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VGLUT3-p.A211V variant fuses stereocilia bundles and elongates synaptic ribbons

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Abstract DFNA25 is an autosomal-dominant and progressive form of human deafness caused by mutations in the SLC17A8 gene, which encodes the vesicular glutamate transporter type 3

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(VGLUT3). To resolve the mechanisms underlying DFNA25, we studied phenotypes of mice harbouring the p.A221V mutation in humans (corresponding to p.A224V in mice). Using auditory brainstem response and distortion product otoacoustic emissions, we showed progressive hearing loss with intact cochlear amplification in the VGLUT3A224V/A224V mouse. The summing potential was reduced, indicating the alteration of inner hair cell (IHC) receptor potential. Scanning electron microscopy examinations demonstrated the collapse of stereocilia bundles in IHCs, leaving those from outer hair cells unaffected. In addition, IHC ribbon synapses underwent structural and functional modifications at later stages. Using super-resolution microscopy, we observed oversized synaptic ribbons and patch-clamp membrane capacitance measurements showed an increase in the rate of the sustained releasable pool exocytosis. These results suggest that DFNA25 stems from a failure in the mechano-transduction followed by a change in synaptic transfer. The VGLUT3A224V/A224V mouse model opens the way to a deeper understanding and to a potential treatment for DFNA25.

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Abstract figure legend: Human deafness DFNA25 is caused by a point mutation in the SLC17A8 gene encoding the vesicular glutamate transporter type 3 (VGLUT3). In the mouse harbouring the point mutation found in humans, the inner hair cells are unable to transduce the acoustic stimulation into neural messages.

Key points
- The vesicular glutamate transporter type 3 (VGLUT3) loads glutamate into the synaptic vesicles of auditory sensory cells, the inner hair cells (IHCs).
- The VGLUT3-p.A211V variant is associated with human deafness DFNA25.
- Mutant mice carrying the VGLUT3-p.A211V variant show progressive hearing loss.
- IHCs from mutant mice harbour distorted stereociliary bundles, which detect incoming sound stimulation, followed by oversized synaptic ribbons, which release glutamate onto the afferent nerve fibres.
- These results suggest that DFNA25 stems from the failure of auditory sensory cells to faithfully transduce acoustic cues into neural messages.

Introduction

Mutations in the SLC17A8 gene are associated with DFNA25, an autosomal-dominant and progressive form of human deafness (Ruel et al. 2008; Ryu et al. 2016). Vesicular glutamate transporters (VGLUTs) are responsible for glutamate loading into synaptic vesicles, a critical step to achieve synaptic transfer (El Mestikawy et al. 2011). SLC17A8 encodes the vesicular glutamate transporter type 3 (VGLUT3; Fremeau et al. 2002; Gras et al. 2002; Takamori et al. 2002). VGLUT3 is expressed in small subsets of neurons in the central nervous system and in the inner ear (Ruel et al. 2008; Seal et al. 2008; El Mestikawy et al. 2011; Zhang et al. 2011). In zebrafish, VGLUT3 is expressed in the ear and lateral line organ, especially in hair cells, which convert mechanical stimuli into glutamate release onto afferent fibres (Obholzer et al. 2008). While the loss of VGLUT3 does not alter the microphonic receptor, reflecting the mechano-transducer current at the hair cell stereociliary bundle, it results in the absence of activity in the posterior lateral line ganglion. Accordingly, the VGLUT3 deletion abolishes the vestibulo-ocular and acoustic startle reflexes (Obholzer et al. 2008). In mice, the inner hair cells (IHCs), which transduce sound stimulation into glutamate secretion, express VGLUT3 (Ruel et al. 2008; Seal et al. 2008). In addition, it was recently established that outer hair cells (OHCs), which amplify acoustic input within the cochlea, also express VGLUT3 (Weisz et al. 2021). The genetic ablation of SLC17A8 leads to the calcium-triggered exocytosis of empty synaptic vesicles, making the auditory afferent fibres silent (Ruel et al. 2008; Seal et al. 2008; Akil et al. 2012; Kim et al. 2019). Thus, the loss of VGLUT3 in mice results in the lack of auditory brainstem responses, reflecting a failure in the ascending auditory pathway activation. In VGLUT3-null mice, distortion product otoacoustic emissions (DPOAEs) that are a signature of the activity in OHCs are unaffected.
In addition, it has been reported that ribbon bodies, the organelles surrounded by a monolayer of synaptic vesicles, are abnormally thin and elongated in the absence of VGLUT3 (Kim et al. 2019). However, the complete deafness in VGLUT3− mice conflicts with the variable onset and progression rate of DFNA25.

