Sensory Neurons Do Not Induce Motor Neuron Loss in a Human Stem Cell Model of Spinal Muscular Atrophy

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Abstract

Spinal muscular atrophy (SMA) is an autosomal recessive disorder leading to paralysis and early death due to reduced SMN protein. It is unclear why there is such a profound motor neuron loss, but recent evidence from fly and mouse studies indicate that cells comprising the whole sensory-motor circuit may contribute to motor neuron dysfunction and loss. Here, we used induced pluripotent stem cells derived from SMA patients to test whether sensory neurons directly contribute to motor neuron loss. We generated sensory neurons from SMA induced pluripotent stem cells and found no difference in neuron generation or survival, although there was a reduced calcium response to depolarizing stimuli. Using co-culture of SMA induced pluripotent stem cell derived sensory neurons with control induced pluripotent stem cell derived motor neurons, we found no significant reduction in motor neuron number or glutamate transporter boutons on motor neuron cell bodies or neurites. We conclude that SMA sensory neurons do not overtly contribute to motor neuron loss in this human stem cell system.

Introduction

Spinal muscular atrophy (SMA) is an autosomal recessive disorder that leads to muscle weakness, respiratory distress, paralysis, and early death due at least in part to loss of motor neurons in the spinal cord. SMA is most often caused by deletion of the survival motor neuron (SMN) gene. Humans have two copies of SMN: a telomeric copy (SMN1) and a centromeric copy (SMN2) [1]. SMN1 produces a full-length protein found in both the cytoplasm and the nucleus and is involved in biogenesis of RNA proteins, RNA transcription, and pre-mRNA splicing [2–4]. In SMA, total full-length SMN protein is drastically reduced due to the absence of SMN1, and SMN2 is not able to fully compensate due to generation of an alternative spliced protein (SMNΔ7) [5–8].

Although SMN is lost in all cell types, it remains to be fully understood why motor neurons are particularly vulnerable. Recent studies have suggested that SMN is important in U12-dependent splicing events necessary for proper motor neuron function [9], but evidence also suggests that other cell types are affected, including astrocytes, sensory neurons, Schwann cells, and skeletal muscle that may each contribute to or exacerbate motor neuron loss [10–18]. In this regard, we have shown that motor neurons generated from SMA patient derived induced pluripotent stem cells (iPSCs) show significant loss through an apoptotic process by 6 weeks in culture [19,20]. Moreover, we recently showed that astrocytes derived from SMA iPSCs are activated and exhibit abnormal calcium homeostasis and reduced growth factor production prior to the overt motor neuron loss [16]. Other studies using human patients and nerve biopsies have shown reduced nerve conduction velocity and inexcitability of sensory neurons [21–23]. Similarly, there is evidence from a mouse model of SMA that dorsal root ganglia (DRG) sensory neurons have reduced neurite outgrowth compared to control [11] and spinal afferent synaptic connections onto the motor neurons are reduced before the onset of significant motor neuron loss [12,14]. Loss of the proprioceptive sensory neurons and primary afferent boutons was more severe for synapses formed on motor neurons projecting to proximal muscles [12,14], which is consistent with the pronounced atrophy observed in the limbs, but the significance of these data is debated as other studies demonstrated that sensory neuron deficits in SMA were a consequence of motor neuron dysfunction, rather than a cause [10,13,24,25]. Data from Drosophila have shown that SMN replacement is necessary in sensory neurons and interneurons to restore motor neuron and muscle function [15]. Finally, very recent studies have found that mRNAs related to synaptic formation and sensory-motor circuitry are dysregulated in SMA mice spinal cord prior to motor neuron loss [26]. Decades of research confirm that direct and indirect sensory afferent innervations onto spinal cord motor neurons, such as in the spinal reflex circuit, are critical to motor neuron function and subsequent motor output [27]. Synaptic activity is critical for neuronal survival through inhibition of apoptotic cascades [28–30], and significant evidence exists suggesting that SMA motor neurons die through an apoptotic mechanism [20,31–33]. Taken together, these data suggest that non-motor neuron cell types may actively contribute to the SMA disease phenotype by directly

