Hidden hearing loss in a Charcot-Marie-Tooth type 1A mouse model

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Hidden hearing loss (HHL), a recently described auditory neuropathy characterized by normal audiometric thresholds but reduced sound-evoked cochlear compound action potentials, has been proposed to contribute to hearing difficulty in noisy environments in people with normal hearing thresholds and has become a widespread complaint. While most studies on HHL pathogenesis have focused on inner hair cell (IHC) synaptopathy, we recently showed that transient auditory nerve (AN) demyelination also causes HHL in mice. To test the effect of myelinopathy on hearing in a clinically relevant model, we studied a mouse model of Charcot-Marie-Tooth type 1A (CMT1A), the most prevalent hereditary peripheral neuropathy in humans. CMT1A mice exhibited the functional hallmarks of HHL together with disorganization of AN heminodes near the IHCs with minor loss of AN fibers. These results support the hypothesis that mild disruptions of AN myelination can cause HHL and that heminodal defects contribute to the alterations in the sound-evoked cochlear compound action potentials seen in this mouse model. Furthermore, these findings suggest that patients with CMT1A or other mild peripheral neuropathies are likely to suffer from HHL. Furthermore, these results suggest that studies of hearing in patients with CMT1A might help develop robust clinical tests for HHL, which are currently lacking.

Introduction

Many people report having hearing difficulty in noisy environments despite presenting with normal audiometric thresholds (1–5). There is substantial evidence associating this auditory pathology with aging or with a history of "mild" noise exposures that only induce temporary increases in hearing thresholds (6–13). Studies on animal models demonstrate that such mild noise exposures do not result in hair cell (HC) loss but instead destroy a subset of synapses between inner HCs (IHCs) and spiral ganglion neurons (SGNs) (14). Moreover, in animals, IHC synaptopathy precedes HC loss during aging, even in the absence of noise overexposure (14–20). Physiological recordings show that animals with IHC synaptopathy have normal hearing thresholds but reduced amplitudes of sound-evoked cochlear compound action potentials (15, 16, 18, 21), a condition that has been termed hidden hearing loss (HHL) (22–24). Importantly, we recently demonstrated that young mice with IHC synaptopathy independent of noise exposure have defects in temporal auditory processing (25), and histopathological studies have shown that age-related synaptopathy also occurs in humans (26, 27), suggesting that HHL might contribute to the hearing difficulties in people with normal thresholds. The compelling information generated from noise-exposed and aging animals led to the sense that IHC synaptopathy might be the only cause of HHL. However, more recently, we showed that transient auditory nerve (AN) demyelination causes a similar hearing deficit in mice without affecting IHC synapses (28), suggesting that demyelinating peripheral neuropathies could also cause HHL. To begin to test this possibility in clinically relevant experimental models, we focused on mouse models of Charcot-Marie-Tooth type 1 (CMT1) disease, the most common hereditary peripheral neuropathy in humans.

CMT1 is a type of CMT, a group of hereditary peripheral neuropathies that are associated with a range of genetic lesions and pathogenic mechanisms (29, 30). CMT1, which is considered a demyelinating CMT, includes multiple subtypes (from A to E) arising from alterations in diverse myelin-related genes (31). CMT1A, the predominant subtype (>50% of CMT1 cases), is caused by the duplication of *PMP22*, a gene that encodes an integral membrane glycoprotein essential for the formation and maintenance of compact

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work.

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myelin (32, 33). Interestingly, recent studies suggest that patients with CMT1A might suffer from hearing difficulties consistent with HHL (34–37). However, this has not been confirmed due to the lack of validated clinical HHL tests in humans (23, 38). In contrast, patients with CMT1E, which is caused by a wide variety of *PMP22* point mutations (31) and leads to more severe neuropathy phenotypes with earlier onset, suffer from overt hearing loss (elevated auditory thresholds) (39–43). Furthermore, a CMT1E mouse model, also referred to as Trembler-J, has been shown to have profound deafness including AN fiber loss (44, 45).

Here we used CMT1A (46) and CMT1E (47) mouse models to explore the specific effect of these hereditary peripheral neuropathies on inner ear structure and function. Our functional studies demonstrate that, as shown by others (44, 45), CMT1E mice present with early-onset overt hearing loss with profound auditory threshold shifts. In contrast, CMT1A mice exhibit the distinct electrophysiological features of HHL (i.e., normal auditory thresholds but reduced sound-evoked compound action potential amplitudes). However, in contrast to mice with IHC synaptopathy, sound-evoked compound action potential latencies are longer in CMT1A mice. At the structural level, whereas CMT1E present severe loss of AN fibers, the cochlear defects of CMT1A mice are primarily disorganization of AN heminodes near the IHCs and a subtle but significant mislocalization of some nodes of Ranvier (NoR) to the area close to the heminodes.

