determinant of preferential vulnerability, where fast fatigable motor units degenerate first, followed by fast fatigue resistant, whereas slow motor units are relatively spared (Pun et al., 2006; Hegedus et al., 2007). Second, MNs that control eye movements (oculomotor, trochlear, and abducens) and sphincter muscles (Onuf's nucleus), as well as gamma MNs that innervate intrafusal muscle fibers are selectively resistant to death (Iwata and Hirano, 1978; Ferrucci et al., 2010; Lalancette-Hebert et al., 2016). Comparative analyses have highlighted important differences in transcriptional profiles, electrical and synaptic properties, and neuromuscular biology and innervation patterns between the resistant and vulnerable MNs (Frey et al., 2000; Nimchinsky et al., 2000; Hedlund et al., 2010; Comley et al., 2015; Venugopal et al., 2015; Nijssen et al., 2017). These inherent differences indicate local and long-range defects in sensorimotor circuits of vulnerable MNs in ALS (Durand et al., 2006). For instance, a circuit-specific difference in the ALS-resistant oculomotor neurons involves a lack of Ia muscle spindle afferent proprioceptive inputs (Spencer and Porter, 1988), which are a principal source of glutamatergic excitation during muscle stretch reflexes. A lack of Ia spindle afferent inputs to spinal gamma MNs was suggested as a mechanism of disease resistance in ALS mouse models (Schneider et al., 2012). Consistent with that observation, reducing Ia proprioceptive muscle spindle afferents partly contributed to α -MN survival. Therefore, identification and characterization of circuit-specific vulnerability to disease progression in ALS (Brownstone and Lancelin, 2018) could help develop effective therapeutic strategies.

To test whether circuit-specific dysfunction involves proprioceptive sensory neurons, we leveraged the unique architecture of the brainstem trigeminal sensorimotor circuitry involved in jaw control. We examined the proprioceptive Ia afferents in the pontine mesencephalic nucleus (Mes V) in the SOD1^{G93A} mouse model at P11 $\pm$ 3 when the jaw motor pools are reported to be dysregulated (Venugopal et al., 2015). Our results show that Mes V neurons present electrical abnormalities and arrhythmic burst discharge patterns, associated with a marked reduction in Nav1.6-type Na⁺ currents. Rescue of these Na⁺ currents restored normal rhythmic burst patterns (see also Venugopal et al., 2019). Concomitant examination of trigeminal ganglion neurons and retinal ganglion neurons (RGNs) confirmed exclusive changes only in the proprioceptive Mes V neurons. Using a computational modeling approach, we show a functional consequence of sensory abnormality on downstream motor integration predictive of imminent reflex dysfunction in ALS.

Materials and Methods

Transgenic mice expressing high levels of human SOD1^{G93A} (mutant SOD1 or mSOD1) and their WT littermates were used for all the experiments (JAX strain: 002726 B6SJL-Tg (SOD1*G93A)1Gur/J). All animal protocols were approved by the Institutional Animal Care and Use Committee at UCLA. Experiments were performed at postnatal week 2 (8- to 14-d-old mice of either gender) when the rhythmic jaw movements and suckling behavior are fully developed (Turman, 2007). Genotype of mice was determined by standard PCR technique using tails samples (Laragen). Experimental preparations and techniques include the following: (1) live brainstem slices to conduct *in vitro* whole-cell current-clamp, voltage-clamp, and dynamic-clamp electrophysiology from Mes V sensory neurons; (2) acutely dissociated trigeminal ganglion neurons to conduct current-clamp experiments; (3) live whole retinal preparation to conduct current-clamp experiments; (4) fixed cryosectioned coronal pontine sections for Nav1.6 protein quantification; and (5) computational model of Mes V-TMN network to investigate a functional consequence of sensory abnormality on motor discharge.

In vitro patch-clamp electrophysiology

Brainstem slice preparation for Mes V electrophysiology

Brain slices were prepared and used for conducting whole-cell current-, voltage-, and dynamic-clamp electrophysiology in the brainstem primary sensory neurons of the trigeminal Mes V nucleus. Pups were anesthetized using isoflurane vapor inhalation and decapitated. The head was immediately immersed in carboxygenated (95% O₂/5% CO₂), ice-cold sucrose cutting solution composed of the following (in mM): 194 sucrose, 30 NaCl, 4.5 KCl, 1.2 NaH₂PO₄, 26 NaHCO₃, 10 glucose, and 1 MgCl₂. The pontine brainstem was rapidly extracted and adhered to the cutting chamber of a vibratome platform at the rostral end (DSK Microslicer; Ted Pella); the brainstem was vertically supported by an agar block. The cutting chamber was filled with ice-cold carboxygenated cutting solution. Beginning at the caudal level where the exit of the facial nerve was markedly visible, 3 or 4 coronal pontine slices (~250 μ m thick) were cut and placed in the carboxygenated incubation solution at room temperature, composed of the following (in mM): 124 NaCl, 4.5 KCl, 1.2 NaH₂PO₄, 26 NaHCO₃, 10 glucose, 2 CaCl₂, and 1 MgCl₂ (Schurr et al., 1988). The pH of the incubation solution was maintained at 7.28 $\pm$ 0.2.

Trigeminal ganglia (TG) extraction and acute dissociation of trigeminal ganglion neurons (TGNs) for electrophysiology

To evaluate excitability changes in the nonproprioceptive neurons of the trigeminal system of the mSOD1 mice, we performed acute dissociation of the TG (Malin et al., 2007; Xu et al., 2010; Yamamoto et al., 2013). Pups were decapitated under isoflurane anesthesia similar to that in Brainstem slice preparation for Mes V electrophysiology. The TG were bilaterally removed with the aid of a dissection microscope and transferred into ice-cold (4°C) modified Tyrode's solution containing the following (in mM): 130 NaCl, 20 NaHCO₃, 3 KCl, 4 CaCl₂, 1 MgCl₂, 10 HEPES, and 12 glucose, with antibiotic/antimycotic solution (0.5%; Thermo Fisher Scientific). The ganglia were then minced and incubated in collagenase (1 mg/ml, Type I; Thermo Fisher Scientific) for 40 min and then in collagenase with trypsin/EDTA (0.2%; Thermo Fisher Scientific) for another 40 min at 37°C. The TG cells were then washed twice with the modified Tyrode's solution and triturated gently using fire-polished Pasteur glass pipettes. Finally, the cell suspension was mixed with BSA (15%; Thermo Fisher Scientific) and centrifuged at 900 rpm for 10 min to remove myelin and debris. The pellet was resuspended with Neurobasal A (Thermo Fisher Scientific) containing B27 (2%; Thermo Fisher Scientific), L-glutamine (0.2%; Thermo Fisher Scientific), and antibiotic/antimycotic solution (0.1%), and cells were plated onto glass coverslips coated with poly-D-lysine/laminin (Thermo Fisher Scientific). The cells were then incubated at 37°C in a humidified 5% CO₂ chamber, and whole-cell patch-clamp electrophysiology was conducted ~24 h after plating (Chen et al., 2008; Marchenkova et al., 2016).

Retina extraction and preparation for electrophysiology

To ascertain whether visual sensory neurons are susceptible to early changes in excitability in ALS mice, we extracted whole retinas from

mSOD1 and WT mice for patch-clamp electrophysiology. For retinal extraction, pups were deeply anesthetized using isoflurane inhalation, and the eyes were denucleated and placed in an ice-cold solution containing the following (in mM): in room temperature-oxygenated solution containing the following (in mM): 124 NaCl, 4.5 KCl, 1.2 NaH₂PO₄, 26 NaHCO₃, 10 glucose, 2 CaCl₂, and 1 MgCl₂. The pH of the incubation solution was maintained at 7.28 ± 0.2 , with osmolarity adjusted to 300 ± 5 mOsm. Each retina was exposed by a single cut along the ora serrata. An additional cut was made along the optic disk to dissect the eyecup into two halves (Wang et al., 1997; Qu and Myhr, 2011). The retina was isolated from pigment epithelium and was stored for 1 h in oxygenated incubation solution at room temperature before whole-cell patch-clamp recording.

Whole-cell current-clamp recording

Whole-cell current-clamp recording was performed in three sets of sensory neurons: the trigeminal proprioceptive Mes V neurons, trigeminal ganglion mechanoreceptive/nociceptive neurons, and the RGNs. Axopatch-1D patch-clamp amplifier (Molecular Devices) and pCLAMP acquisition software (version 9.2, Molecular Devices) were used. Silver chloride electrode (Ag/AgCl wire) and 3 mM KCl agar-bridge were used for suitable grounding and electrical isolation. Patch electrodes were fabricated from borosilicate glass capillary tubing (1.5 mm OD, 0.86 mm ID) using a model P-97 puller (Sutter Instrument). Tip resistance was 3–5 M Ω when filled with a pipette internal solution containing the following (in mM): 135 K-glucuronate, 5 KCl, 0.5 CaCl₂, 5 HEPES (base), 5 EGTA, 2 Mg-ATP, and 0.3 Na-ATP (Gupta et al., 2012), pH 7.25 ± 0.2 , osmolarity adjusted to 290 ± 5 mOsm. The external recording solution consisted of the following (in mM): 140 NaCl, 4 KCl, 2 CaCl₂, 10 HEPES (base), 2 MgCl₂, and 10 glucose.