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affecting motor neuron survival, but further investigation is needed. Traditional in vitro and in vivo animal models are readily available to study neurodegenerative diseases, and the ground-breaking iPSC technology has opened up additional avenues of exploration into human development and disease [34–36]. Here we generated sensory neurons from SMA and control iPSCs in an effort to determine if SMA iPSC-derived sensory neurons directly contribute to motor neuron loss. We followed an established sensory neuron differentiation protocol [37] and show that SMA and control iPSCs generate similar numbers of peripherin positive sensory neurons, with small populations of these neurons being nociceptors and proprioceptors. Using direct co-culture methods, we asked whether SMA iPSC-derived sensory neurons caused loss of healthy iPSC-derived motor neurons indicating a direct role of sensory neurons in SMA disease progression. Neither total motor neuron number nor vesicular glutamate transporter staining on healthy iPSC-derived motor neurons was significantly reduced when cultured with SMA iPSC-derived sensory neurons. Therefore, we conclude that interactions with SMA iPSC-derived sensory neurons are not sufficient to induce motor neuron loss in a human iPSC system.

Materials and Methods

Cell culture

Two independent control iPSC lines (4.2 and 21.5) and two independent SMA iPSC lines (3.6 and 7.12) were used [19,20,38]. iPSCs were grown in feeder free conditions on Matrigel substrate (BD Biosciences) in Nutristem medium (Stemgent). Neural stem cells (EZ Spheres) were generated by lifting intact colonies after disase treatment (1 mg/ml Gîbco) and placing them directly into a human neural progenitor growth medium (Stemline, Sigma) supplemented with 100 ng/ml basic fibroblast growth factor (FGF-2, Millipore), 100 ng/ml epidermal growth factor (EGF, Miltenyi Biotec), and 5 µg/ml heparin (Sigma) in ultra-low attachment flasks and were passaged weekly using a chopping technique as previously described [39].

Sensory neuron differentiation

Induction of sensory neurons was accomplished using an established protocol [37]. Briefly, human iPSC colonies were grown to confluence on poly-ornithine/Matrigel coated coverslips. Once 80–90% confluence was reached, the Nutristem medium was replaced with knockout serum replacement (KSR) medium supplemented with 1 µM dorsomorphin (Tocris) and 10 µM SB431542 (Stemgent) on days 0–5. KSR medium was prepared by supplementing 920 ml of Knockout DMEM, with 150 ml knockout serum replacement, 1 mM L-glutamine, 100 µM MEM nonessential amino acids, 1% antibiotic-antimycotic (anti-anti) (all from Life Technologies), and 0.1 mM β-mercaptoethanol (Sigma). Cells were fed daily and N2 medium (1:1 DMEM/F12, 1% N2, 1% NEAA, heparin (1 mg/ml), 1% anti-anti) supplemented with retinoic acid (0.1 µM; Sigma) for one week followed by supplementation with retinoic acid (0.1 µM) and purmorphamine (1 µM; Stemgent) for an additional week changing half the medium every 3–4 days. After two weeks of differentiation, the spheres were dissociated with accutase (Millipore) and plated onto 2 week old sensory neuron cultures at a density of 30,000 cells per coverslip. Long term medium consisted of N2 medium supplemented with BDNF (25 ng/ml), GDNF (25 ng/ml), NGF (25 ng/ml), ascorbic acid (200 µg/ml), cAMP (1 µM). The medium was replaced every 2 days.

Immunocytochemistry

Plated cells were fixed in 4% paraformaldehyde (PFA; Fisher Scientific) for 20 minutes at room temperature and rinsed with PBS. Non-specific labeling was blocked and the cells permeabilized with 5% normal goat serum (Millipore) and/or 5% normal donkey serum (Millipore) and 0.2% Triton X-100 (Sigma) in PBS for 30 minutes at room temperature. Cells were rinsed with PBS and then incubated with primary antibodies for one hour at room temperature or overnight at 4°C. Cells were subsequently labeled with the appropriate fluorescently-tagged secondary antibodies. Hoechst nuclear dye was used to label nuclei. Primary antibodies used were rabbit anti-Peripherin (Millipore, AB1530), mouse anti-BIII Tubulin (Tuj1, Promega, G7121), rabbit anti-NTRK1 (Millipore, 06-574), rabbit anti-TRPV1 (Novus Biologicals, NB1P-97417), mouse anti-GFAP (Cell Signaling, 3670), rabbit anti-Parvalbumin (Calbiochem, PC255L), mouse anti SMI32R (Covance, SMI-32R), guinea pig anti-VGlut1 (Millipore, AB5905), and myelin protein zero (Protein Tech, 10572-1-AP). Secondary antibodies included donkey anti-mouse AF488 (Invitrogen, A21202), goat anti-rabbit Rhod (Invitrogen, R6394), and goat anti-guinea pig AF480 (Life Technologies, A11073).