The rare variant VGLUT3-p.A211V was identified as a cause of DFNA25 (Ruel et al. 2008). This variation does not impair the glutamate vesicular accumulation, neither the quantal release nor the quantal content in hippocampal autaptic neurons (Ramet et al. 2017). In addition, investigation of a mouse line harbouring the p.A221V allele in humans (p.A224V in mouse, VGLUT3A224V/A224V mice) showed that this variant leads to a strong decrease of VGLUT3 expression (≈-70%) in terminals from the central nervous system neurons without altering VGLUT3-piloted behaviour (Ramet et al. 2017). Thus, mechanisms depending on the VGLUT3-p.A211V variant and underlying DFN25 auditory deficits in humans are not yet clearly understood. To tackle this point, we herein investigated the auditory system of VGLUT3A224V/A224V mice, the architecture of the IHC stereocilary bundle in terminals from the central nervous system neurons, without altering VGLUT3-piloted behaviour (Ramet et al. 2017). As previously described (Ramet et al. 2017). A point mutation was introduced in exon 5 of the mouse SLC17A8 gene: a GCG (coding for an alanine) was exchanged for a GTG (coding for a valine). Mice were bred in-house and maintained on a C57Bl6/J genetic background. Heterozygote mice were bred to generate VGLUT3A224V/A224V and wild-type littermates.

Genotyping

Genotypes of mice were determined by polymerase chain reaction (PCR) analysis of genomic DNA using FastStart PCR Master Mix (Roche Applied Science, Penzberg, Germany) as previously described (Ramet et al. 2017). In brief, tail or toe biopsies were digested overnight at 55°C in 300 μl of lysis buffer (containing in mM: 100 Tris HCl pH 8.5; 5 EDTA; 0.2% SDS; 200 NaCl; pH: 8.5) with 100 μg/ml of proteinase K (Promega, Madison, WI, USA). Samples were centrifuged for 5 min at 12,000 g and supernatants were collected. DNA was precipitated by addition of 500 μl of isopropanol. Samples were centrifuged for 10 min at 12,000 g and supernatants were discarded. DNA was washed with 500 μl of EtOH 70% and centrifuged for 5 min at 12,000 g. After evaporation of the EtOH, DNA was suspended in 100 μl of water. The PCR was conducted with the following thermal cycle program: one cycle of 94°C for 8 min; 40 cycles of 94°C for 40 s, 56°C for 30 s, 72°C for 40 s; final elongation step at 72°C for 7 min. The PCR primers were the following: p1, 5′-GGGAGGGAAAGCCAGGAAAAGG-3′; and p2, 5′-GGACACGCTCAGTAGCTGTGACCCAG-3′ for the wild-type and the mutated allele, yielding bands of 219 and 306 bp, respectively, which were visualized on a 2% agarose gels.

Ethical approval

Experiments were carried out in accordance with the animal welfare guidelines 2010/63/EC of the European Council Directive regarding the care and use of animals for experimental procedures and comply with the principles and regulation described in the Editorial by Grundy (2015). Animals were housed in facilities accredited by the French ‘Ministère de l’Agriculture et de la Forêt’ (Agreement C-34-172-36) and the experimental protocol was approved (Authorization CEEA-LR-12111) by the Animal Ethics Committee of Languedoc-Roussillon (France).

In vivo recordings

Mice were anaesthetized by an intraperitoneal injection of a mixture of Zoletil 50 (40 mg/kg) and Rompun 2% (3 mg/kg). The rectal temperature was measured with a thermistor probe and maintained at 37.1°C ± 1°C, using a heated underblanket (Homeothermic Blanket Systems, Harvard Apparatus). Heart rate was monitored via EKG electrodes. Within 20 min of the end of the physiological session, animals were sacrificed by cervical dislocation while still under deep anaesthesia.
Auditory brainstem response and distortion product otoacoustic emission recordings