Mes V neuron identification. The proprioceptive Mes V nucleus was identified bilaterally in the coronal slices as a dorsally located ellipsoid region, ventrolateral to the aqueduct in the caudal pontine slices at the level of VII nerve exit, and in the lateral periphery of the periaqueductal gray in further rostral pontine slices (Wu et al., 2001; Enomoto et al., 2006; Seki et al., 2017). The Mes V neurons were easily distinguished on the basis of their large, ellipsoid soma with ~ 30 – 50 μ m diameter (Henderson et al., 1982; Del Negro and Chandler, 1997; Enomoto et al., 2006). In a subset of experiments, sensory Mes V neurons were identified using anterograde labeling by injection of 1% AlexaFluor-568 (Thermo Fisher Scientific) in the jaw closer muscles (20 μ l on each side) (see Fig. 1).

TGN identification. The dissociated TGNs were plated on a coverslip, and cells were identified for whole-cell patch-clamp recording using differential interference contrast microscopy. Medium to large isolated cell bodies with diameter > 10 μ m were selected. During initial experiments, the dissociated cells were labeled using NeuN stain (Neurotrace) for morphology identification and yield estimation. The pipette internal solution was filled with 1% Texas Red (Invitrogen) to label the recorded cells in a subset of experiments (see Fig. 7).

RGN identification. A single retinal quarter was placed in the patch-clamp recording chamber with the ganglion cell layer on the top. Using a recording glass pipette tip, membrane of the surface layer was penetrated and removed by moving the pipette back and forth to expose the retinal ganglion cell layer. The internal limiting membrane encasing the ganglion cell layer was carefully dissected using an empty patch pipette, and access was gained to a selected cell for patch-clamp recording (Murphy and Rieke, 2006; Margolis and Detwiler, 2007). Larger cells in the ganglion cell layer (~ 20 μ m diameter) were targeted in the flat-mount retina, and recordings were performed in the dark. We did not test cellular response to light versus dark, and we performed all our recording in normal ACSF without any synaptic blockers.

Whole-cell voltage-clamp recording

Voltage-clamp experiments were performed in Mes V neurons to evaluate changes in voltage-gated Na⁺ currents in the mSOD1 mice, compared with their WT littermates as controls. Recordings were performed using a pipette internal solution composed of the following (in mM): 130 CsF, 9 NaCl, 10 HEPES, 10 EGTA, 1 MgCl₂, 3 K₂-ATP, and 1 Na-GTP.

The external recording solution consisted of the following (in mM): 131 NaCl, 10 HEPES, 3 KCl, 10 glucose, 2 CaCl₂, 2 MgCl₂, 10 tetraethylammonium (TEA)-Cl, 10 CsCl₂, 1 4-AP, and 0.3 CdCl₂. Use of TEA, 4-AP, and CdCl₂ allowed isolation of voltage-gated Na⁺ currents by pharmacologically blocking the voltage-gated K⁺ and Ca²⁺ currents (Del Negro and Chandler, 1997; Enomoto et al., 2006). For acceptable recordings, each cell's uncompensated series resistance (R_s) was monitored throughout, and only recordings with $R_s < 10\%$ of the input resistance (R_{inp}) were included. Voltage-clamp protocols were used to activate the Nav1.6-type persistent and resurgent voltage-gated Na⁺ currents, both of which are critical for Mes V neuron excitability (Wu et al., 2005; Enomoto et al., 2007). To activate the resurgent Na⁺ current, the voltage-clamp protocol consisted of a brief 3 ms step to 30 mV from an initial holding potential of -90 mV, followed by 100 ms steps from 0 to -90 mV in steps of 10 mV, and then a return to the holding potential of -90 mV (Raman and Bean, 2001; Enomoto et al., 2006). The slower low-voltage-activated persistent Na⁺ current protocol included a slow voltage ramp from -90 to 30 mV for 1 s, followed by a step to -40 mV (Do and Bean, 2003; Wu et al., 2005). Following these protocols, these Na⁺ currents were abolished using 0.5 μ M TTX, and leak-subtracted Na⁺ current magnitude was quantified (see Fig. 3).

Real-time dynamic-clamp electrophysiology

Real-time closed-loop dynamic-clamp electrophysiology was adopted for *in silico* knock-in of Nav1.6-type Na⁺ currents in Mes V neurons. Briefly, the Linux-based Real-Time eXperimental Interface (RTXI version 1.3) was used to implement dynamic clamp, running on a modified Linux kernel extended with the Real-Time Applications Interface, which allows high-frequency, periodic, real-time calculations (Lin et al., 2010). The RTXI computer interfaced with the electrophysiological amplifier (Molecular Devices, Axopatch 200A in current-clamp mode) and the data acquisition PC via a PCIe-6251 board (National Instruments). Computation frequency was 20 kHz. Brainstem slices were perfused with oxygenated recording solution (~ 2 ml/min) at room temperature ($\sim 22^\circ\text{C}$ – 24°C) while secured in a glass-bottom recording chamber mounted on an inverted microscope with differential interface contrast optics (Carl Zeiss, Axiovert 10). Current-clamp (and dynamic-clamp) data were acquired and analyzed using custom-made software (G-patch, Analysis) with sampling frequency 10 kHz and cutoff filter frequency 2 kHz. The conductance-based Nav1.6-type Na⁺ current models were developed in our laboratory as detailed by Venugopal et al. (2019).

Experimental design and statistical analyses

Animals were divided into two main experimental groups: those carrying the human-SOD1^{G93A} transgene (mSOD1) and their age-matched non-carrier WT littermate controls. Statistical analyses were performed using Excel (Microsoft) and Systat 13 (Systat Software). Groups were compared using two-tailed unpaired Student's *t* test, two-way repeated-measures ANOVA to analyze group (WT vs mSOD1), and treatment effects (e.g., different levels of current application), and χ^2 proportionality test for comparison of distribution of cell types in the two animal groups (WT vs mSOD1). For two-way repeated-measures ANOVA, we assumed equal variance and normal distribution and first conducted a *post hoc* univariate *F* test for pairwise comparisons of treatment effect for each genotype. The statistical significance obtained using the *F* test, which is based on a regression model, was further verified against a two-tailed unpaired Student's *t* test, which indeed gave a more conservative estimate of the statistical significance we report here (for details, see results relating to Figs. 1*h*, 3*c–f*, 6*g*, and 7*e*). $p < 0.05$ was considered statistically significant.

Data acceptance criteria and analytical approaches

Patch-clamp recording was performed in whole-cell configuration following rupturing of a giga-ohm seal. Cells with uncompensated series resistance > 10 M Ω and resting potential more positive than -50 mV were discarded. Further data acceptance criteria included input resistance ≥ 100 M Ω (measured as the slope, voltage/current at the end of

100 ms current pulses near rest $V_{\text{rest}} \pm 10$ mV) and action potential height ≥ 80 mV (measured from spike threshold to peak). Spike characteristics, including height, half-width, after-hyperpolarization, and input threshold ($I_{\text{threshold}}$), were determined using response to a 10 ms current pulse. Average spike frequencies used for frequency-current relationships were determined for increasing steps of 1 s current pulses, typically up to $3 \times I_{\text{threshold}}$. In our Mes V discharge patterns, we distinguished rhythmic bursting activity from tonic spiking by constructing the interevent interval (IEI) histogram with bin width of 1 to 5 ms. If the IEI histogram had a clear single peak at ~ 20 ms, the cells were classified as tonic cells, whereas cells with additional peak(s) at larger IEI values ≥ 100 ms corresponding to the interburst intervals (IBIs) were grouped as bursting neurons. In the bursting Mes V neurons, we used 40 ms as the cutoff for IEI to further separate the interspike intervals (ISIs) from IBIs; all tonically spiking neurons had visibly regular IEIs < 50 ms with a sharp peak in the IEI histograms between 10 and 20 ms. These analyses were performed using Clampfit 10.5. Autocorrelation was used as a measure of rhythmicity of membrane voltage responses in Clampfit 10.5.

Nav1.6 fluorescent immunohistochemistry

Tissue preparation

Pups were deeply anesthetized using intraperitoneal injection of sodium pentobarbital (2.2 $\mu\text{g/g}$ BW). Using a 25-gauge needle connected to a pressure-controlled perfusion system (~ 200 mmHg), an adequate volume of freshly prepared, chilled 4% PFA (fixative) was transcardially perfused until clear drainage from the right atrium was noted. Subsequently, the mouse was decapitated, and the brain was extracted and incubated in 30% sucrose solution at 4°C for 48 h. The pontine brainstem block was cut and placed caudal side down within an RNase-free mold and embedded in optimal cutting temperature compound (Tissue TEK-OCT compound, Thermo Fisher Scientific), flash frozen using dry ice, and stored at -80°C .