Together, these results support the hypothesis that mild disruptions of AN myelination can cause HHL and that AN heminodal defects are responsible for the reduced amplitude and longer latencies of sound-evoked compound action potentials seen in the mouse models (28, 48). Our results also support the notion that patients with CMT1A are likely to suffer from HHL and that this type of hearing disorder might affect patients with other types of peripheral neuropathies, including Guillain-Barré syndrome. Based on these findings, we propose that studies of hearing in patients with CMT1A might help in the design and validation of robust clinical tests for HHL. This would circumvent the difficulties encountered in the attempts to develop HHL tests based on self-reporting on historical noise exposure, which have resulted in conflicting conclusions (1, 38, 49).

Results

CMT1A mice have progressive HHL. To determine the auditory phenotypes of CMT1A and CMT1E mice, we recorded distortion product otoacoustic emissions (DPOAEs) and auditory brainstem responses (ABRs) at 1, 2, 3, and 4 months of age (Figure 1A) in the mutants and their WT C57BL/6J littermate controls. ABR waveforms (Figure 1B) reflect the sound-evoked electrical activity along the ascending auditory pathway, from the activation of IHCs and SGNs in the cochlea up to the inferior colliculus in the CNS. ABR thresholds provide information about hearing sensitivity, whereas the amplitude of the first peak of the ABR waveform reflects the magnitude of the sound-evoked SGN activation. DPOAEs are sounds generated by the inner ear in response to sound stimulation, reflecting the activity of outer HCs (OHCs). Together, these measurements provide insight into the integrity of the cochlea. For example, increased ABR thresholds without a change in DPOAEs provides strong evidence for auditory neuropathy (i.e., defects in the function of IHCs and/or SGNs without OHC defects).

As anticipated for C57BL/6J WT, control mice show stable DPOAE and ABR thresholds during their first 4 months of life (Figure 1, C and D) and a small but statistically significant age-related decline in ABR peak I amplitudes (Figure 1E, Figure 2A, Supplemental Figure 1, and Supplemental Figure 2A; supplemental material available online with this article; https://doi.org/10.1172/jci.insight.180315DS1). Their ABR peak I latencies become shorter by 2 months of age, possibly due to the maturation of AN myelination (50, 51) (Figure 1F and Supplemental Figure 2B). Consistent with published reports (44, 45), CMT1E mice have higher ABR thresholds at all ages and frequencies tested (Figure 1D), while their DPOAE thresholds are mildly increased compared with WT mice at 4 months of age only (Figure 1C). Furthermore, CMT1E ABR peak I amplitudes are considerably lower and the latencies longer than in WT mice (Figure 1, E and F; Figure 2A; and Supplemental Figure 1). These findings support a profound auditory neuropathy as early as 1 month of age in CMT1E mice, which then progresses to include OHC dysfunction. In contrast, CMT1A mice have normal ABR and DPOAE thresholds at all ages (Figure 1, C and D), but their ABR peak I amplitudes are already smaller than those in WT by 1 month of age and decline further over time (Figure 1E, Figure 2A, and Supplemental Figure 1), the hallmarks of progressive HHL. Moreover, ABR peak I latencies are longer than those in WT mice at 2 month of age and do not get shorter at later time points (Figure 1F and Figure 2A), suggesting a lack of myelin maturation. The ABR

Figure 1. CMT1A mice have progressive hidden hearing loss, whereas CMT1E have early-onset overt hearing loss. (**A**) Experimental design. CMT1A, CMT1E, and WT C57BL/6J littermates were used. ABRs and DPOAEs were recorded at 1, 2, 3, and 4 months of age, and after the last tests, cochleae were collected and processed for either transmitted electron microscopy or confocal microscopy to evaluate structural changes at the level of the peripheral auditory nerve. (**B**) ABR waveform illustrating the criteria used to measure ABR peaks. Peak I amplitudes are measured relative to the baseline, while peak II to peak V are measured from the top to the bottom right of each peak, respectively. (**C**) CMT1A mice have normal DPOAE thresholds at all time points, while CMT1E mice show mild DPOAE threshold shifts starting at 4 months of age. (**D**) CMT1A mice have normal ABR thresholds while, in CMT1E animals, they are increased at all time points when compared with WT littermates. (**E**) ABR peak I amplitudes are progressively reduced in both CMT1 groups compared with WT mice; however, these reductions are more severe in CMT1E mice (Supplemental Figure 1). (**F**) ABR peak I latencies are longer in both CMT1 groups compared with WT mice, yet CMT1E animals show a more severe phenotype compared with CMT1A mice. WT, *n* = 14–19 mice; CMT1A, *n* = 8–11 mice; CMT1E, *n* = 6–12 mice. DPOAE threshold, ABR threshold, ABR peak I amplitude, or ABR peak I latency at each individual time point were evaluated by 2-way ANOVA followed by Tukey's multiple-comparison test. ABR peak I amplitudes and latencies were measured at 80 dB SPL. ***P* < 0.01; ****P* < 0.001; *****P* < 0.0001 versus WT mice; †††*P* < 0.001; ††††*P* < 0.0001 versus CMT1A mice. Data are shown as mean ± SEM.