For auditory brainstem response (ABR), the acoustic stimuli consisted of 10 ms tone bursts, with a 8 ms plateau and 1 ms rise/fall time, delivered at a rate of 20.4/s with alternate polarity by a JBL 2426H loudspeaker in a calibrated free field. Stimuli were presented by varying intensities from 80 to 0 dB SPL, in 10 dB steps. Stimuli were generated and data acquired using MATLAB (MathWorks, Natick, MA, USA) and LabVIEW (National Instruments, Austin, TX, USA) software. The potential difference between vertex and mastoid intra-dermal needles was amplified (5000 times, VIP-20 amplifier), sampled (at a rate of 50 kHz), filtered (bandwidth of 0.3–3 kHz) and averaged (600 times). Data were displayed using LabVIEW software and stored on a computer (Dell T7400). ABR thresholds were defined as the lowest sound intensity, which elicits a clearly distinguishable response. For DPOAE recordings, an ER-10C S/N 2528 probe (Etymotic Research), consisting of two emitters and one microphone, was inserted in the left external auditory canal. Stimuli were two equilevel (65 dB SPL) primary tones of frequency f1 and f2 with a constant f2/f1 ratio of 1.2. The distortion 2f1-f2 was extracted from the ear canal sound pressure and processed by the HearID auditory diagnostic system (Mimosa Acoustic) on a computer (Hewlett Packard). The probe was self-calibrated for the two stimulating tones before each recording. f1 and f2 were presented simultaneously, sweeping f2 from 20 to 2 kHz in quarter octave steps. For each frequency, the distortion product 2f1-f2 and the neighbouring noise amplitude levels were measured and expressed as a function of f2.

Electrocochleography

A retroauricular incision of the skin was performed on anaesthetized mice and the left tympanic bulla was opened. Cochlear potentials were recorded with a silver positive electrode placed on the round window membrane. The acoustic stimuli were identical to those used to elicit ABRs except for the alternate polarity. Gross cochlear potentials were amplified (2500 times, VIP-20 amplifier), sampled (at a rate of 50 kHz), filtered (bandwidth of 0.001–20 kHz), averaged (50–300 times), displayed with LabView software and stored on a computer (Dell T7400). The signal was then digitally filtered using MATLAB software with a low-pass filter (cut-off frequency 3.5 kHz) to measure the compound action potential (CAP) and the summating potential (SP).

Patch-clamp recordings

After cervical dislocation of mice (1–2 months and 6 months of age), IHCs of the apical coil of freshly dissected organs of Corti were patch-clamped at their baso-lateral face at room temperature in tight perforated-patch configurations. The dissection solution contained the following in mM: 5.36 KCl, 141.7 NaCl, 1 MgCl2-6H2O, 0.5 MgSO4•7H2O, 10 Hepes and 10 d-glucose. For Ca2+ current and capacitance measurement recordings, the extracellular solution contained the following in mM: 2.8 KCl, 105 NaCl, 1 MgCl2-6H2O, 2 CaCl2, 10 Hepes, 35 TEA-Cl, 1 CsCl, 10 d-glucose. The pipette solution for perforated patch-clamp recordings of Ca2+ currents contained the following in mM: 135 Cs-glutamate acid, 10 TEA-Cl, 10 4-AP, 1 MgCl2-6H2O, 10 Hepes and 400 μg/ml amphotericin B. Solutions were adjusted to pH 7.3 and had osmolarities between 290 and 310 mOsm/kg H2O. All chemicals were obtained from Sigma (St. Louis, MO, USA) with the exception of amphotericin B (Calbiochem, La Jolla, USA). Patch pipettes were pulled from borosilicate glass capillaries ( KWik Fil, WPI, Worcester, MA, USA) with a two-step vertical puller PIP 6 (HEKA Elektronik, Lambrecht, Germany) and coated with silicone elastomer (Sylgard).

Ca2+ current recordings

Currents were low-pass filtered at 5 kHz and sampled at 10 kHz for exocytic cell membrane capacitance change (∆Cm) and at 40 kHz for Ca2+ current recordings. Ca2+ current was isolated using P/p protocols (10 leak pulses with amplitudes of 20% of the original pulse from a holding potential of -117 mV). Cells that displayed a membrane current exceeding -50 pA at -87 mV were discarded from the analysis. No Rs compensation was applied. All voltages were corrected for liquid junction potentials calculated between pipette and bath (-17 mV). Mean resting capacitance, series and membrane resistances are indicated in Table 1.