Imaging and quantification

At least five images were taken on each of at least three different fluorescently labeled coverslips per timepoint using a Nikon inverted microscope and Spot imaging software. All experiments

Calcium imaging

iPSC-derived sensory neuron cultures were cultured for 2 weeks and functionally tested using ratiometric live-cell calcium imaging using dual-wavelength fluorescent calcium indicator FURA-2AM (Invitrogen) to detect intracellular calcium levels. Cells were loaded with 2.5 µL/ml FURA-2AM in 2% BSA (Sigma) for 1 hour at room temperature, washed in extracellular buffer for 20 minutes, and mounted onto a perfusion chamber and superfused with extracellular buffer at 6 ml/min. Coverslips were superfused with extracellular buffer containing 50 mM KCl for 30 secs. Fluorescence images were taken with a cooled CCD camera (CoolSNAP FX, Photometrics, Tucson, AZ) before and after stimulation. Metafluor imaging software was used to detect and analyze intracellular calcium changes throughout the experiment (Molecular Devices, Sunnyvale, CA) where a ≈20% increase in intracellular calcium from baseline constituted a response. Extracellular buffer contained 150 mM NaCl, 10 mM HEPES, 5.6 mM KCl, 1 mM MgCl₂, and 2 mM CaCl₂ (all from Sigma).

Motor neuron differentiation/co-culture

Motor neuron differentiation from the EZ spheres was performed as previously described [19,39]. Briefly, the neural progenitor growth medium was replaced with neural induction medium (1:1 DMEM/F12, 1% N2, 1% NEAA, heparin (1 mg/ml), 1% anti-anti) supplemented with retinoic acid (0.1 µM; Sigma) for one week followed by supplementation with retinoic acid (0.1 µM) and purmorphamine (1 µM; Stemgent) for an additional week changing half the medium every 3–4 days. After two weeks of differentiation, the spheres were dissociated with accutase (Millipore) and plated onto 2 week old sensory neuron cultures at a density of 30,000 cells per coverslip. Long term medium consisted of N2 medium supplemented with BDNF (25 ng/ml), GDNF (25 ng/ml), NGF (25 ng/ml), ascorbic acid (200 µg/ml), cAMP (1 µM). The medium was replaced every 2 days.
were repeated at least 2 times. The images were analyzed for antigen specificity using MetaMorph Software (Molecular Devices Inc.). Data were statistically analyzed via one-way ANOVA or Student’s t-test as appropriate with Prism software, \( \alpha = 0.05 \) (GraphPad). Data are presented as the average \( \pm \) S.E.M. using two independent control lines and two independent SMA lines.

**Ethics statement**

The privacy of the fibroblast cell sources from which the iPSCs were derived is maintained by Coriell Institute for Medical Research. The use of iPSCs has been approved by the Stem Cell Research Oversight Committee at the Medical College of Wisconsin.