waveforms (Figure 2A) also show that the summating potentials (SP), which represent the activation of IHCs, are normal in CMT1A mice but are significantly reduced in 4-month-old CMT1E, likely reflecting the OHC dysfunction seen at this age. The waveforms also illustrate that CMT1E mice have a progressive

Figure 2. Distinct effects of CMT1A and CMT1E on signaling along the ascending auditory pathway. (**A**) Mean ABR waveforms recorded at 1, 2, 3, and 4 months of age show a progressive delay and reduction of the first ABR peak in both CMT1 mouse models. CMT1E phenotype is more severe than in CMT1A mice, affecting the later ABR peaks along the ascending auditory pathway. ABRs shown here are group means in response to 16 kHz tone pips at 80 dB SPL. Data are shown as mean ± SEM. (**B**) Quantification of ABR amplitudes in peak II, III, IV, and V at 4 months of age shows that CMT1E mice have decreased signaling along the ascending auditory pathway while, in CMT1A mice, this signaling is normal compared with WT mice. ABR peak II–V amplitudes were evaluated at 16 kHz cochlear frequency and at suprathreshold levels (80 dB SPL). WT, *n* = 15 mice; CMT1A, *n* = 10 mice; CMT1E, *n* = 12 mice. One-way ANOVA followed by Tukey's multiple-comparison test was used to evaluate statistical differences among the experimental groups. ****P* < 0.001; *****P* < 0.0001. Data are shown as mean ± SEM.

decrease in signaling along the ascending auditory pathway (Figure 2, A and B), whereas ABR peaks II, III, IV, and V in CMT1A mice are normal at 4 months of age (Figure 2, A and B), suggestive of the homeostatic compensation in the auditory brainstem and midbrain seen in animals with partial peripheral deafferentation (52–56). Remarkably, after 2 months of age, the functional defects in CMT1A are identical to those we previously found in mice following transient AN demyelination — i.e., normal threshold, reduced ABR peak I amplitudes, and longer peak I latencies (28).

CMT1A and CMT1E mice have distinct inner ear axonal and myelin pathologies. To explore the structural basis for the functional phenotypes in the CMT1 mice, inner ears were harvested after the final ABR/ DPOAE recording and were subjected to several levels of histological analysis. Transmission electron microscopy of cross sections through the osseous spiral lamina (OSL, the bony structure encasing the AN fibers; Supplemental Figure 3) in the midcochlear region (~16 kHz) demonstrated that CMT1A cochleas have a small but significant reduction in myelinated axon density (18%), whereas axonal loss is much

larger in CMT1E mice (~78%) (Figure 3, A and B, and Supplemental Figures 3 and 4). Interestingly, while the diameters of the remaining axons in CMT1E cochleas are like those seen in WT, axonal diameters in CMT1A ears are reduced (Figure 3C). In contrast, measurements of g-ratio (the ratio of the inner versus outer layer diameter of the myelin sheath) reflect that myelin is thinner (larger g-ratio) in CMT1E auditory axons (Figure 3, D and E), whereas CMT1A cochleas have a more complex myelin phenotype, with small-diameter axons exhibiting thicker myelin, resulting in a steepening of the regression line in the g-ratio versus axon diameter graph (Figure 3, D and E). In addition, in CMT1A, we observed some axons with mild myelin compaction defects, but myelin abnormalities were more dramatic in CMT1E mice (Supplemental Figure 4). These results demonstrate that CMT1A and CMT1E mutations cause distinct AN axonal and myelin structural changes, which are similar to those seen in other peripheral nerves (46, 57–62), with CMT1A mice presenting a less severe phenotype.

CMT1A results in AN heminodes abnormalities. The auditory physiology phenotypes in CMT1A mice i.e., normal threshold, reduced ABR peak I amplitudes, and longer peak I latencies (Figure 1, Figure 2A, and Supplemental Figure 1) — are remarkably similar to those we previously recorded in mice that had undergone remyelination after transient ablation of Schwann cells (28). In the latter case, the only AN structural abnormality that correlates with the physiological findings is the disarray of AN heminodes, the nodal structures at the peripheral ends of the terminal Schwann cells adjacent to the IHCs. AN heminodes have been proposed to function as the SGN action potential initiation site and to play a critical role in synchronous neural transmission (48, 63–65). To evaluate the effect of CMT1A and CMT1E on nodal structures, we used laser confocal microscopy with antibodies targeted to Ankyrin-G (a nodal protein marker; ref. 66), Caspr (a paranodal marker; ref. 66), and Myelin basic protein (MBP) to visualize myelin (67) on cochleas from 4-month-old mice.