Capacitance measurement recordings

Cell membrane capacitance (Cm) was measured using the Lindau–Neher technique (Lindau & Neher, 1988), implemented in the software-lockin module of Patchmaster (HEKA Elektronik) combined with compensation of pipette and resting cell capacitances by the EPC-10 (HEKA Elektronik) compensation circuits. A 1 kHz, a 70 mV peak-to-peak sinusoid was applied about the holding potential of -87 mV. Given the sinewave amplitude (70 mV) and the holding potential of the hair cells (Vm=-87 mV), the voltage command may reach...
the calcium current activation threshold at the most positive excursion of the sinewave. This might lead to errors in the membrane capacitance estimates, as the membrane resistance would change. However, the 1 kHz frequency of the sinewave should minimize the calcium current contamination, as the slow kinetics of the calcium channels would prevent it from following the 1 kHz cycle-to-cycle voltage command. $\Delta C_{m}$ was estimated as the difference of the mean $C_{m}$ over 400 ms after the end of the depolarization (the initial 250 ms was skipped) and the mean prepulse capacitance (400 ms). Mean $\Delta C_{m}$ estimates present grand averages calculated from the mean estimates of individual IHCs.

**Immunohistochemistry, confocal and super-resolution microscopy**

Immunohistochemistry was performed on whole-mount preparations of organs of Corti. Mice were decapitated after rapid cervical dislocation and their cochleas were removed from the temporal bone and dissected in phosphate buffered saline (PBS, containing in mM: 130 NaCl, 2.68 KCl, 10 Na$_2$HPO$_4$, 1.47 KH$_2$PO$_4$, pH 7.4) solution. The mid-apical cochlear turns were then fixed for 15 min in 4% paraformaldehyde diluted in PBS; afterwards, they were immuno-histochemically processed as a whole-mount. The tissues were rinsed three times for 10 min in PBS, then preincubated for 1 h in blocking solution (10% goat serum, 0.3% Triton), then incubated overnight at 4°C in the incubation solution (1% goat serum, 0.1% Triton) with primary antibodies or antisera. The primary antibodies used, and their respective dilutions were: CtBP2 1:500 (BD Transduction Laboratories, San Jose, CA, USA; Cat No.:612044), Homer1 1:300 (Merck Millipore, Cat. No. ABN37) and VGLUT3 1:300 (Synaptic Systems, Cat.No. 135 204). Tissues were then rinsed three times for 10 min in wash buffer (containing in mM: 15.8 Na$_2$HPO$_4$, 2.9 NaH$_2$PO$_4$, 0.1% Triton X-100, 450 NaCl) and incubated for 2 h in the incubation solution with fluorescently labelled secondary antibodies, rhodamin-phalloidin (Molecular probes, Eugene, OR, USA) and Hoechst dye (Invitrogen, Carlsbad, CA, USA). They were finally rinsed four times for 10 min in wash buffer and mounted in Dako fluorescence mounting medium (code-s3023, Agilent Technologies, Inc. Santa Clara, CA, USA). Tissues were examined with the Zeiss LSM880 (Zeiss, Oberkochen, Germany) airyscan or confocal microscopes of the Montpellier Resource Imaging facility (Montpellier, France). Image stacks were then processed with ImageJ software (Wayne Rasband, National Institutes of Health, USA). The quantifications of ribbons and synapses, that is, juxtaposed spots of the pre-synaptic ribbon component RIBEYE and post-synaptic density protein Homer, were performed in Z-stack confocal images using a 3D custom algorithm (Bourien et al. 2014). Mean ribbon and synapse estimates present grand averages calculated from the mean estimates of individual cochleas.