**Results and Discussion**

**Sensory differentiation and characterization from SMA and control iPSCs**

To examine the functional properties of SMA sensory neurons, we utilized human induced pluripotent stem cells (iPSCs) derived from SMA patients and healthy individuals [19,20,30]. Control and SMA iPSCs were cultured as adherent monolayers for sensory neuron induction following a previously established protocol [37]. This protocol allows for the efficient generation of peripheral positive sensory neurons within 2 weeks of differentiation. Peripherin labels thin, unmyelinated sensory neurons, which includes both nociceptive and proprioceptive neurons. The cellular compositions derived from the differentiations were analyzed by immunostaining with specific cellular markers at 2, 4, and 6 weeks of culture. Importantly, both control and SMA iPSCs acquired a sensory neuronal phenotype as identified by βIII tubulin (TuJ1) and peripherin (Figure 1A). Quantification of TuJ1 and peripherin positive neurons resulted in approximately 35% TuJ1+ and 20% peripherin+ cells over the course of the differentiation period from both control and SMA iPSCs (Figure 1B, C). Previous studies have found decreased neurite extension from isolated DRG sensory neurons from SMA mice [11] as well as axon degeneration from the sural nerve in human SMA patients [23]. However, neither of these characteristics was observed in human SMA iPSC-derived sensory neurons (Figure 1A, D). Interestingly, we used ratiometric live cell calcium imaging to test functional responses to KCl depolarization stimulation and found a significant reduction in the intracellular calcium response in SMA iPSC-derived sensory neurons compared to control iPSC-derived sensory neurons (Figure 1E–G), which agrees with data collected from one SMA patient indicating a reduced neuronal excitability [22]. The reason for some of the discrepancies between data presented here and previous reports [11,23] is unknown. The sensory neuron differentiation protocol generates neurons similar to DRG neurons [37], which facilitates comparisons with the mouse study [11], but species differences and/or culture conditions may contribute to different results. Additionally, the sural nerve studied by Rudnik-Schoneborn and colleagues [23], which is a sensory nerve found in the leg that innervates the foot may be too distinct from DRG-like neurons to allow direct comparisons.

Peripherin-positive sensory neurons consist of multiple subtypes serving distinct sensory functions such as proprioceptors for limb position and nociceptors for recognizing noxious (i.e. painful) stimuli. We therefore aimed to delineate the subtypes present within our cultures. We first assessed the differentiations for the presence of nociceptors using immunostaining for neurotrophic tyrosine kinase receptor type 1 (NTRK1), also known as TrkA (Figure 2A). We found approximately 20% of the TuJ1 positive neurons in both SMA and control iPSC cultures expressed NTRK1 (Figure 2B). Since previous studies have shown proprioceptive sensory neurons to be affected in SMA [12,14], we examined whether any proprioceptors were present following differentiation. We used immunostaining for parvalbumin, a marker for large type I proprioceptive sensory neurons and found approximately 5–10% of the neuronal population were parvalbumin positive (Figure 3A, B). Importantly, these data show that unaffected control and SMA iPSC-derived sensory neurons are capable of generating similar numbers of nociceptive and proprioceptive sensory neurons with no significant SMA iPSC-derived sensory neuron loss over time compared to control, regardless of sensory neuron subtype (Figures 1–3). It should be noted that the sensory neuron differentiation in our hands was not as efficient as previously described by others [37]. This could be due to inherent variability of iPSCs, but because all control and SMA iPSC lines used here generated consistent sensory neuron populations, this seems unlikely. However, another possible reason for lowered differentiation efficiency could be due to differences in quantification techniques used; differentiation percentages were based on immunocytochemistry results in unsorted cell cultures rather than by FACS as done by Chambers et al [37]. Nevertheless, our data show no difference between control and SMA iPSCs in regards to sensory neuron differentiation.

Because the differentiation protocol generates a mixed population of cells, it is not fully clear what other cellular subtypes are present in the cultures. However, we did analyze the differentiations for the presence of glial cells using GFAP as a marker of Schwann cells, which revealed abundant GFAP+ cells dispersed throughout the culture (Figure 4). Although we cannot completely rule out these are GFAP+ astrocytes, the differentiation protocol proceeds through a Sox10+ neural crest developmental stage (Figure S1 and [37]), so these are more likely to be Schwann cells. We have previously shown that SMA iPSC-derived astrocytes exhibit an activated morphology and show defects in calcium signaling and growth factor expression compared to control iPSC-derived astrocytes [16], which may directly contribute to motor neuron loss. SMA mouse Schwann cells have shown significant myelination defects [40], which necessitates further investigation into iPSC-derived Schwann cell function. However, the stable iPSC-derived sensory neuron cultures over time suggests that functional impairments in SMA iPSC-derived Schwann cells may be minimal or that SMA iPSC-derived sensory neurons are more resistant than motor neurons to glial-induced damage.