Whereas AN heminodes in WT mice are tightly clustered within the 20 μm segments of the myelinated axon closest to the IHCs, heminode clustering is disrupted in CMT1A cochleas (Figure 4, A and B), with elongated Caspr⁺ regions and Ankyrin-G⁺ regions located farther from the peripheral end of the heminodes relative to those in WT. Furthermore, similar to what we observed following transient demyelination-remyelination (28), there are numerous NoR in the area directly adjacent to heminodes in CMT1A cochleas, whereas very few NoR are present in this area in WT (Figure 4, B and D). Nevertheless, the overall NoR density in the peripheral AN processes of CMT1A mice is similar to that in WT (Figure 4, C and E), suggesting that, overall, Schwann cell internode lengths are not affected in these mutants. As can be expected from the severe AN axonal loss in CMT1E mice, the heminodal disruption in these mutants is more severe (Figure 4, A and B). Accurate quantification of heminodes and NoR density in CMT1E mice was not possible due to the profound AN axon loss.

CMT1A and CMT1E mice have different degrees of IHC-SGN synapse loss without affecting HC number. Finally, to evaluate the potential contribution of HC or IHC synapse loss to the auditory physiology defects of the mutant mice, we quantified the number of IHCs and OHCs as well as IHC-SGN synaptic density in 4-month-old mice from each cohort. Notably, IHC and OHC survival were not affected by either of the 2 CMT1 mutations (Figure 5, A and C). In concordance with the severe axonal loss (Figure 3, A and B), CMT1E mutants have a large (58%–61%) reduction in IHC-SGN synaptic density compared with WT (Figure 5, B and D). In contrast, CMT1A mice exhibit a more modest (15%) reduction in IHC-SGN synaptic density in the mid-high frequency area (Figure 5, B and D), which may also reflect the small degree of axonal loss in these mutants (Figure 3, A and B, and Supplemental Figures 3 and 4).

Discussion

The concept of HHL was first proposed in 2011 (24), and since then, research on the mechanism of its pathogenesis has centered around IHC synaptopathy (14). Here we provide evidence that a mouse model of CMT1A, the most prevalent hereditary peripheral myelin disorder, exhibits the functional hallmarks of HHL (i.e., normal auditory thresholds but reduced ABR peak I amplitudes). The observation that a mouse model of CMT1E, which is caused by different alterations in the same gene as CMT1A (*PMP22*), has overt hearing loss (like some CMT1E patients) together with a distinct set of structural deficits in the cochlea, corroborates the specificity of the CMT1A phenotype. Furthermore, the notion that myelin disorders are an alternative etiology for HHL is supported by the remarkable phenotypic similarities between mice following transient AN demyelination (28) and the CMT1A model (i.e., normal ABR and DPOAE thresholds) reduced ABR peak I amplitudes together with longer peak I latencies. This idea is also supported by

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Figure 3. CMT1A and CMT1E mice have distinct myelin and axonal pathologies. (**A**) Representative electron micrographs of sections through the OSL showing an axon bundle (top), a high-magnification image of the axons (middle), and an individual myelinated axon image (bottom) from 4-month-old WT (left), CMT1A (center), and CMT1E (right) mice. Scale bars: 4 μm (top), 1 μm (middle), 200 nm (bottom).(**B**) CMT1A and CMT1E animals have different degrees of myelinated axons loss — 19% and 78%, respectively — compared with WT mice. Dots represent the number of myelinated axons per 100 μm² cross section through the OSL at 16 kHz per mouse (*n* = 3 cochleae per genotype). (**C**) Frequency histogram showing that CMT1A mice have a larger number of small diameter axons than WT mice. Inset: CMT1A mice have a 16% reduction in axon diameter versus WT. Each dot represents the averaged axon diameter per experimental animal (*n* = 3 cochleae per genotype). (**D**) CMT1E mice have decreased myelin thickness (g-ratio = 0.735) compared with WT (g-ratio = 0.633) and CMT1A mice (g-ratio = 0.633). Each dot represents the averaged g-ratio per experimental animal (*n* = 3 cochleae per genotype) (**E**) CMT1E hypomyelination is independent of axon diameter while CMT1A mice have an increased steepening in the regression line in the g-ratio versus axon diameter graph (*n* = 3 cochleae per genotype at the 16 kHz region; WT axons = 201–253 per cochlea, CMT1A axons = 138–198 per cochlea, CMT1E axons = 62–146 per cochlea). Quantification of myelinated axon densities, axon diameters, and g-ratios were analyzed by 1-way ANOVA followed by Tukey's multiple-comparison test. Linear regression and differences among slopes of axon diameters versus g-ratio plots were obtained with GraphPad Prism simple linear regression test. **P* < 0.05; *****P* < 0.0001. Data are shown as mean ± SEM.

the fact that the longer ABR peak I latencies observed in both myelinopathy models is absent from mice with noise-induced or age-related IHC synaptopathy (14, 28).