For stimulated emission depletion (STED) microscopy imaging, secondary goat anti-guinea pig Alexa-488 1:1000 (Cat.-Nr. A11039; Thermo Fisher Scientific), goat anti-mouse IgG, Abberior STAR RED 1:200 (STRED-1001-500UG, Abberior GmbH) and goat anti-rabbit IgG, Abberior STAR-580 1:200 (ST580-1002-500UG, Abberior GmbH) antibodies were applied for 2 h at room temperature. After mounting the specimen in Dako mounting medium, image acquisition was performed using an Abberior Instruments Expert Line STED microscope (Abberior Instruments GmbH; based on an Olympus IX83 inverted microscope) in confocal and/or STED mode using a 1.4 NA UPlanSApo 100x oil immersion objective lens. We employed two pulsed lasers 580 nm (red) and 647 nm (far-red) for excitation and a pulsed 775 nm laser for STED. Image stacks were acquired with Impositor Software, keeping xy pixel sizes of 30 $\times$ 30 nm and step sizes of 50 nm (3D-STED). xy pixel sizes in 2D-STED were kept at 30 $\times$ 30 nm. The microscope is housed at the Montpellier Resource Imaging (MRI) facility (Montpellier, France). Image stacks were then processed using ImageJ (Fiji) software (Wayne Rasband, National Institutes of Health, USA) and ribbon size analysis was achieved by using custom written script in the MATLAB software over

<table>
<thead>
<tr>
<th>Genotype</th>
<th>VGLUT3$^{+/+}$</th>
<th>VGLUT3$^{A224V/A224V}$</th>
<th>VGLUT3$^{+/+}$</th>
<th>VGLUT3$^{A224V/A224V}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cm (pF)</td>
<td>9.5 ± 0.7</td>
<td>8.9 ± 0.3</td>
<td>7.3 ± 0.4</td>
<td>6 ± 0.7</td>
</tr>
<tr>
<td>Rm (GΩ)</td>
<td>1.5 ± 0.1</td>
<td>1.6 ± 0.1</td>
<td>1.7 ± 0.3</td>
<td>1.6 ± 0.3</td>
</tr>
<tr>
<td>Rs (MΩ)</td>
<td>29.3 ± 1.5</td>
<td>28.7 ± 1.1</td>
<td>32.3 ± 2.1</td>
<td>33.9 ± 2</td>
</tr>
</tbody>
</table>

Numbers of cells recorded: 19 and 21 IHCs from VGLUT3$^{+/+}$ and VGLUT3$^{A224V/A224V}$ mice at 1–2 months, respectively; 14 IHCs and 9 IHCs from VGLUT3$^{+/+}$ and VGLUT3$^{A224V/A224V}$ mice at 6 months, respectively.

Abbreviations: Cm, resting membrane capacitance; Rm, membrane resistance; and Rs: series resistance.
Table 2. Numbers of mice, cochleae and hair cells examined in scanning electron microscopy in VGLUT3+/+, VGLUT3+/A224V and VGLUT3A224V/A224V mice at 1, 2, 4 and 6 months of age

<table>
<thead>
<tr>
<th>Age</th>
<th>1 month</th>
<th>2 months</th>
<th>4 months</th>
<th>6 months</th>
</tr>
</thead>
<tbody>
<tr>
<td>VGLUT3+/+</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Three mice</td>
<td>Three cochleae</td>
<td>Two mice</td>
<td>Two mouse cochleae</td>
<td>Six mice</td>
</tr>
<tr>
<td>IHCs: 815</td>
<td>OHCs: 2931</td>
<td>IHCs: 472</td>
<td>OHCs: 1860</td>
<td>IHCs: 1366</td>
</tr>
<tr>
<td>VGLUT3+/A224V</td>
<td>Two mice</td>
<td>Three cochleae</td>
<td>Three mouse cochleae</td>
<td>Two mouse cochleae</td>
</tr>
<tr>
<td>IHCs: 743</td>
<td>OHCs: 3171</td>
<td>IHCs: 1140</td>
<td>OHCs: 4580</td>
<td>IHCs: 552</td>
</tr>
<tr>
<td>VGLUT3A224V/A224V</td>
<td>Five mice</td>
<td>Six cochleae</td>
<td>Five mouse cochleae</td>
<td>Four mouse cochleae</td>
</tr>
<tr>
<td>IHCs: 1217</td>
<td>OHCs: 5256</td>
<td>IHCs: 732</td>
<td>OHCs: 3015</td>
<td>IHCs: 1140</td>
</tr>
</tbody>
</table>

IHC: inner hair cell; OHC: outer hair cell.