Cryo-sectioning and fluorescent immunohistochemistry

Pontine brainstem was sectioned at 20 μm thickness beginning approximately at the level of the exiting VII nerve, and 6–8 sections per animal were mounted on glass slides (Thermo Fisher Scientific), ensuring inclusion of the rostro-caudal extent of the Mes V area. Fluorescent immunohistochemistry was immediately performed on slides by marking contours around each section using a MINI PAP PEN (Invitrogen). Every experiment was simultaneously run on WT and mSOD1 pair(s) to ensure similar tissue treatment for subsequent fluorescence quantification. Following three 10 min washes with $1 \times$ PBS, sections were incubated in a blocking buffer (10% normal donkey serum and 0.3% Tween-20 in $1 \times$ PBS) for 1 h at room temperature. Red fluorescence-conjugated Nav1.6 primary antibody derived from rabbit (Alomone Labs) was added (1:200 dilution) and incubated for 24 h at 4°C . Appropriate dilution was determined based on initial dilution series experiments (1:100, 1:200, and 1:500) with suitable negative controls. Following primary incubation, further wash steps (5 min, 3 times in $1 \times$ PBS) and green fluorescence-conjugated NeuN stain (Neurotrace) were added (1:200 dilution) for 30 min to enable identification of neuronal cell bodies. Sections were rinsed for 5–10 min with $1 \times$ PBS and coverslipped for imaging.

Imaging and quantification of Nav1.6 intensities

Imaging was performed using a fully automated epifluorescence microscope (Keyence, Bz 9000e). First, the Mes V nuclei were bilaterally identified with NeuN stain at low magnification ($2\times$), ventrolateral to the periaqueductal gray. The identified cells within the nucleus were then imaged at $60\times$ magnification (1/15 s exposure time for NeuN and 2 s exposure time for Nav1.6 with standard excitation settings), and z stacks were collected at 0.5 μm depth resolution with green (NeuN) and red (Nav1.6) filters. The z-projected images were then overlaid to generate a merged image to localize Nav1.6 label on Mes V neurons. Quantification of fluorescent intensities in Mes V neurons was performed using open-source ImageJ software (Schindelin et al., 2012). First, 12 ± 6 Mes V neurons with clearly visible NeuN-stained cell bodies and nuclei were

manually traced as ROIs in each image. These ROIs were then overlaid on the z-projected Nav1.6 image (see Fig. 4). The mean ROI intensity and area were quantified for each cell/ROI using the ImageJ menu-driven functions.

Computational model of simplified sensorimotor network

Our simplified sensorimotor network consisted of conductance-based models for sensory Mes V neuron and trigeminal MN (TMN). All model conductances followed the well-known Hodgkin-Huxley formalism (Hodgkin and Huxley, 1952). The model was implemented using XPPAUT (Ermentrout, 1997) script and will be made available upon request. The model consists of an excitatory network input from the Mes V to the TMN and the model assumptions are detailed below.

The Mes V neuron model consisted of our recently updated version consisting of both persistent and resurgent Na^+ currents that were reduced in the mSOD1 mouse in this study (Venugopal et al., 2019). Briefly, the model incorporates a minimal set of ionic conductances essential for producing rhythmic bursting and for maintaining cellular excitability in these neurons (Wu et al., 2005). These include a potassium leak current I_{leak} , a 4-AP-sensitive delayed-rectifier type potassium current I_K , and a Nav1.6-type sodium current I_{Na} with three separable components: fast, persistent, and resurgent currents. Each of these model currents is based on our previous experimental work (Chandler et al., 1984; Del Negro and Chandler, 1997; Wu et al., 2001, 2005; Enomoto et al., 2006; Venugopal et al., 2019).

The TMN model is based on well-established two-compartment MN models, which include dendritic Ca^{2+} currents (e.g., Booth et al., 1997; Venugopal et al., 2011b). This is based on known physiology of MNs, which richly express noninactivating dendritic L-type Ca^{2+} currents, also known as the persistent Ca^{2+} current (CaP) in both the brainstem and spinal cord (Lee and Heckman, 1998; Carlin et al., 2000; Theiss et al., 2007). We chose such a model to reproduce dendritic plateau potentials, membrane bistability (Booth et al., 1997), and also since the CaP plays a crucial role in synaptic amplification (Hultborn et al., 2003) and motor control (Lee et al., 2003; Heckman et al., 2005). The somatic compartment consisted of the action potential generating fast sodium, delayed-rectifier type potassium, high-voltage-activated N-type Ca^{2+} and N-type Ca^{2+} -activated potassium conductances (Hsiao et al., 1998, 2005). The plateau-generating L-type persistent Ca^{2+} or CaP, CaP-activated potassium and a persistent sodium conductance were confined to the dendritic compartment. Both compartments also consisted of a potassium leak conductance.

A lumped synaptic variable modeled the excitatory synaptic input from the Mes V to TMN model. This assumption was based on known anatomy and physiology of Mes V-TMN connections. For example, nearly 80% of Ia muscle spindle afferents from the Mes V nucleus synapse on to proximal and distal dendrites of TMNs (Yoshida et al., 2017). To incorporate such dendritic excitation, we modeled the TMN with two electrically coupled morphological compartments (soma and lumped dendrite) (Booth et al., 1997; Venugopal et al., 2011b, 2012) and confined synaptic input to the dendritic compartment. Second, we tuned the strength of the model excitatory conductance to generate sensory-coupled spikes in the TMNs since Mes V neurons provide strong monosynaptic glutamatergic excitation to TMNs (Chandler, 1989; Del Negro and Chandler, 1998; Turman et al., 1999, 2000). For simplicity, we assumed no other form of synaptic excitation or inhibition. This allowed us to exclusively examine the consequence of irregular sensory patterns as observed in our data, on MN discharge. The lumped synaptic variable represented a classic excitatory synapse modeled as follows (Venugopal et al., 2011a):

$$s' = \alpha_s(1 - s)s_\infty(V) - \beta_s s$$

where s is the synaptic variable, $\alpha_s = 2$ is the rate constant for fast rise of s , and $\beta_s = 0.05$ is the decay rate constant for relatively slower decay of s (Heckman and Binder, 1991; Trueblood et al., 1996; Powers et al., 2012). The steady-state voltage-dependent function $s_\infty(V)$ is given by the following:

Table 1. Membrane properties of Mes V neurons

	Burst		Tonic	
	WT (n = 18)	mSOD1 (n = 19)	WT (n = 14)	mSOD1 (n = 7)
RMP, mV	−56.9 ± 1.5	−54.8 ± 1.0	−53.6 ± 1.6	−52.8 ± 1.4
R _{in} , MΩ	211.5 ± 25.9	253.1 ± 19.3	264.2 ± 16.9	300.1 ± 18.2
C _m , pF	49.2 ± 3.1	50.3 ± 2.7	42.0 ± 2.1	42.2 ± 2.2

Table 2. Action potential properties of Mes V neurons^a

	Burst		Tonic	
	WT (n = 18)	mSOD1 (n = 19)	WT (n = 14)	mSOD1 (n = 7)
AP height, mV	89.3 ± 3.1	90.2 ± 4.3	92.5 ± 4.1	93.5 ± 3.8
Half-width, ms	1.3 ± 0.13	1.3 ± 0.10	1.2 ± 0.08	1.4 ± 0.09 (<i>p</i> = 0.025)
AP threshold, mV	−50.1 ± 2.1	−44.6 ± 1.1 (<i>p</i> = 0.008)	−44.7 ± 2.1	−41.4 ± 1.1
AP current, pA	40.7 ± 5.4	57.1 ± 7.2 (<i>p</i> = 0.04)	41.4 ± 4.0	50.0 ± 7.7
AHP peak, mV	−11.0 ± 0.9	−8.1 ± 0.8 (<i>p</i> = 0.009)	−9.0 ± 0.7	−8.5 ± 0.7

^a*p* values are based on Student's *t* test.

$$s_{\infty}(V) = \frac{1}{\left(1 + e^{\frac{-(V+15)}{0.02}}\right)}$$

where, *V* is the voltage of the presynaptic neurons (here, Mes V neuron). The EPSC was included in the dendritic compartment of the TMN model and is given by the following conductance-based equation:

$$I_{syn} = g_{syn}s(V_d - V_e)$$

where *V_d* is the TMN dendritic voltage and *V_e* = −20 mV is the reversal potential for an excitatory synaptic current. The value of maximal synaptic conductance, *g_{syn}*, was set such that presynaptic Mes V bursts generated postsynaptic TMN firing frequencies in the range observed in SOD1^{G93A} TMNs (Venugopal et al., 2015).