Here we also show that, like in mice following transient AN demyelination (28), cochlear heminodes are disrupted and some NoR are mislocalized in CMT1A mice. Moreover, this structural phenotype correlates

Figure 4. CMT1A and CMT1E mice have nodes of Ranvier and heminodes abnormalities. (**A**) Representative low-magnification images showing that the AN heminodes adjacent to the organ of Corti (top) and the AN myelin in the OSL (bottom) are disrupted in CMT1A (center) and CMT1E (right) cochleae at 4 months old compared with WT mice (left). Scale bar: 70 μm. (**B** and **C**) High-magnification images showing the heminodes and nodes of Ranvier and the nodes of Ranvier at the 16 kHz cochlear region from WT (left), CMT1A (center) and CMT1E (right) mice. Scale bars: 10 μm. (**D**) CMT1A mice have increased number of nodes of Ranvier–like structures in the 20 μm area below the disrupted heminodes compared with WT (WT, *n* = 10 cochleae; CMT1A, *n* = 12 cochleae). (**E**) The density of nodes of Ranvier at the level of the OSL in CMT1A mice is not affected. Samples were immunolabeled for paranodes (Caspr), nodes of Ranvier (Ankyrin-G) and myelin (MBP). WT, *n* = 4 cochleae; CMT1A, *n* = 5 cochleae. One cochlea from each mouse was analyzed, and quantifications were done at the ~16 kHz region. Nodes of Ranvier density in either the 20 μm below the heminodal area or in the OSL region was analyzed by 2-tailed *t* test. Quantification of heminodes and nodes of Ranvier in CMT1E mice was not possible due to the severe axonal damage exhibited. ****P* < 0.001. Data are shown as mean ± SEM.

> with the reduced sound evoked SGN activation in both models, supporting the crucial role of the AN heminodes as the SGN action potential initiation site and its importance for fast and synchronized AN spike generation as has been previously suggested in computational models or electrophysiological and immunostaining studies (48, 63–65). Importantly, our mathematical model of HHL caused by myelinopathy suggest that the heminodal defects might cause AN fiber spike generation failure, which could contribute to the reduced amplitude of the first peak of the ABR waveform (48, 63–65). The finding that CMT1A results in AN heminode defects also raises the possibility that defects in other heminodes (e.g., those close to neuromuscular synapses) or other action potential initiation sites (e.g., axon initial segments in sensory neurons) might contribute to other sensory and motor phenotypes in CMT1A.

> The distinct AN myelin and axonal phenotypes in each CMT1 mouse model are also remarkably similar to those observed in other peripheral nerves, suggesting that the specific *PMP22* mutations exert common pathological effects across myelinating glia. For example, AN in CMT1A cochleas exhibit an increased proportion of smaller axons, a slight reduction in the number of myelinated fibers, hypermyelination of

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SGN synapse loss without hair cell loss. (**A** and **B**) Representative confocal images of hair cells (**A**) and IHC-SGN synapses (**B**) at the 16 kHz region of WT (top), CMT1A (middle), and CMT1E (bottom) cochleae immunolabeled for hair cells (MyoVIIa), presynaptic ribbons (Ctbp2), and postsynaptic receptor patches (GluR2). Scale bars: 30 μm (left), 10 μm (right). (**C**) Inner (left) and outer (right) hair cells survival is not affected by these 2 different CMT1 mutations. WT, *n* = 6 cochleae; CMT1A, *n* = 6 cochleae; CMT1E, *n* = 5 cochleae. One cochlea from each animal was imaged, with 3 adjacent *Z* stacks acquired at each specific cochlear region. (**D**) CMT1A mice have mild IHC-SGN synapse loss at 16 and 32 kHz areas, compared with WT mice. CMT1E IHC-SGN synapse density is more severely reduced than in CMT1A. WT, *n* = 3–7 cochleae; CMT1A, *n* = 3–7 cochleae; CMT1E, *n* = 3–7 cochleae. One cochlea from each animal was imaged, with 3 adjacent *Z* stacks acquired at each specific cochlear region. IHC-SGN synapse density was analyzed by 1-way ANOVA followed by Šidák's multiple-comparison test. Quantification of inner and outer hair cells was analyzed by Kruskal-Wallis test followed by Dunn's multiple-comparison test. **P* < 0.05; ***P* < 0.01; *****P* < 0.0001. Data are shown as mean ± SEM.