**SMA iPSC-derived sensory neurons do not induce motor neuron loss**

Data from SMA animal models suggest a potential role for sensory neurons in motor neuron demise [12,14,15,26]; therefore, we used direct co-culture methods to test whether SMA iPSC-derived sensory neurons cause loss of healthy control iPSC-derived motor neurons. We followed established differentiation protocols to generate motor neurons from control iPSC lines well characterized in our lab [16,19,20,39]. Control iPSCs were patterned toward motor neurons for 2 weeks followed by direct plating onto 2 week differentiated sensory neuron cultures derived from control and SMA iPSCs. Once plated, the cells were cultured together for an additional 2–6 weeks to allow for maturation. Using immunostaining at 4, 6, and 8 weeks of total differentiation (i.e. 2, 4, and 6 weeks of co-culture), we assessed the numbers of motor neurons and sensory neurons in co-culture utilizing the motor neuron marker SMI-32 and the peripheral sensory neuron marker peripherin. Consistent with being cultured alone (Figure 1), control and SMA iPSC cultures showed similar numbers of...
sensory neurons in the presence of control motor neurons (Figure 5A, B). Moreover, control motor neurons revealed no statistical difference in cell survival when cultured in the presence of either control or SMA iPSC-derived sensory neurons (Figure 5C). Although a significant reduction in calcium response to KCl induced depolarization was found in SMA iPSC-derived sensory neurons (Figure 1E), this does not appear to negatively impact motor neuron survival. The sensory neuron differentiation medium contains NGF as well as higher concentrations of GDNF and BDNF than what is typically found in the motor neuron differentiation medium.
differentiation medium. Culture of SMA iPSC-derived motor neurons alone in the sensory neuron differentiation medium does not rescue motor neuron death (Figure S2). Therefore, the absence of a motor neuron phenotype in control motor neuron/SMA sensory neuron iPSC co-cultures is likely not due to culture medium supplementation.

Figure 2. Generation of nociceptive neurons from control and SMA iPSCs. (A) Control and SMA iPSCs generate nociceptive neurons as indicated by NTRK1 (red) and Tuj1 (green). Nuclei are labeled with Hoechst nuclear dye (blue). (B) There was no difference in neuron differentiation between control and SMA iPSC cultures at either 4 or 6 weeks of differentiation. n.s. = not significant by ANOVA. Scale bar = 50 μm. doi:10.1371/journal.pone.0103112.g002

Figure 3. Generation of proprioceptive neurons from control and SMA iPSCs at 6 weeks of differentiation. (A) Both control and SMA iPSCs generate proprioceptive neurons as indicated by parvalbumin (red) and Tuj1 (green). Nuclei are labeled with Hoechst nuclear dye (blue). (B) There was no difference in parvalbumin+ neuron number between control and SMA iPSC cultures. n.s. = not significant by t-test. Scale bar = 20 μm. doi:10.1371/journal.pone.0103112.g003
SMA iPSC-derived sensory neurons do not affect VGlut1 abundance

Previous work has shown reduced VGlut1+ boutons on the dendrites of SMA motor neurons [10,12,14,25]; therefore, we sought to determine if this was also the case in the SMA/control iPSC co-culture system. To test whether there was a global decrease in VGlut1, we immunostained SMA sensory neuron/control motor neuron co-cultures with SMI-32 and VGlut1 and examined the number of VGlut1+ puncta on both the neurites and cell bodies of SMI-32+ motor neurons (Figure 6A). Quantification of VGlut1 resulted in a trend toward fewer VGlut1+ puncta on control iPSC-derived motor neurons cultured in the presence of SMA iPSC-derived sensory neurons over 6–8 weeks in culture; however, this trend was not statistically different from co-cultures of control iPSC-derived sensory neurons with control iPSC-derived motor neurons (Figure 6B, C). It is possible that after longer times in co-culture the reduction in VGlut1+ boutons would reach significance. However, culturing differentiated iPSC for extended periods of time can lead to sub-optimal culture conditions which may compromise neuronal health and complicate data interpretations. SMA iPSC-derived motor neurons are significantly reduced after 6 weeks in culture [19,20], but we did not test here whether the addition of SMA iPSC-derived sensory neurons would exacerbate this phenotype. Instead, we tested whether VGlut1 synaptic loss on motor neurons occurs in the absence of sensory neuron co-culture. At 4 weeks of motor neuron differentiation we found that VGlut1+ puncta are significantly reduced on both the neurites and cell bodies of SMA iPSC-derived motor neurons compared to control (Figure S3A, B). These data are consistent with the idea that synapse loss precedes motor neuron death [14], but the loss of VGlut1+ puncta is precipitated by the motor neuron and not the sensory neuron. Taken together, our data suggest that SMA iPSC-derived sensory neurons have a negligible effect on motor neuron afferent innervation and are not sufficient to induce motor neuron loss in this iPSC system.