Results

Impaired excitability of SOD1^{G93A} proprioceptive Mes V neurons

First, we examined whether the electrical excitability of the jaw muscle proprioceptive neurons in the trigeminal Mes V nucleus shows abnormalities in the mSOD1 mice. Using suitable morphological and anatomical selection criteria, we conducted *in vitro* whole-cell patch-clamp recording and evaluated the basic membrane properties, action potential characteristics, and responses to increasing step depolarizations. The comparative membrane properties in mSOD1 and WT are summarized in Tables 1 and 2. In Figure 1*a*, we illustrate the location of the anterogradely labeled jaw closer muscle spindle afferents in the brainstem Mes V nucleus (green), obtained from injection of dye into the jaw closer muscles. Concomitantly, the jaw closer motor pools are also retrogradely labeled specifically in the dorsolateral trigeminal motor nucleus (Mot V) as shown in Figure 1*a*, ventrolateral to Mes V. In Figure 1*b*, we present an example of an anterogradely labeled Mes V sensory neuron, which is dye-filled during whole-cell patch-clamp recording. Figure 1*d* shows the age distribution of mice from which our dataset was obtained. In Figure 1*c*, we show representative examples of the two most commonly observed patterns of action potential discharges in our dataset from both WT and mSOD1 mice (Brocard et al., 2006; Enomoto et al., 2006). These included the following: (1) rhythmic burst discharge consisting of repetitive sequences of high-frequency spikes (~80 Hz), followed by periods of quiescence on the order of 100–1000 ms (Wu et al., 2001; Brocard et al., 2006); and (2) a tonic spiking pattern with short ISIs, on the order of 10 ms. Our empirical criteria for classifying cells as bursters were

based on the distributions of spike intervals (see Experimental design and statistical analyses). In the mSOD1 bursting Mes V neurons, we noted marked increase in the current and voltage thresholds for spike generation (Fig. 1*e,f*) compared with the WT. The mean ± SD for *I_{threshold}* was 40.7 ± 5.4 pA and 57.1 ± 7.2 pA, for *V_{threshold}* was −50.1 ± 2.1 mV and −44.6 ± 1.1 mV for WT and mSOD1, respectively. Corresponding *p* values from a two-sample Student's *t* test were 0.04 for *I_{threshold}* and 0.008 for *V_{threshold}* comparisons. The bursting neurons in the mSOD1 mice also showed diminished spike frequencies in response to increasing current injections (Fig. 1*h*). We quantified the average spike frequencies for the last 500 ms of 1 s step depolarization of increasing magnitude. A factorized two-way repeated-measures ANOVA was used to examine the effects of genotype (Factor 1), current injections (Factor 2), and any interaction effects on measured spike frequencies (dependent variable). We noted a significant between-subjects genotype effect (*df* = 1, *F* ratio = 28.996, *p* = 1.8E-05) and within subjects injected current effect (*df* = 5, *F* ratio = 38.484, *p* = 1.8E-016); there was no genotype × current interaction effect (*df* = 5, *F* ratio = 1.678, *p* = 0.145), suggesting that increasing current intensities similarly enhanced spike frequencies in both WT and mSOD1. A *post hoc* unpaired, two-tailed Student's *t* test confirmed significant reductions in spike frequencies for injected current magnitude ≥ 150 pA (Student's *t* test, 150 pA: *p* = 0.0091, 200 pA: *p* = 0.01, 250 pA: *p* = 0.0003, 300 pA: *p* = 0.01). Furthermore, a greater proportion of mSOD1 Mes V neurons presented burst-like activity (Fig. 1*i*). A χ^2 proportionality test yielded *p* = 0.0074. Conversely, tonically spiking neurons were less perturbed and did not present any statistically significant *I_{threshold}* or *V_{threshold}* changes (Fig. 1*e,g*); however, at higher current injections ≥ 250 pA, these neurons also showed reduced firing frequencies similar to bursting neurons (Fig. 1*h*) (Student's *t* test, 250 pA: *p* = 0.008, 300 pA: *p* = 0.0174); the ANOVA statistics for between-subjects genotype differences was as follows: *df* = 1, *F* ratio = 11.8688, *p* = 0.0073, and for increasing current intensities was as follows: *df* = 5, *F* ratio = 27.202, *p* = 1.45E-012. We noted that these neurons also showed an increase in spike width (summary, Tables 1, 2). Moreover, the proportion of tonically firing cells was reduced in the mSOD1 Mes V dataset (Fig. 1*i*). Together, Mes V neurons in the SOD1^{G93A} mouse are hypoexcitable with a large propensity toward burst formation in action potential trains.

Arrhythmic spike and burst patterns of SOD1^{G93A} Mes V neurons

This overall reduction in action potential generation capability and increased burst discharge in the mSOD1 Mes V neurons was accompanied by further abnormalities in bursting properties. In Figure 2*a*, we illustrate a rhythmic burst pattern in a WT Mes V neuron (left) compared with irregular burst patterns in a mSOD1 Mes V neuron (right). The corresponding IEI time series highlights such irregularities in Figure 2*b*. In the rhythmic WT burst pattern, the IBIs are well demarcated from the ISIs. However, this distinction is less obvious in the mutant pattern; and as noted, the IBI regularity from one burst to another is also uncertain. The ISIs also presented significant irregularities observed for the shorter spike intervals. We used the autocorrelation function of the membrane voltage and compared its second peak between WT and mSOD1 cells using a time lag, τ = 100 ms to examine regularity of spikes (Rieke et al., 1999). This time window allowed examination of interspike irregularities within a typical burst. Figure 2*c* illustrates representative examples of the autocorrelation function for WT (left) and mSOD1 (middle) Mes V neurons.

Figure 1. Impaired excitability in intrinsically bursting mSOD1 Mes V neurons. **a**, Schematic represents the Ia proprioceptive sensory and motor nuclei of the trigeminal jaw control circuit identified using anterograde/retrograde tracer injection in jaw closer muscles in the SOD1^{G93A} mouse (or mSOD1). Left half of a coronal pontine section is shown. Mes V, Trigeminal mesencephalic sensory nucleus; Mot V, trigeminal motor nucleus. White dotted enclosures represent both nuclei. **b**, Left, Anterogradely labeled Mes V neurons (green). Middle, Dye-filled (red) Mes V neuron during patch-clamp recording. Right, A merged image highlighting the double-labeled neuron. **c**, Representative examples of discharge patterns in two classes of Mes V neurons. Left, A rhythmic bursting neuron (maroon). Right, A tonic spiking neuron (grey). These discharge patterns are induced by a 200 pA depolarizing step current for 20 s duration. **d**, Bar charts represent the age distribution of the mice, including 24 WT mice ($n = 32$) and 21 mSOD1 mice ($n = 26$), where n is the number of cells for all the data presented. **e**, A single near-threshold spike in a WT (blue) and a mSOD1 (red) Mes V neuron generated in response to a brief (5 ms) step depolarization. The corresponding injected current threshold ($I_{threshold}$) (bottom) and membrane voltage threshold ($V_{threshold}$) for spike generation (top, horizontal lines and arrows) are highlighted (magenta). **f**, **g**, Dot plots represent $I_{threshold}$ (left) and $V_{threshold}$ (right) in bursting (**f**) and tonic (**g**) Mes V neurons. Blue and red horizontal lines indicate average group values. **f**, For bursting neurons, * $p < 0.05$ (above horizontal line), for $I_{threshold}$ and $V_{threshold}$. **g**, For tonic spiking neurons, NS (above black horizontal line) indicates no statistically significant group differences between WT and mSOD1 cells. **h**, Spike frequency-injected current responses in bursting (left) and tonic spiking (right) Mes V neurons: blue represents WT; red represents mSOD1. * $p < 0.05$ (above horizontal line), between WT and mSOD1 cells for injected current ≥ 150 pA. **i**, Pie charts represent proportional distribution of bursting and tonic spiking neurons in WT and mSOD1 mice. * $p < 0.05$.

A two-sample unpaired Student's t test with assumed equal variance showed that the second peak of autocorrelation between the two groups was significantly different. A $p = 0.0029$ was noted with mean WT peak $\pm$ SD as 0.44 ± 0.04 , and the mean mSOD1 peak $\pm$ SD as 0.30 ± 0.03 . We also note that the subsequent peaks are almost completely suppressed, further highlighting irregularity in spike patterns in the mutant. Additionally, the burst duration and IBI lengths were reduced while the ISIs were increased (Fig. 2d): for WT and mSOD1, respectively, the mean burst duration $\pm$ SD were 0.41 ± 0.11 s and 0.06 ± 0.01 s, mean IBI $\pm$ SD were 0.89 ± 0.14 s and 0.29 ± 0.03 s, and mean ISI $\pm$ SD were 17.2 ± 0.34 ms and 26.4 ± 2.37 ms. Together, the proprioceptive Mes V neurons present early impairment in action potential generation and in the ability to sustain regular burst discharge in the SOD1^{G93A} mouse.