Figure 5. CMT1A and CMT1E mice have different degrees of IHC-

small-diameter fibers, and a steepening of the regression line in the g-ratio versus axon diameter graph, all features that have been reported in other peripheral nerves of this mouse model and also in CMT1A rat models (46, 57, 58, 60). In contrast, sciatic nerves from CMT1E mice exhibit hypomyelination, much like what we and others have observed in the cochlea of this model (59, 61, 62). Furthermore, the AN phenotypes in both models are consistent with findings in other nerves in patients with CMT1 (68).

Given the lack of validated diagnostic tools for HHL in humans (23), our findings in the CMT1A mouse model provide strong support for the handful of clinical observations suggesting patients with CMT1A have HHL-like phenotypes. An early study of children with CMT1A reported that most have normal audiological thresholds, but some have lower ABR amplitudes and longer latencies (36). Furthermore, more than 60% of these children have defects in auditory perception, temporal processing, and speech recognition, consequences that have now been predicted for humans with HHL (1–9). More recent studies reported that adult patients with CMT1A have normal auditory thresholds but defects in speech perception in noisy backgrounds, reflected by lower temporal and spectral resolution than healthy controls (34, 35). Importantly, our results also suggest that studying patients with genetic disorders such as CMT1A is likely to facilitate the development of better tests for HHL that could be validated using the power of phenotype-genotype relationships.

The auditory function deficits of CMT1A and CMT1E mice are profoundly different but, in each case, are similar to those seen in patients with the respective mutations (34–37, 39–43, 69). Thus, we infer that it is likely that the cochlear structural phenotypes in each patient population correlate to those we see in each mouse model as well. Specifically, our findings suggest that patients with CMT1E most likely have severe SGN neuron loss, whereas patients with CMT1A are likely to retain most SGNs but have heminode disruptions. If this is the case, it would imply that patients with CMT1E are unlikely to benefit much from hearing aids or cochlear implants, whereas patients with CMT1A might be helped by hearing augmentation. This could in part explain the different efficacy outcomes observed in some patients with CMT who have cochlear implants (70, 71).Given the growing evidence that hearing loss can contribute to cognitive decline and dementia (72–74), it would also be important that neurologists who care for patients with peripheral neuropathy include auditory testing in clinical workups and consider hearing-enhancement devices in treatment. Furthermore, our observations that a mouse model of myelinopathy has the hallmarks of HHL suggest that this might be a frequent comorbidity in other peripheral demyelinating neuropathies, e.g., Guillain-Barré syndrome. In this context, therapies that promote AN remyelination and heminode reorganization could help to prevent or slow the progression of hearing impairments associated with CMT1A or other peripheral demyelinating neuropathies (75–79).

Methods

Sex as a biological variable. Our study examined male and female mice, and no differences were found between sexes.

Animals. *B6.Cg-Tg (PMP22) C3Fbas/J* mice (PMP22-C3 or C3-PMP, here termed CMT1A; The Jackson Laboratory, stock no. 030052), *B6.D2-pmp22Tr-j/J* mice (Trembler-J, here termed CMT1E; The Jackson Laboratory, stock no. 002504), and C57BL/6J WT littermates (1 to 4 months old, male and female) were used in this study. CMT1A mice express 3 copies of WT human peripheral myelin protein 22 (PMP22) gene, mimicking CMT1A (46). CMT1E mice have a semidominant point spontaneous mutation (i.e., a T to C transition at nucleotide position 47) resulting in the substitution of a leucine by a proline in the first transmembrane domain of the PMP22 protein (47). This point mutation causes peripheral demyelination that mimics CMT1E disease.

Auditory function tests. ABRs and DPOAEs were performed on mice anesthetized with a mixture of ketamine (100 mg/kg, i.p.) and xylazine (20 mg/kg, i.p.). All groups of animals were tested at 1, 2, 3, and 4 months of age.

For ABR recordings, acoustic stimuli were delivered through a closed acoustic system, consisting of 2 sound sources (CDMG15008-03A, CUI) and an electret condenser microphone (FG-23329-PO7, Knowles) as an in-dwelling probe microphone. Three needle electrodes were placed into the skin at the dorsal-midline: 1 close to the neural crest, 1 behind the left pinna, and 1 at the base of the tail (ground). ABR potentials were evoked with 5 ms tone pips (0.5 ms rise-fall, with a \cos^2 envelope, at 40 s⁻¹) delivered to the eardrum at log-spaced frequencies from 5.6 kHz to 42.25 kHz. The response was amplified (10,000×) and filtered (0.3–3 kHz) with an analog-to-digital board in a PC-based data-acquisition system. The sound level was raised in 5 dB steps from 20 to 80 dB SPL. At each level, 1,024 responses were averaged (with stimulus polarity alternated) after "artefact rejection" above 15 μV.