**Figure 4.** Peripheral glial cells are present in the sensory neuron cultures. GFAP+ glial cells (green) are simultaneously generated during the sensory neuron differentiation in both control and SMA iPSC cultures. Nuclei are labeled with Hoechst nuclear dye (blue). Scale bar = 50 μm. doi:10.1371/journal.pone.0103112.g004

**Figure 5.** The presence of SMA iPSC-derived sensory neurons does not induce motor neuron loss at 4, 6, and 8 weeks of differentiation. (A) Co-culture of control iPSC-derived SMI-32+ (green) motor neurons (MN) with either SMA iPSC-derived peripherin+ (red) sensory neurons (SN) or control iPSC-derived peripherin+ (red) SNs does not induce motor neuron loss. Nuclei are labeled with Hoechst nuclear dye (blue). Quantification is shown in B and C. n.s. = not significant by ANOVA. Scale bar = 50 μm. doi:10.1371/journal.pone.0103112.g005
Conclusion

Using a human iPSC system, we show that SMA and control iPSCs can be differentiated at equivalent levels into peripheral sensory neurons containing markers of both nociceptors and proprioceptors. As opposed to SMA iPSC-derived motor neurons that die around 6 weeks in culture [19,20], there is no reduction in neurite length or loss of SMA iPSC-derived sensory neurons over time, which is consistent with the more selective motor neuron loss observed in mouse models and human SMA patients. Moreover, neither the presence of SMA iPSC-derived sensory neurons nor the slight reduction in VGluT1+ bouton numbers on healthy motor neurons is sufficient to induce motor neuron loss. Together with the evidence of VGluT1+ bouton loss on SMA iPSC-derived motor neurons in the absence of sensory neurons, these results suggest that degeneration of the sensory-motor system is likely caused by motor neuron deficits rather than sensory neuron deficits.

Supporting Information

Figure S1 GFAP+ glial cells (green) produced during the differentiation process express Sox10 (red) indicating these are likely Schwann cells. Images in A and B are taken from two different SMA iPSC lines. Nuclei are labeled with Hoechst nuclear dye (blue). Scale bar = 50 μm. (TIF)

Figure S2 Culturing SMA iPSC-derived motor neurons in sensory neuron medium does not prevent motor neuron loss. The average number of SMI-32+/Tuj1+ motor neurons at 6 weeks of differentiation was significantly reduced in SMA iPSC cultures maintained in either the standard motor neuron (MN) medium or in the sensory neuron (SN) medium compared to control iPSCs. There was no difference between the two growth conditions for the SMA iPSC cultures. **p = 0.0051 by ANOVA. n.s. = not significant by ANOVA. (TIF)

Figure S3 VGluT1+ puncta are reduced on SMA iPSC-derived motor neurons in the absence of sensory neuron innervation. Four
week differentiated SMA iPSC-derived motor neurons exhibit significantly fewer VGlut1+ puncta on both the cell soma (A) and neurites (B) compared to control iPSC-derived motor neurons. *p = 0.0267 by Student’s t-test; **p = 0.0032 by Student’s t-test. (TIF)

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Author Contributions
Conceived and designed the experiments: AJS ADE. Performed the experiments: AJS ADE. Analyzed the data: AJS ADE. Contributed reagents/materials/analysis tools: ADE. Contributed to the writing of the manuscript: AJS ADE.