Disrupted firing patterns of SOD1^{G93A} proprioceptive Mes V neurons are linked to reduced Nav1.6 Na⁺ currents and channel protein

To identify the ionic basis for the observed impairments in action potential bursts, we tested whether the voltage-gated Na⁺ currents, which are essential for spike generation and burst control in these neurons, are compromised (Wu et al., 2005; Enomoto et al., 2006; Yang et al., 2009). Our previous report using the Nav1.6 subunit KO mouse demonstrated significant reductions in TTX-sensitive Na⁺ currents, which paralleled impairment in spike discharge (Enomoto et al., 2007). These Na⁺ currents, including the persistent and resurgent components, are important for generation and maintenance of burst discharge in Mes V neurons (Wu et al., 2005; Enomoto et al., 2006; Venugopal et al., 2019). Therefore, we examined whether the observed impairment in

both excitability and burst discharge in a large majority of the mSOD1 Mes V neurons could be explained by reductions in the functional expression of these Nav1.6 Na⁺ channel currents. To test this, we used the *in vitro* voltage-clamp approach combined with suitable pharmacology and measured the Nav1.6 channel mediated persistent and resurgent Na⁺ currents in WT and mSOD1 Mes V neurons (see Cryo-sectioning and fluorescent immunohistochemistry). As shown in Figure 3*a–d*, the mSOD1 Mes V neurons showed significantly reduced current density (pA/pF) of voltage-gated persistent and resurgent Na⁺ current components, compared with WT. As shown in Figure 3*c*, the peak persistent Na⁺ currents were downregulated at subthreshold activation voltages of -60 and -50 mV (mean $\pm$ SD current densities at -60 mV were -1.01 ± 0.22 pA/pF and -0.43 ± 0.10 pA/pF, and at -50 mV were 1.15 ± 0.21 pA/pF and -0.66 ± 0.11 pA/pF for WT and mSOD1 cells, respectively). Corresponding *p* values based on a Student's *t* test were 0.03 and 0.044 at -60 and -50 mV, respectively. Similarly, as shown in Figure 3*d*, the peak resurgent Na⁺ currents were also downregulated at -40 and -50 mV at which these currents are typically maximally activated (Enomoto et al., 2006, 2007) (mean $\pm$ SD current densities at -50 mV were -5.13 ± 0.282 pA/pF and -3.23 ± 0.51 pA/pF, and at -40 mV these values are -5.69 ± 0.84 pA/pF, and -3.93 ± 0.47 pA/pF for WT and mSOD1 cells, respectively). Corresponding *p* values for Student's *t* test were 0.0387 and 0.0424 at -50 and -40 mV, respectively. Such reductions in current density, however, were not accompanied by altered voltage-gating properties in the measured Na⁺ currents between WT and mSOD1 Mes V neurons, as noted by the normalized conductance Boltzmann curves (Fig. 3*e,f*).

Next, using fluorescent immunohistochemistry, we tested whether the observed decrease in the Nav1.6 Na⁺ currents in the mSOD1 mouse could result from a downregulation of Nav1.6 Na⁺ channels on Mes V neuronal membrane (Enomoto et al., 2007). We quantified the Nav1.6 protein expression in Mes V neuron cell bodies colocalized with NeuN stain. The NeuN-stained Mes V neurons were readily detected based on their oval cell bodies and rostro-caudal distribution in the dorsal pontine sections, ventrolateral to the periaqueductal gray (Fig. 4*a*, inset). As noted in the figure, the Nav1.6 protein expression was discernable in both WT and mSOD1 Mes V neurons. The three NeuN images (green) shown as level A–C illustrate the representative caudal-to-rostral sections used in our analysis. For immunofluorescent quantification, we manually outlined the NeuN-stained Mes V cell bodies (ROIs) in which the nuclei were clearly visible under high magnification (60 $\times$) (see Materials and Methods). These ROIs were then overlaid on the Nav1.6 protein-stained

Figure 2. Arrhythmic burst discharge in mSOD1 Mes V neurons. *a*, Representative examples showing rhythmic bursting in a WT (blue) Mes V neuron (left), and arrhythmic bursting in a mSOD1 (red) Mes V neuron (right). *b*, Time series plots of IELs corresponding to the examples shown in *a* highlight IEL irregularities (IELs are shown on log scale). *c*, Autocorrelation function of the membrane voltage in a WT (left) and mSOD1 Mes V neuron (middle); second peak is highlighted. Right, Dot plots represent the second peaks in WT (blue) and mSOD1 (red) bursting neurons. *d*, Left, Inset, The timing properties of bursts. Dot plots represent significantly reduced burst duration (BD), IBIs, and increased ISIs in the mSOD1 (red) Mes V neurons compared with WT (blue). *c, d*, $n = 14$ (WT) and $n = 16$ (mSOD1), where n is the number of bursting cells. * $p < 0.05$ (above the black horizontal line).

image (Fig. 4*b*) and the ROI areas, and mean intensities were quantified using the open-source ImageJ software (Schneider et al., 2012). Figure 4*c* shows a scatterplot of ROI areas and mean intensities for WT (black circles) and mSOD1 (red circles) Mes V neurons to illustrate that there was no sampling bias in our cell samples. There was an overall positive correlation between ROI/cell area and mean intensities in both WT and mSOD1 Mes V neurons (trend lines have positive slopes in Fig. 4*c*). However, such a correlation was weaker in the mutant. Statistical comparison of ROI areas and mean intensities between WT and mSOD1 revealed significantly diminished mean intensities in the mutant (mean $\pm$ SD: 36.02 ± 13.18 a.u. for WT and 27.01 ± 12.61 a.u. for mSOD1; $p < 0.001$ using two-sample Student's *t* test) (Fig. 4*d*; box plots show first, second, and the third quartiles, and error bars indicate $1.5\times$ deviations from the median). However, the WT and mSOD1 ROI areas were not statistically different (data not shown), whereas the mean intensities per unit area (a.u./ μm^2) were also diminished in the mutant ($p < 0.001$). These results indicate that the reduced functional expression of Nav1.6 Na⁺ currents in the mSOD1 Mes V neurons are associated with a downregulation of Nav1.6 protein expression.

Figure 3. Reduced Nav 1.6-type Na⁺ currents in mSOD1 Mes V neurons. **a, b**, Representative current traces from whole-cell voltage-clamp experiments showing a reduction in TTX-sensitive persistent Na⁺ (**a**) and resurgent Na⁺ (**b**) currents in mSOD1 Mes V neuron (red), compared with WT (blue). Top, Voltage-clamp protocol. Bottom, Current responses. Voltage-clamp protocol in **a** consists of a ramp voltage from -90 to 30 mV at 100 mV/s, and in **b**, consists of a brief, 5 ms step from -90 to 30 mV, to evoke a transient Na⁺ current, followed by repolarization to voltages between -90 and 0 mV to elicit resurgent Na⁺ current due to open-channel unblock. **c, d**, Current-voltage relationships of the peak persistent (**c**) and resurgent (**d**) Na⁺ current densities in mSOD1 Mes V neurons (red). **c**, Asterisk (above the black horizontal line) indicates statistically significant reduction in peak current amplitudes at subthreshold activation voltages between -60 and -50 mV for persistent Na⁺ peaks in mSOD1 (red) compared with WT (blue). **d**, Asterisk (above the black horizontal line) indicates statistically significant reduction in peak current amplitudes at repolarization voltages of -50 and -40 mV for resurgent Na⁺ peaks in mSOD1 (red) compared with WT (blue). **e, f**, Boltzmann curves fit to the normalized peak conductances as a function of activation voltages for persistent (**e**) and resurgent (**f**) Na⁺ currents in WT (blue) and mSOD1 (red) Mes V neurons. **g**, Bar charts represent the age distribution of the mice, including 8 WT mice ($n = 14$) and 6 mSOD1 mice ($n = 11$), where n is the number of cells for all the data presented in **c–f**.

Rescue of Nav1.6 Na⁺ currents restored normal burst patterns and rhythmicity in SOD1^{G93A} proprioceptive Mes V neurons

To test whether rescuing the Nav1.6 Na⁺ current impairment can restore the normal burst discharge in the mSOD1 Mes V neurons, we used an *in vitro* dynamic-clamp approach. This included real-time injection of conductance-based Nav1.6 Na⁺ currents into a mSOD1 Mes V neuron during whole-cell current-

clamp recording of burst discharge (Fig. 5a). Figure 5b illustrates that addition of such realistic computer-generated Nav1.6 Na⁺ persistent and resurgent currents into an irregularly bursting mSOD1 Mes V neuron converted an irregular burst pattern to a more rhythmic pattern. The red traces in Figure 5b show the membrane voltage of the mSOD1 Mes V neuron during default burst discharge, and the black trace in between shows the modified burst discharge upon addition of the Nav1.6-type Na⁺ currents. To evaluate whether the rescue of Nav1.6-type Na⁺ currents in the mSOD1 neurons also restored the normal burst patterns similar to WT, we quantified the time intervals between spikes (IEIs) and demonstrate that the irregularities in mSOD1 IEIs were abolished (Fig. 5c) (Venugopal et al., 2019). Second, we measured the second peak of the autocorrelation function of the membrane potential and show that the rescue of Na⁺ currents restored the rhythmicity to WT values in a dose-dependent manner (Fig. 5d). Last, Na⁺ current addition also reinstated the time intervals and duration to WT values (Fig. 5e).

Together, the muscle spindle afferent neurons in the mSOD1 Mes V nucleus present early impairment in excitability, including hypoexcitability and arrhythmicity, which are associated with reductions in the voltage-gated Na⁺ currents and Nav1.6 Na⁺ channels.

