LETTER TO THE EDITOR



Lypd1-DTR/+: A New Mouse Model for Specifically Damaging the Type Ic Spiral Ganglion Neurons of the Cochlea

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Dear Editor,

Inner ear auditory neurons, which are also referred as to spiral ganglion neurons (SGNs), bridge sound-receptor hair cells (HCs) to the cochlear nucleus in the brainstem [1]. Two types of SGN exist: type I SGNs (95%), which innervate inner HCs (IHCs), and type II SGNs (only 5%), which form synapses with outer HCs (OHCs) [1, 2]. Recent transcriptomic studies have revealed the heterogeneity within type I SGNs, which can be further divided into three distinct subtypes: Ia, Ib, and Ic [3-5]; these three subtypes first emerge at E16 and are refined over the first postnatal week [6, 7]. Although no molecular markers have thus far been found to show complete "ON" or "OFF" expression patterns in the subtypes, several genes display marked expression gradients among Ia, Ib, and Ic [3–5]. Ly6/Plaur domain containing 1, which is encoded by Lypd1 (also known as Lynx2), belongs to the Ly-6/neurotoxin superfamily and is

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involved in modulating the activity of nicotinic acetylcholine receptors [8]. Notably, *Lypd1* is expressed at the highest level in type Ic SGNs and minimally expressed in Ib and Ia SGNs in adult cochleae [3–5]. Moreover, two intrinsic transcription factors, Neurod1 and Runx1, promote the entry of undifferentiated SGN precursors into the Ic or Ic/Ib differentiation track [6, 9]. Lastly, IHC-driven activity through synapses between IHCs and SGNs is also required in the diversification of type I SGNs [3, 4]. Vesicular glutamate transporter type 3 (vGlut3, encoded by *Slc17a8*) is essential for glutamate release from IHCs [10, 11], and, accordingly, *vGlut3^{-/-}* mice show drastically reduced percentages of type Ic SGNs and overproduction of type Ia SGNs expressing high levels of Calb2 [3, 4].

Type Ic SGNs present a few notable features: First, they have a low spontaneous firing rate [3]; second, they form synapses with larger presynaptic ribbons at the modiolar side of IHCs [3–5]; and third, they are selectively vulnerable to aging [3]. Thus, degeneration of type Ic SGNs in mice is expected to closely mimic human age-related hearing loss (ARHL), or presbycusis, the most common form of sensory impairment among older adults [12]. However, a mouse model with specific damage of type Ic SGNs is not yet available.

Here, we generated a new mouse model: $Lypd1*3\times$ V5-P2A-DTR/+ (Lypd1-DTR/+, in short); we applied the diphtheria toxin (DT)/DT receptor (DTR) system, which is widely used to achieve cell-type-specific damage in the inner ear [13]. By using CRISPR/Cas9-mediated homologous recombination in mouse zygotes [14], we inserted a DNA fragment containing 3×V5-P2A-DTR immediately before the stop codon (TGA) of *Lypd1* (Fig. S1A–C). Thus, DTR would be expressed, in principle, under the control of endogenous *Lypd1* promoters and enhancers. Moreover, because three V5 tags are fused to the C-terminus of Lypd1 here, a V5 antibody can be used to visualize Lypd1 protein expression. Tail-DNA PCR analysis readily distinguished wild-type (WT), heterozygous, and homozygous knockin (KI) littermates (Fig. S1D). Southern blotting using a DTR probe (blue line in Fig. S1C) further revealed that, besides being correctly targeted into the *Lypd1* locus as designed, the targeting vector was randomly inserted in one unknown location in the mouse genome. Because random insertion frequently occurs in inactive regions of the genome, we suspected that *Lypd1*-DTR/+ could still be used to specifically damage Lypd1⁺ type Ic SGNs. Alternatively, *Lypd1*-DTR/+ could at least serve as a pseudo-transgenic mouse line, as confirmed by our assays below.