DPOAEs, in response to 2 primary tones of frequency f1 and f2, were recorded at $(2 \times f1) - f2$, with $f2/f1 = 1.2$ and the f2 level 10 dB lower than the f1 level. Stimuli were raised in 5 dB steps from 20 to 80 dB. The ear canal sound pressure was amplified and digitally sampled at 4 μs intervals. DPOAE thresholds were defined as the lower SPL where $(2f1 - f2) - (2f1 - f2Nse) > 0$, where f2Nse represents the f2 base recording noise floor.

Both ABR and DPOAE recordings were performed using the EPL cochlear function test suite (Mass Eye and Ear). ABR thresholds, ABR peak I amplitudes and latencies, ABR waveforms, and DPOAE thresholds were analyzed with ABR peak Analysis software (Mass Eye and Ear) and Microsoft Excel.

Plastic sections and transmission electron microscopy. Otic capsules from 4-month-old CMT1A, CMT1E, and WT mice were dissected and fixed in 1.25% paraformaldehyde, 2.5% glutaraldehyde (EMS, 16320), and 100 mM cacodylate (EMS, 11552) overnight, followed by osmification in 1% osmium tetroxide (EMS, 19150) for 45 minutes and decalcification in 5% EDTA (Thermo Fisher Scientific, BP118-500), 1% glutaraldehyde, and 0.1M PBS (Thermo Fisher Scientific, 10010-023) for 5 days. Cochleae were microdissected, and whole mount tissues were gradually dehydrated in ethanol (70%, 95%, and 100% steps) and embedded in araldite resin. The embedded samples were degassed for 2 hours and hardened at 60°C for 5 days. For myelinated axons density counts, semi-thin cross sections of AN fibers were made at the level of the OSL through the 16 kHz cochlear region. All myelinated fibers in the OSL cross sections were counted at 60× magnification. The number of fibers was then divided by the axonal bundle extent of OSL included in that section, to arrive at an estimate of the number of myelinated fibers per $100 \mu m^2$ of organ of Corti. For g-ratio and axon diameter estimations, ultra-thin (70 nm) OSL cross-sections were prepared for transmission electron microscopy. Ultra-thin sections containing 16 kHz cochlear region were sequentially poststained with 6% w/v uranyl acetate and 4.4% w/v lead citrate. Transmission electron microscopy was performed on JEOL 1400-plus electron microscope (JEOL USA). Multiple nonoverlapping regions of the AN fiber cross-sections were imaged at 600×, 3,000×, and 10,000× magnification.

The circumference of each axon and axon + myelin sheath were measured using ImageJ software g-ratio plugin (version 1.53c, NIH) on 3,000× magnification images. The g-ratios were calculated as g-ratio $=$ (axon area)/(axon area + myelin sheath area). Axon diameters were calculated using the same software. Axons with circularity $(4 \times \pi \times \text{area/perimeter}^2)$ under 0.6 were excluded from analysis. All electron microscopy images are representative of 3 individual mice per group.

Immunostaining and confocal imaging. Inner ear tissues from 4-month-old CMT1A, CMT1E, and WT mice were dissected and fixed in 4% paraformaldehyde (Alfa Aesar, A11313.36) in 0.01M phosphate-buffered saline (PBS) for 2 hours at room temperature, followed by decalcification in 5% EDTA at 4°C for 5 days. Then, cochlear tissues were microdissected and permeabilized by freeze-thawing in 30% sucrose in PBS. The microdissected tissues were incubated in blocking buffer containing 5% normal horse serum and 0.3% Triton X-100 in PBS for 1 hour. Tissues were then incubated in primary antibodies (diluted in 1% normal horse serum and 0.3% Triton X-100 in PBS) at 37°C overnight. The primary antibodies used in this study were as follows. For NoR, heminodes and myelin staining, anti-Caspr (NeuroMab, clone K65/35; 1:1,000), anti–Ankyrin G (gift from Paul Jenkins laboratory, Department of Pharmacology, University of Michigan; 1:500), anti-MBP (MilliporeSigma, MAB386; 1:1,000) were used; for HCs and IHC-SGN synapses, anti-Ctbp2 (BD Biosciences, 612044; 1:200), anti-GluR2 (MilliporeSigma, MAB397; 1:1,000), and anti-MyoVIIa (Proteus Biosciences, 25-6790; 1:100) were used. Tissues were then incubated with appropriate Alexa Fluor–conjugated fluorescent secondary antibodies (Invitrogen; 1:1,000 diluted in 1% normal horse serum and 0.3% Triton X-100 in PBS; for NoR, heminodes and myelin, AF488 IgG1 [A-21121], AF568 [A11036], and AF647 [A-21247]; for HCs and IHC-SGN synapses, AF488 IgG2a [A-21131]; AF568 IgG1 [A-21124]; and AF647 IgG [A-21244]) for 1 hour at room temperature. The tissues were mounted on microscope slides in ProLong Diamond Antifade Mountant (Thermo Fisher Scientific). All pieces of each cochlea were imaged at low power (10× magnification) to convert cochlear locations into frequency (tonotopic mapping) using a custom plug-in to ImageJ (1.53c NIH) available at the website of the Eaton-Peabody Laboratories (EPL). Cochlear tissues from the \sim 16 kHz region were used for further analyses unless stated otherwise. Confocal *Z* stacks of cochlear tissues were taken using a Leica SP8 confocal microscope.