Mechanoreceptive and nociceptive trigeminal sensory neurons lack early excitability changes in the SOD1^{G93A} mouse

Next, to assess whether the altered excitability is restricted to the proprioceptive muscle spindle afferents in the mSOD1 Mes V nucleus, we examined the nonproprioceptive primary sensory neurons in the TG. The sensory architecture of the trigeminal system segregates the A α -type proprioceptive neurons in the Mes V and the A β mechanoreceptors, A δ nociceptors, and C-fiber type neurons in the peripheral TG (Lazarov, 2002). Therefore, we examined whether the different types of sensory neurons in the TG were affected differently by SOD1 mutation. To

conduct *in vitro* patch-clamp electrophysiology, we performed acute dissociation of the TG in P8–P14 WT and mSOD1 mice, age-matched with mice used to examine Mes V neurons (Fig. 6a,b; see also Trigeminal ganglia (TG) extraction and acute dissociation of trigeminal ganglion neurons (TGNs) for electrophysiology). In both WT and mSOD1 TG neurons, we were able to distinguish the A β from the A δ nociceptive neurons based on action potential duration and presence of a hump, as shown in

Figure 4. Immunofluorescent quantification of Nav1.6 protein expression in WT and mSOD1 Mes V cells. **a**, Representative images showing three rostro-caudal levels of coronal brainstem sections, stained with NeuN (left column of images). Scale bar, 500 μm . In these images, the Mes V nucleus is highlighted with dashed contours. At Level B, the subjacent trigeminal motor nucleus (mot V) is highlighted. For each level, comparative images consisting of Mes V neurons from WT and mSOD1 mice are shown at 60 $\times$ magnification. Green represents NeuN. Red represents Nav1.6 protein. Merged images are shown enlarged with white arrows highlighting representative Mes V neurons. Left, Bottom, Boxed inset represents the pontine levels at which sections were collected. **b**, Representative image showing ROIs drawn at ~ 12 Mes V neurons for immunofluorescence quantification. White arrows indicate three representative ROIs. **c**, Scatterplot showing WT (black circles) and mSOD1 (red circles) values of ROI area and mean intensity quantification. Trendlines show a positive linear regression. **d**, Box plots represent mean ROI intensity per cell for WT (black) and mSOD1 (red), respectively. *n* values indicate number of cells obtained from 4 WT and 4 mSOD1 mice across 9 and 8 sections, respectively. A two-tailed Student's *t* test was used for statistical comparison.

Figure 6c (Xu et al., 2010; Kim et al., 2011). A small proportion of C-fiber type neurons were also present in our dataset, which were distinguished from the A-type neurons based on a lack of membrane sag during the hyperpolarizing current injections (Fig. 6c: a white block arrow in the left two panels highlight membrane sag). However, these cells exhibited single spikes and were comparatively fewer in both WT and mSOD1 datasets to yield meaningful statistical assessment and, hence, were excluded from further analysis. Interestingly, there were no proportional shifts in the sample dataset among these different types of neurons between WT and mSOD1 mice as shown in Figure 6d. We compared the basic membrane properties and action potential characteristics of WT and mSOD1 TG neurons as shown in Tables 3 and 4, the spike threshold current and voltage as shown in Figure 6e. None of these properties was different between WT and mSOD1. The average spike frequency responses to increasing steps of depolarization also did not present any significant changes w.r.t. geno-

type (Fig. 6f,g; frequency calculation and statistical tests and assumptions were similar to Mes V neurons and *p* values were > 0.05 and *F* ratio ~ 0). Last, the autocorrelation function of the membrane voltage (Fig. 6h) did not show arrhythmicity in the mSOD1 TG neurons compared with WT. The second peak of the autocorrelation function was similar, and subsequent peaks were also similarly discernable with the autocorrelation time lag of 100 ms. These results suggest that nonproprioceptive trigeminal sensory neurons are resistant to early excitability changes in the SOD1^{G93A} mouse model for ALS.

Early excitability changes are absent in the SOD1^{G93A} RGNs

Last, we investigated whether the peripheral visual sensory RGNs show altered excitability in the mSOD1 mice. For these experiments, we used a whole retinal preparation and conducted whole-cell patch-clamp electrophysiology *in situ* in the RGNs in P8–P14 mice (Fig. 7a,b; see also Retina extraction and prepara-

Figure 5. Rescue of Nav1.6-type sodium currents in mSOD1 Mes V neurons using dynamic clamp. **a**, Schematic shows the dynamic-clamp setup used to introduce conductance-based models of Nav1.6-type Na^+ currents into mSOD1 Mes V neurons in real time during whole-cell patch-clamp recording. I_{dyn} is the computer-generated model Na^+ current in combination with a step depolarization to drive the patched Mes V neuron. V_m is the measured membrane voltage. **b**, Representative traces showing the membrane voltage in a bursting mSOD1 Mes V neuron with control/default behavior (left, red), followed by addition of Nav1.6-type currents (middle, black), which restores WT-like rhythmic bursting; subsequent removal of added currents returns default mSOD1 behavior (right red) in this neuron. Bottom, Green trace represents I_{dyn} . **c**, Time series plots of IEIs (log scale) for the three different conditions in **b**. Each dot represents an interval between two consecutive spikes. **d**, Autocorrelation function of the membrane voltage for default mSOD1 (left) and with addition of Nav1.6 currents (middle). The height of the second autocorrelation peak highlights rhythmicity (arrowhead). Right, Measured second peak values for 4 mSOD1 cells. Dashed horizontal line indicates average WT values. Gray shaded region represents $\pm$ SD. **e**, Treatment effects on burst characteristics, including burst duration (BD), IBIs, and ISIs within bursts (ISI) shown for various mSOD1 cells tested under the different conditions as in **b**. Dashed lines indicate average WT values. Gray regions represent $\pm$ SD.

tion for electrophysiology). The rationale for examining RGN excitability was threefold: (1) to test whether functional abnormalities were limited to muscle spindle afferent proprioceptive neurons while absent in the visual sensory neurons, similar to the mechanoreceptive and nociceptive TG neurons; (2) to ascertain whether excitability changes were not confounded by the type of preparation (*in vitro* dissociated TGNs vs *in situ* RGNs); and (3) to clarify whether postnatal excitability changes in other sensory neurons are equally immune in parallel with maturing peripheral sensory systems in mice (Cabanes et al., 2002; Chen et al., 2009). Similar to the Mes V and TG neurons, we com-

pared the membrane properties (Tables 5, 6), action potential thresholds (Fig. 7c), spike frequency-injected current responses (Fig. 7d,e), rhythmicity/regularity of membrane voltage (Fig. 7f), and proportional distribution of apparent electrophysiological subtypes of RGNs in the WT and mSOD1 mice (Fig. 7g,h). We found that none of the above properties tested was different between the two groups (all the statistical tests were similar to Mes V neurons). These results confirm that the early excitability changes are limited to muscle spindle afferent proprioceptive neurons in the cranial sensory system of the SOD1^{G93A} mouse model for ALS.

Figure 6. Excitability of primary sensory neurons in the TG of mSOD1 mice. *a*, Schematic showing the trigeminal sensory nuclei along the midbrain–brainstem regions. Primary 1a afferent neurons in the TG (red dashed rectangle) were acutely dissociated, and whole-cell current-clamp recordings were performed. *b*, Top left, Example showing NeuN stain (green) used to identify the dissociated TG neurons. Top right, Example of a dissociated TG neuron filled with Texas red 568 dye during whole-cell recording. Bottom left, Merged image. Bottom right, Bright field (BF) image. Scale bars, 50 μ m. *c*, TG neurons were classified into three types: A β , A δ , and C (for criteria, see Results). Open arrows indicate membrane sag during a 1 s hyperpolarizing step pulse in A β and A δ types. Black arrow indicates a hump in A δ action potential. *d*, Percentage of the subtypes of TG neurons in WT and mSOD1 mice are not significantly different. *e*, Inset, The current ($I_{\text{threshold}}$) and voltage ($V_{\text{threshold}}$) thresholds for action potentials in WT (blue) and mSOD1 (red) TG neurons. Dot plots represent comparison of these properties between WT (blue) and mSOD1 (red) within the A β and A δ groups with no statistical significance. *f*, Representative membrane voltage responses to increasing levels of 1 s current injection (bottom traces) in repetitively firing WT and mSOD1 TG neurons (both A δ type). *g*, Spike frequency–injected current responses in all the A δ TG neurons that showed repetitive firing. *h*, Autocorrelation function of membrane potential in WT (left) and mSOD1 (middle) TG neurons; second peaks (arrows) are highlighted. Right, Dot plots represent the second peaks for all the cells with no significant difference between WT (blue) and mSOD1 (red) TG neurons. *i*, Bar charts represent the age distribution of the mice, including 8 WT mice ($n = 36$) and 7 mSOD1 mice ($n = 27$), where n is the number of cells for all the data presented in *e–h*.