We first assessed the degree to which V5 expression recapitulated the previously reported expression pattern of endogenous Lypd1 [3–5]. Triple staining of V5, the pan-SGN marker Map2, and the glial-cell marker Sox2 revealed that V5⁺/Map2⁺ SGNs accounted for $26.8 \pm 1.8\%$ (n = 3) of the total Map2⁺ SGNs in *Lypd1*-DTR/+ mice at P28, which matched the 24–25% reported in the adult WT cochlea. To

further show that V5 expression can faithfully reflect Lypd1 expression, we compared the proportion of $V5^+/Map2^+$ SGNs between Lypd1-DTR/+ (Fig. 1A-A''') and Lypd1-DTR/+; vGlut3^{-/-} mice at P28 (Fig. 1B-B'''). As expected, the percentage of V5⁺/Map2⁺ SGNs dropped significantly (P < 0.001), from 26.8 \pm 1.8% (n = 3) in Lypd1-DTR/+ to $6.6 \pm 1.5\%$ (*n* = 3) in *Lypd1*-DTR/+; *vGlut3*^{-/-} (Fig. 1C). Conversely, triple staining of Map2, Calb2 (a marker enriched in type Ia SGNs), and Sox2 showed that the percentage of Calb2⁺/Map2⁺ SGNs increased from $65.4 \pm 4.9\%$ (n = 3) in Lypd1-DTR/+ to 86.1 ± 2.4% (n = 3) in Lypd1-DTR/+; vGlut3^{-/-} at P28 (Fig.1D–F). Notably, V5 (Lypd1) was not detected in any Sox2⁺ glial cell. Thus, our results agreed with the previous finding that the percentage of type Ic SGNs is drastically lower in *vGlut3^{-/-}* mice than in control mice [3, 4].

Dual labeling for V5 and Calb2 further revealed a minimal overlap of their expression in SGNs at P28 (Fig. S2); Calb2 is enriched in type Ia and Ib SGNs but is minimally expressed in type Ic SGNs [3–5]. Most V5+ SGNs did not



Fig. 1 Visualization of type Ic SGNs by using V5 expression in the *Lypd1*-DTR/+ model. **A–B**^{***} Triple staining of Map2, V5, and Sox2 in *Lypd1*-DTR/+ (**A–A**^{***}) and *Lypd1*-DTR/+; $vGlut3^{-/-}$ (**B–B**^{***}) mice. All V5⁺ cells express the pan-SGN marker Map2. Yellow arrows: V5⁺/Map2⁺ SGNs. V5 expression was not detected in any Sox2⁺ glial cells. **C** Percentages of V5⁺/Map2⁺ SGNs (type Ic) among all SGNs: *Lypd1*-DTR/+ (red) *vs Lypd1*-DTR/+; *vGlut3^{-/-}* (black) mice at P28. ****P* <0.001. **D–E**^{***} Triple staining of Map2,

Calb2, and Sox2 in *Lypd1*-DTR/+ (**D**-**D**^{**}) and *Lypd1*-DTR/+; *vGlut3*^{-/-} (**E**-**E**^{**}) mice at P28. Calb2 (Calretinin) is expressed in a fraction of Map2⁺ SGNs, but not in any Sox2⁺ glial cells. Yellow arrows: Calb2⁺/Map2⁺ SGNs. **F** Percentages of Calb2⁺/Map2⁺ SGNs among all SGNs: *Lypd1*-DTR/+ (red) *vs Lypd1*-DTR/+; *vGlut3*^{-/-} (black) mice at P28. n = 3 for each model, **P* <0.05. Scale bars, 20 µm.

express Calb2 (white arrows in Fig. S2A–A''), and only 5.3 \pm 2.5% of the V5⁺ SGNs were Calb2⁺ (yellow arrows in Fig. S2B–B''); moreover, Calb2 expression in these V5⁺ SGNs was lower than that in other nearby Calb2⁺ SGNs that did not express Lypd1 (V5) (Fig. S2B–B''). Similarly, among the Calb2⁺ SGNs, only 1.8 \pm 0.9% of the cells were V5⁺. Collectively, these results showed that despite the random vector insertion in the *Lypd1*-DTR/+ mouse genome, the V5 expression pattern resembled the previously reported *Lypd1* mRNA expression pattern in adult SGNs [3–5]. Thus, we reason that DTR is primarily expressed in type Ic SGNs and that *Lypd1*-DTR/+ is a suitable model for ablating Lypd1⁺ Ic SGNs through DT administration at adult ages.