30 IHCs (n = 4 cochleas) at 2 months, 73 IHCs (n = 4 cochleas) at 4 months, 61 IHCs (n = 4 cochleas) at 6 months from VGLUT3+/+ mice and 39 IHCs (n = 3 cochleas) at 2 months, 118 IHCs (n = 3 cochleas) at 4 months, 83 IHCs (n = 4 cochleas) at 6 months from VGLUT3A224V/A224V mice. Measurement of ribbon size was obtained by (i) surrounding the ribbon using contour function (iso-intensity line fixed at mean background + 2 SD) and (ii) quantifying length of long axis (a_lg) and short axis (a_sh) using the fit_ellipse function (least-squares criterion). The elliptical surface (S_{el}) was then calculated using the equation S_{el} = (a_lg \times a_sh \times \pi) / 4.

Electron microscopy

Scanning (SEM) and transmission (TEM) electron microscopy were done for the anatomical examination of cochlear hair cells. For both techniques, the animals were decapitated under deep anaesthesia (pentobarbital, 50 mg/kg), their cochleae were removed from the temporal bone. For SEM, PBS-washed cochleae were fixed with 2.5% glutaraldehyde in phosphate buffer, pH 7.2 for 2 h at room temperature, followed by washing in phosphate buffer. The bony capsule of the cochlea was dissected out, and the stria vascularis as well as the tectorial and Reissner's membranes were removed. Fixed cochleae were dehydrated using a graded ethanol series (15–100%), followed by critical point drying with CO2. Subsequently, the samples were sputter-coated with an approximatively 10 nm thick gold film and then examined under a scanning electron microscope (Hitachi S4000) using a lens detector with an acceleration voltage of 10 kV at calibrated magnifications. The numbers of mice, cochleae and total numbers of IHCs and OHCs are given in the Table 2.

For TEM, the cochleae were perfused with a solution of 2.5% glutaraldehyde in PHEM buffer (1X, pH 7.4) and immersed in the same fixative for 1 h at room temperature, then overnight at 4°C. Samples were then rinsed in PHEM buffer and post-fixed in a 0.5% osmic acid for 2 h in the dark and at room temperature. After two rinses in PHEM buffer, the cells were dehydrated in a graded series of ethanol solutions (30–100%). The cells were embedded in EmBed 812 using an Automated Microwave Tissue Processor for Electronic Microscopy, Leica EM AMW. Thin sections (70 nm; Leica-Reichert Ultracut E) were collected at different levels of each block. These sections were counterstained with uranyl acetate 1.5% in 70% ethanol and lead citrate and observed using a Tecnai F20 transmission electron microscope at 200 KV. Both SEM and TEM were carried out at the CoMET facility (MRI, INM, Montpellier, France).

Data analysis

Data were analysed using Igor Pro (WaveMetrics, Lake Oswego, OR, USA), MATLAB and R (R core team) software. All these data are expressed as means ± standard deviation (SD) and were compared by two-tailed Mann–Whitney Wilcoxon’s test, Student’s t test or a two-sample test for equality of proportions.

Results

Auditory neuropathy in VGLUT3A224V/A224V mouse

We first examined the synchronous activation of the ascending auditory system of wild-type (VGLUT3+/+) mice, heterozygous (VGLUT3+/A224V) mice and homozygous (VGLUT3A224V/A224V) mice (Fig. 1). In heterozygous mice, we observed a slight threshold shift.
Deficit of hair cell transduction in DFNA25

(mean threshold at 1 month: 28.2 ± 5.81 dB SPL vs. 33.6 ± 1.88 dB SPL in VGLUT3+/+ and VGLUT3+/+A224V mice, respectively, P = 0.003, two-tailed Mann–Whitney Wilcoxon’s test; at 6 months: 43.4 ± 6.1 dB SPL vs. 53 ± 3.8 dB SPL in VGLUT3+/+ and VGLUT3+/+A224V mice, respectively, P = 0.001, two-tailed t test; Fig. 1F–J). The alteration in the ABR waveform was associated with a progressive threshold shift (mean threshold at 1 month: 28.2 ± 5.8 dB SPL vs. 35.4 ± 10.2 dB SPL in VGLUT3+/+ and VGLUT3+/+A224V mice, respectively, P = 0.007; at 6 months: 43.4 ± 6.1 dB SPL vs. 67.7 ± 6 dB SPL in VGLUT3+/+ and VGLUT3+/+A224V mice, respectively, P = 1.10^{-6}, two-tailed Mann–Whitney Wilcoxon’s test; Fig. 1F–J). To determine whether the hearing loss was due to defective OHCs, we probed the DPOAEs. We observed robust DPOAEs in the heterozygote and homozygote mice, indicating that OHC mechanical activity was essentially preserved (2f1-f2 @ f2 = 11.9 kHz: 39.6 ± 7.3 dB SPL, 38.5 ± 4.3 dB SPL and 44.4 ± 7.6 dB SPL in 6-month-old VGLUT3+/+ mice, VGLUT3+/+/A224V mice, and VGLUT3+/+A224V mice, respectively; Fig. 1K–O). Altogether, VGLUT3+/+A224V mice showed a progressive hearing loss with functional cochlear amplification.