Table 3. Membrane properties of TG neurons

	A β		A δ		C	
	WT ($n = 18$)	mSOD1 ($n = 13$)	WT ($n = 15$)	mSOD1 ($n = 11$)	WT ($n = 3$)	mSOD1 ($n = 3$)
RMP, mV	-53.2 ± 0.9	-54.2 ± 0.9	-50.8 ± 0.3	-52.6 ± 1.1	-62.5 ± 2.5	-63.0 ± 3.0
R_{in} , M Ω	293.7 ± 86.4	164.6 ± 22.9	413.9 ± 50.1	373.2 ± 56.8	224.1 ± 50.6	308.8 ± 123.4
C_{m} , pF	32.8 ± 4.6	35.1 ± 4.2	20.3 ± 3.7	18.4 ± 4.8	21.8 ± 4.2	27.4 ± 8.8

Table 4. Action potential properties of TG neurons

	Aβ		Aδ		C	
	WT (n = 18)	mSOD1 (n = 13)	WT (n = 15)	mSOD1 (n = 11)	WT (n = 3)	mSOD1 (n = 3)
AP height, mV	107.8 ± 2.9	108.1 ± 3.8	113.8 ± 4.3	112.1 ± 2.6	104.5 ± 7.9	105.0 ± 8.0
Half-width, ms	1.3 ± 0.16	1.5 ± 0.25	2.6 ± 0.19	2.7 ± 0.33	2.3 ± 0.04	3.6 ± 0.9
AP threshold, mV	−27.6 ± 1.3	−26.6 ± 1.4	−19.4 ± 1.4	−20.3 ± 2.1	−23.6 ± 7.8	−20.4 ± 3.0
AP current, pA	546.7 ± 118.7	524.6 ± 103.2	212.7 ± 33.9	244.5 ± 75.5	210.0 ± 37.9	200.0 ± 61.1
AHP peak, mV	−5.9 ± 0.8	−5.1 ± 0.8	−7.1 ± 0.7	−6.3 ± 0.6	−4.3 ± 1.2	−3.1 ± 2.1

Figure 7. Excitability of primary sensory neurons in the retina of mSOD1 mice. **a**, Schematic showing retina extraction for patch-clamp recording. **b**, Bar charts represent the age distribution of the mice, including 7 WT mice ($n = 13$) and 5 mSOD1 mice ($n = 10$), where n is the number of cells for all the data presented in **c–h**. **c**, Inset, Current ($I_{\text{threshold}}$) and voltage ($V_{\text{threshold}}$) thresholds for action potentials in WT (blue) and mSOD1 (red) RG neurons. Dot plots represent comparison of these properties between WT (blue) and mSOD1 (red) RG neurons tested. **d**, Representative membrane voltage responses to increasing levels of 1 s current injection (bottom) in repetitively firing WT and mSOD1 RG neurons. **e**, Spike frequency–injected current responses in all the RG neurons that showed repetitive firing. **f**, Autocorrelation function of membrane potential in WT (left) and mSOD1 (middle) RG neurons. Second peaks (arrows) are highlighted. Right, Dot plots represent the second peaks for all the cells with no significant difference between WT (blue) and mSOD1 (red) TG neurons. **g**, RG neurons were classified into three types: single spikers (SS), rate adapting (RA), and tonic spiking (for details, see Results). **h**, Percentage of the subtypes of RG neurons in WT and mSOD1 mice are not significantly different.

Computer modeling predicts motor dysfunction due to perturbed proprioceptive sensory gating

Difficulty in chewing and swallowing is a common clinical feature observed in ALS patients with both spinal and bulbar onset (Riera-Punet et al., 2018a,b). In the SOD1^{G93A} mouse model, mastication is severely impaired early during disease develop-

ment (Lever et al., 2009). During normal mastication, the Mes V neurons relay sensory feedback from the jaw muscle spindles to adjust the activity and force generated in the jaw-closer muscles. Nearly 80% of the glutamatergic projections from the Mes V neurons synapse on the jaw-closer trigeminal motor pools and provide muscle stretch reflex inputs to the TMNs (Yoshida et al.,

Table 5. Membrane properties of retinal ganglion cells

	WT (<i>n</i> = 13)	SOD1 (<i>n</i> = 10)
RMP, mV	−55.4 ± 2.0	−54.1 ± 2.4
R_{in} , MΩ	462.3 ± 19.3	436.2 ± 24.4
C_m , pF	29.0 ± 3.3	29.3 ± 4.4

Table 6. Action potential properties of retinal ganglion cells

	WT (<i>n</i> = 13)	mSOD1 (<i>n</i> = 10)
AP height, mV	67.6 ± 3.9	66.3 ± 5.3
Half-width, ms	3.8 ± 0.5	4.0 ± 0.7
AP threshold, mV	−50.6 ± 1.5	−51.6 ± 2.2
AP current, pA	22.7 ± 4.7	22.2 ± 4.6
AHP peak, mV	−6.1 ± 0.4	−6.4 ± 0.8

2017). It is likely that the observed abnormalities in the mSOD1 Mes V excitability alter the jaw stretch reflex control. To examine how irregular discharge in the mSOD1 Mes V neurons might modify motor discharge, we used a computational model of a simplified sensorimotor network. Our model consisted of a realistic sensory Mes V neuron, which provides strong monosynaptic excitation to a postsynaptic TMN (Trueblood et al., 1996). We used an oversimplified, yet realistic, construct and tested how irregularities in ongoing burst patterns in the sensory Mes V neuron, such as during rhythmic jaw movements, could exclusively modulate the discharge patterns in a TMN. In Figure 8*a*, we first illustrate rhythmic burst discharge, such as in a WT Mes V neuron reproduced by the model. Such a regular pattern was converted into irregular/arrhythmic discharge observed in the mSOD1 Mes V neurons by a 25%–50% reduction in the Mes V persistent and resurgent Na⁺ conductances, and addition of subthreshold stochastic inputs to enhance burst irregularities (Fig. 8*b*) (see also Venugopal et al., 2019). We further assumed that the TMN model displays dendritic Ca²⁺ currents, which can mediate plateau potentials and membrane bistability, often observed in brainstem and spinal MNs (Hsiao et al., 1998, 2005; Lee and Heckman, 1998). The model Mes V directly excited the dendritic compartment of the TMN, which in turn depolarized the electrically coupled TMN soma. We further assumed that such depolarization drives dendritic Ca²⁺-mediated plateau potentials, as would occur during NMDA receptor activation (Hsiao et al., 2002; Manuel et al., 2012). Activation of Ca²⁺ plateau depolarized the TMN soma and enabled sensorimotor synchronization driving downstream motor discharge (Fig. 8*c*). However, when the sensory patterns were irregular, such synchronization was perturbed (Fig. 8*d*). In particular, the shorter burst intervals mimicking mSOD1 discharge patterns induced instances of asynchronous self-sustained discharge in the TMN (Fig. 8*d*, dashed boxes). In our model, this was due to inadequate deactivation of the plateau causing dendritic Ca²⁺ currents in the MN between sensory bursts. Although under *in vivo* conditions, other mechanisms, such as synaptic inhibition and Ca²⁺-activated K⁺ currents, might effectively regulate Ca²⁺ plateaus and jaw reflexes (e.g., Inoue et al., 1994; Hultborn et al., 2003; Li and Bennett, 2007; Venugopal et al., 2012), our results highlight that rhythmic sensory timing, or lack thereof, can modulate sensorimotor synchronization. Together, one putative consequence of proprioceptive sensory irregularities leads to an asynchronous sustained motor discharge and may partly contribute to looming dysfunctions, such as muscle fasciculations, which are common symptoms in ALS (Hirota et al., 2000; Vucic and Kiernan, 2006) (see summary in Fig. 8*e*).

Discussion

In this study, we focused on SOD1^{G93A} mutation-induced modifications to proprioceptive sensory neurons (Mes V cells), at an early time point when MN dysregulation was previously reported (Venugopal et al., 2015). There are four main findings: (1) an early circuit-specific reduced excitability exclusively in the trigeminal proprioceptive sensory neurons in comparison with mechanoreceptive, nociceptive, and visual sensory neurons in age-matched mSOD1 mice; (2) impaired bursting in the proprioceptive Mes V neurons associated with a downregulation of Nav1.6 Na⁺ currents and ion channels; (3) rescue of normal burst patterns in the mSOD1 Mes V neurons upon restoration of Na⁺ currents; and (4) computational model-based prediction of an effect of aberrant sensory gating on MN discharge patterns. Considering these results, we discuss the consequences on disease development and progression.