Next, we characterized the SGN cell-death pattern in *Lypd1*-DTR/+ mice treated with or without a single dose of DT (25 ng/g body weight) at P31 and then analyzed at P38. Compared with *Lypd1*-DTR/+ mice not treated with DT (control, "no DT") (Fig. 2A–A"'), DT-treated *Lypd1*-DTR/+ mice showed the loss of a majority of V5⁺ SGNs (Fig. 2B–B"'); in the control mice, the percentage of V5⁺ (Lypd1⁺) type Ic SGNs was $25.2 \pm 0.3\%$ (n = 3), but this

dropped markedly, to $0.8 \pm 0.3\%$ (n = 3), in DT-treated mice (Fig. 2C). In accordance with this specific loss of Ic SGNs, the total number of Map2⁺/Sox2⁻ SGNs in the DT-treated mice (290.8 \pm 7.4, n = 3) was also significantly lower than that (341.2 \pm 4.4, n = 3) in no-DT-treated mice (Fig. 2D).

Lastly, we quantified the hearing ability of *Lypd1*-DTR/+ mice by measuring the auditory brainstem response (ABR): the hearing thresholds at all frequencies in DT-treated mice (red line in Fig. 2E) were significantly higher than their counterparts in no DT-treated *Lypd1*-DTR/+ mice (green line in Fig. 2E). Thus, the hearing ability was substantially impaired in DT-treated mice. Furthermore, analysis of WT mice treated with an identical single dose of DT revealed no significant difference in ABR threshold/hearing function between the DT-treated WT (blue line in Fig. 2E) and no DT-treated *Lypd1*-DTR/+ mice (green line in Fig. 2E), eliminating the possibility that DT treatment itself affects the hearing of mice.

Two lines of evidence support the view that LypdI-DTR/+ is a reliable model for visualizing type Ic SGNs. First, we found here that the percentage of V5⁺/Map2⁺



Fig. 2 The V5⁺ (Lypd1⁺) SGNs are specifically damaged in adult *Lypd1*-DTR/+ mice after DT treatment. **A–B**^{***} Triple staining of Map2, V5, and Sox2 in *Lypd1*-DTR/+ mice without (**A–A**^{***}) and with single-dose DT treatment at P31 (**B–B**^{***}). All samples were analyzed at P38. Arrows in (**A–A**^{***}): one V5⁺/Map2⁺ type Ic SGN. Almost all V5⁺ SGNs disappeared in DT-treated mice (**B–B**^{***}). C Percentages of V5⁺ SGNs in no-DT-treated (red) and DT-treated

(black) *Lypd1*-DTR/+ mice at P38. ****P <0.0001. **D** Total numbers of Map2⁺ SGNs in no-DT-treated (red) and DT-treated (black) *Lypd1*-DTR/+ mice at P38. **P <0.01. **E** ABR measurements from wild-type mice with DT treatment (blue line) and *Lypd1*-DTR/+ mice without (green line) or with DT treatment (red line). n = 3 for each model, **P <0.001, ***P <0.001, ****P <0.0001. Scale bar, 20 µm.

SGNs was 25.2–26.8%, which is close to the ~24% reported in three previous studies [3–5]. Second, when V5 expression was used as the readout, the percentage of type Ic SGNs was measured to be drastically decreased in $vGlut3^{-/-}$ mutants as reported before [3, 4]. Thus, the V5 expression faithfully recapitulated the expression pattern of endogenous Lypd1. More importantly, the near absence of V5⁺ SGNs in *Lypd1*-DTR/+ mice after treatment with a single dose of DT confirmed that type Ic SGNs were selectively ablated, implicating *Lypd1*-DTR/+ as a useful model in which to mimic human ARHL.

Considerable research effort has been devoted to converting cochlear glial cells into SGNs [15–17] because this represents a promising approach for regenerating SGNs in situ. We previously showed that Neurod1 and Neurog1 expression can convert glial cells into nascent SGNs, but adult glial cells are not responsive to ectopic Neurod1 and Neurog1 expression [16]. Whether pre-damaging endogenous SGNs would increase the reprogramming efficiency of Neurod1 and Neurog1 remains unknown. Ouabain has been used to damage type I SGNs [18], but ouabain does not show selectivity for the type I SGN subtype. To the best of our knowledge, Lypd1-DTR/+ is the first genetic model that allows selective damage of type Ic SGNs. It will be of interest to determine whether glial cells tend to transdifferentiate into type Ic SGNs only if endogenous type Ic SGNs are specifically damaged. In summary, we have successfully established the Lypd1-DTR/+ mouse model that could be used to visualize and ablate type Ic SGNs.

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