NoR and heminodal structure *Z* stacks (0.3 μm step size) were taken under 63× (+4.61× optical zoom) or 40× magnification (for low-magnification images) at the ~16 kHz cochlear region (1 *Z* stack per individual animal). ImageJ/Fiji software (version 1.53c, NIH) was used for *Z* stack processing and quantification of NoR presence either in the 20 μm below the heminodal zone or in the OSL lamina region. For NoR quantification, the number of NoR in each sample was counted manually using an ImageJ/Fiji software multipoint counter tool. For figures, 1 representative image was selected from among the 12–48 images.

Images for HC counts were taken under 40× magnification. For IHC synapse counts, *Z* stacks (0.3 μm step size) were taken under $63 \times (+2.4 \times$ optical zoom) magnification spanning the entire IHC height to ensure all synapses were imaged. Imaging and analyses of cochlear HCs and synapses were performed as previously described in ref. 80. Briefly, ImageJ/Fiji software (version 1.53c, NIH) was used for image processing and quantification. One cochlea from each animal was imaged for each experiment, with 3 adjacent *Z* stacks acquired at each specific cochlear region. For HC quantification, the number of IHCs and OHCs at specific cochlear regions in each animal was determined based on the MyoVIIa channel and counted manually using ImageJ/Fiji software multipoint counter tool. For synapse counts, CtBP2 and GluR2 puncta in each image stack were also captured and counted manually using ImageJ/Fiji software multipoint counter tool. Synaptic counts of each *Z* stack were divided by the number of IHCs, which could be visualized by staining of MyoVIIa antibody. Each individual image usually contained 8–10 IHCs. For figures, 1 representative image was selected from among the 9–24 images from the specific frequency shown.

Statistics. Graphics and statistical tests were performed using GraphPad Prism version 9.3.1 for Windows (GraphPad Software, www.graphpad.com). Data sets with normal distributions were analyzed with parametric tests whereas nonparametric tests were used for sets that did not conform to normality criteria. We used 2-way ANOVA, followed by Tukey's multiple-comparison test to compare either DPOAE thresholds, ABR thresholds, ABR peak I amplitudes, or latencies at each time point (1, 2, 3, and 4 months) among CMT1A, CMT1E, and WT mice (Figure 1, C–F, and Supplemental Figures 1 and 2). One-way ANOVA followed by Tukey's multiple-comparison test was used to evaluate statistical differences in ABR peak II, III, IV, and V amplitudes (Figure 2A). Quantification of myelinated axon densities, quantification of axon diameters, and g-ratios were analyzed by 1-way ANOVA followed by Tukey's multiple-comparison test (Figure 3, B–D). Simple linear regression of axon diameters versus g-ratios and statistical differences among slopes were obtained with GraphPad Prism (Figure 3E). Quantification of NoR density in either the 20 μm below the heminodal area or in the OSL region were analyzed by 2-tailed *t* test (Figure 4, D and E). Quantification of inner and OHCs was analyzed by Kruskal-Wallis test followed by Dunn's multiple-comparison test (Figure 5C). Quantification of IHC-SGNs synapse density was analyzed by 1-way ANOVA followed by Šidák's multiple-comparison test (Figure 5D). *P* < 0.05 was considered as statistically significant.

Study approval. All animal procedures were approved by the IACUC of the University of Michigan, and all experiments were performed in accordance with guidelines and regulations suggested by the *Guide for the Care and Use of Laboratory Animals* (National Academies Press, 2011).

Data availability. All data generated or analyzed during this study are included in the manuscript and in the Supporting Data Values file. Original electron microscopy and confocal images used for calculations, ABR, and DPOAE recordings will be uploaded to Dryad.

Author contributions

LRC and GC designed the research studies. LRC, LJ, MCY, ASD, and NDC conducted experiments. LRC, MCY, ASD, and NDC acquired data. LRC, MCY, ASD, NDC, ZAA, and GC analyzed data. LRC, LJ, MCY, ASD, NDC, and GC wrote the manuscript.

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