Circuit-specific vulnerability in ALS

Selective vulnerability is a hallmark of neurodegenerative diseases (Double et al., 2010); however, the factors that determine such selectivity are elusive and remain poorly understood. In ALS, as also with normal aging (Kanning et al., 2010), MNs within a motor pool present selective and preferential vulnerability (Frey et al., 2000; Pun et al., 2006; Hegedus et al., 2007; Saxena and Caroni, 2011). Specifically, α-MNs forming the fast-fatigable motor units preferentially die followed by fast fatigue-resistant MNs, whereas the neighboring slow α-MNs and γ-MNs remain resistant to degeneration (Hegedus et al., 2007, 2008; Lalancette-Hebert et al., 2016). Although multiple intrinsic factors, such as cell size (Dukkipati et al., 2018), Ca²⁺ buffering capacities (von Lewinski and Keller, 2005), synaptic organization (Nimchinsky et al., 2000; Lorenzo et al., 2006), and gene and protein expression patterns (Brockington et al., 2013; Comley et al., 2015), could govern selective vulnerability, the underlying network architecture can be a crucial determinant. For instance, a lack of muscle spindle afferent terminals on γ- and ocular MNs has been suggested as a mechanism of disease resistance (Keller and Robinson, 1971; Lalancette-Hebert et al., 2016). These afferents relay the proprioceptive sensory feedback, which modulates MN activity (the stretch reflex) during muscle force generation via glutamatergic excitation (e.g., Chandler, 1989; Mentis et al., 2011). Ablation of such spindle afferents significantly delayed MN death and disease progression in the SOD1^{G93A} mouse model (Lalancette-Hebert et al., 2016). Curiously, even in invertebrate SOD1-knock-in model systems, muscle spindle afferents act as an early trigger for MN degeneration (Held et al., 2019). This suggests the possibility that the proprioceptive feedback may indeed represent a phylogenetically conserved pathway of disease vulnerability. Furthermore, in a mouse model for spinal muscular atrophy, a related MN disease, proprioceptive inputs play a predominant role in triggering MN degeneration (Mentis et al., 2011). Advancing these findings, our comparative analysis of excitability of multiple sensory neurons revealed a SOD1 mutation-driven modification exclusively in the proprioceptive sensory neurons. Although non-motor and non-cell-autonomous triggers have been implicated for MN death in ALS (Boillée et al., 2006a), to date, there is no direct evidence for exclusive circuit elements showing intrinsic electrophysiological abnormalities. In combination with our previous report on early dysregulation of MN excitability (Venugopal et al., 2015), the present results implicate a defective sensorimotor network in ALS, which may represent a convergent neuroanatomical pathway across multiple neurodegenerative MN diseases.

Figure 8. Computer-based model predicts irregular sensory-driven motor asynchrony. **a**, Rhythmic burst pattern in a WT Mes V sensory neuron (top) is reproduced in a conductance-based Mes V neuron model (bottom). Top and bottom right, Corresponding time series graphs of IELs (green circles). **b**, Burst irregularities in a mSOD1 Mes V neuron (top trace) is reproduced in the model neuron (bottom) by a 25% reduction in persistent Na⁺ conductance and a 50% reduction in resurgent Na⁺ conductance together with a slight increase in membrane potential noise. Top and bottom right, Corresponding time series graphs of IELs (green circles). **c**, **d**, Simulated postsynaptic membrane potentials in the motor dendrite (cyan) and soma (blue) for WT (**c**) and mSOD1 (**d**) sensory patterns (top, magenta). Dashed rectangles represent frequent occurrences of asynchronous self-sustained motor discharge patterns. **e**, Schematic showing putative motor asynchrony and dysfunctional discharge patterns resulting from burst irregularities in the primary Ia afferent neuron.

Excitability change: an early disease compensation and a marker for vulnerable circuits

Neurodegenerative diseases involve a protracted phase of progressive decline in the functional homeostasis of vulnerable neurons (Roselli and Caroni, 2015). In mouse models of ALS, hyperexcitability represents the earliest form of homeostatic disruption in brainstem, spinal, and corticomotor neurons, which are vulnerable to degeneration (Pieri et al., 2003; Amendola et al., 2007; Bories et al., 2007; van Zundert et al., 2008; Vucic et al., 2009; Quinlan et al., 2011; Venugopal et al., 2015). These results prompted a hyperexcitability-driven excitotoxicity hypothesis for cell death in ALS (Durand et al., 2006; van Zundert et al., 2012). Although attractive, a causal relationship between hyperexcitability and subsequent neurodegeneration is not well supported (e.g., Leroy et al., 2014; Simon et al., 2016), and indeed recent evidence in MNs proposes a transition into hypoexcitability at adult stages (Delestrée et al., 2014; Martínez-Silva et al., 2018). Alternatively, it is likely that early hyperexcitability represents a compensatory mechanism of survival in normally low excitable cells since enhancing intrinsic excitability promotes neuroprotection in these MNs (Saxena et al., 2013). Early compensation is also possible in slow MNs, which normally show high excitability and are resistant to degeneration in ALS. Some of these slow MNs were hypoexcitable early in the SOD1^{G93A} mouse (Venugopal et al., 2015). Such early hypoexcitability in surviving

MNs likely represents a way to moderate energy demands for spike generation, which can be costly (e.g., Le Masson et al., 2014). This in turn could support cell survival in a disease background. A similar strategy could be used by the proprioceptive sensory neurons shown here, which are involved in ongoing reflex control. We further speculate that hypoexcitable shift in the set point operation of neurons within a vulnerable circuitry could be triggered by disease-induced changes in astroglial cells, which monitor extracellular ionic homeostasis (e.g., Ca²⁺ and K⁺) as well as bursting in Mes V neurons (Kadala et al., 2015; Morquette et al., 2015). Such crosstalk leading to compensatory intrinsic excitability changes needs to be further clarified. Notwithstanding a compensatory function, early excitability changes seem to be clear markers of vulnerable circuitry. For instance, disease-resistant MNs (e.g., oculomotor neurons) (Venugopal et al., 2015) or nonproprioceptive sensory neurons as shown here do not present early modifications in excitability. In contrast, a specific set of bursting proprioceptive Mes V neurons were hypoexcitable, which form a monosynaptic jaw stretch reflex circuitry with vulnerable TMNs. Such reduced presynaptic sensory excitability could also be a trigger for compensatory hyperexcitability in the postsynaptic TMNs (Venugopal et al., 2015). However, it is yet unclear whether and how such sensory alterations contribute to MN degeneration (see, e.g., Lalancette-Hebert et al., 2016). Together, early excitability changes are crucial markers of vulner-

able circuitry and may represent an early compensation to meet energy demands and likely support cell survival early during disease development.

Altered activity patterns and ionic mechanisms as pathophysiological substrates

Altered phasic/burst patterns, reduced excitability, and loss of excitation are defining features of vulnerability to neurodegeneration (Roselli and Caroni, 2015). Here we present a unique result highlighting early irregular and impaired burst discharge in proprioceptive sensory neurons forming a vulnerable sensorimotor circuitry. One consequence of irregular bursting could affect sensory gating of postsynaptic motor discharge as shown by our computational model. Second, altered Ia afferents could directly alter static versus dynamic spindle inputs important for activation of slow versus fast persistent Ca^{2+} currents in MNs during a stretch reflex (Manuel et al., 2007). Moreover, impaired burst duration and frequencies together with reductions in persistent Na^{+} current could reduce extracellular Ca^{2+} and affect astrocyte-mediated Ca^{2+} buffering in the Mes V sensory nucleus (Morquette et al., 2015). Alternatively, reduced persistent Na^{+} and bursting could be a direct consequence of mSOD1 astrocyte dysfunction (Su et al., 2001). Curiously, retinal ganglion cells, which also rely on persistent Na^{+} current for repetitive spiking (Boiko et al., 2003; Van Wart and Matthews, 2006a,b), did not show altered rhythmicity as measured by the autocorrelation function. These observations suggest that the mutation could induce observed changes via altered extrinsic factors, such as astrocyte dysfunction or by altered intrinsic factors, such as abnormal protein synthesis contributing to the observed Na^{+} channelopathy in Mes V neurons. Furthermore, in the trigeminal system, the nonproprioceptive sensory neurons in the TG were resistant to abnormalities in the SOD1^{G93A} mouse. Given that the TG and Mes V neurons share ontogeny (Lazarov, 2002), an exclusive change only in Mes V supports extrinsic triggers within the jaw stretch reflex circuitry causing abnormal excitability. However, we note that definitive studies measuring Mes V excitability under physiological conditions during natural movements, such as chewing, which dynamically modulate spindle activity would be essential to fully evaluate functional deficits resulting from proprioceptive afferent dysfunction.

Secondary effects of altered burst patterns could lead to recruitment of distinct types of voltage-dependent Ca^{2+} currents; resulting alterations in intracellular Ca^{2+} levels and kinetics can activate distinct CREB-dependent gene expression (Wheeler et al., 2012) and differentially activate nuclear translocation transcription factors, such as the NFAT family, important in immune response (Hernández-Ochoa et al., 2007). Importantly, reduced excitability can deplete intracellular ER Ca^{2+} stores and disrupt protein homeostasis (e.g., a ubiquitin-dependent degradation of transcription factors important for synaptic plasticity) (Lalonde et al., 2014). Together, we suggest that identifying presymptomatic excitability changes, their upstream physiological triggers (e.g., astrocyte dysfunction), and downstream effectors (e.g., CREB-mediated gene expression changes) can elucidate the cascade of events leading to neurodegeneration. Additionally, excitability changes also represent early markers of disease vulnerability in ALS.

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