Next, we carried out electrocochleography to probe the activity of IHCs and the synchronous activation of afferent fibres (Fig. 2A). CAP was strongly diminished in 1- to 2-month-old VGLUT3+/+A224V mice, consistent with the reduction of the ABR wave 1 (mean CAP N1-P1 amplitude: 203.4 ± 80.6 μV vs. 80.7 ± 37 μV in VGLUT3+/+ and VGLUT3+/+A224V mice, respectively; Fig. 2A).
Fig. 2B). In addition, we observed a smaller SP, reflecting the receptor potential in IHCs (mean SP amplitude: 93.3 ± 57.1 μV vs. 50.2 ± 50.8 μV in VGLUT3+/+ and VGLUT3A224V/A224V mice, respectively, P = 0.007, two-tailed Mann–Whitney Wilcoxon’s test; Fig. 2C). These results implied a selective alteration of the IHC’s function in VGLUT3A224V/A224V homozygous mice.

**Alteration at the stereociliary bundle in VGLUT3A224V/A224V mouse inner hair cells**

Given the reduction in the SP, we examined the ultrastructure of the stereocilia bundle from IHCs at the mid-apical region of the cochlea (e.g. the region encoding frequency up to 20 kHz) using SEM (Fig. 3). In VGLUT3+/+ mice, the apical side of IHCs projected distinct and erected stereocilia (Figs. 3A–D and 4A–D). In heterozygous mice, stereocilia bundles were found to be normal at 1 month of age (Figs. 3E and 4E, M) but a small fraction of IHCs harbouring fused stereocilia was observed at 2 months (2.7 ± 0.7 % vs. 8.6 ± 0.8 % in VGLUT3+/+ and VGLUT3A224V/A224V mice, respectively, P = 4.10⁻⁵, two-sample test for equality of proportions; Figs. 3F and 4F, N). Although some IHCs with a distorted bundle were seen, the number of IHCs with these abnormal stereocilia did not differ between heterozygous and wild-type mice at later ages (Figs. 3G–H and 4G–H, O–P). In homozygous VGLUT3A224V/A224V mice, we found more than 20% IHCs with splayed stereocilia already at 1 month (4.8 ± 0.7 % vs. 23.7 ± 1.2% in VGLUT3+/+ mice and VGLUT3A224V/A224V mice, respectively, P = 2.10⁻¹⁶, two-sample test for equality of proportions; Figs. 3I and 4I, M). Collapsed stereocilia bundles were frequently observed at later stages (9.5 ± 0.8 % vs. 35.1 ± 1.4 % in 4-month-old VGLUT3+/+ and VGLUT3A224V/A224V mice, respectively, P = 2.10⁻¹⁶, two-sample test for equality of proportions; Figs. 3J–L and 4J–L, N–P). By plotting the percentage of normal, altered and missing hair cells as a function of the age, we found that the fraction of IHCs with massive fusion or even absent cilia apparatus significantly increased with age (12.7 ± 1% vs. 61 ± 1.7% in 6-month-old VGLUT3+/+ and VGLUT3A224V/A224V mice, respectively, P = 2.10⁻¹⁶, two-sample test for equality of proportions; Fig. 6A). Together, these data suggest that the threshold shift in VGLUT3A224V/A224V mice is associated with a defective ultrastructure of stereocilia from IHCs.

OHCs had a normal appearance, even though the IHCs with disrupted bundles were located at the same region (Figs 3 and 6B). High magnification SEM micrograph confirmed that the stereocilia bundle in OHCs appeared well-organized, with a staircase pattern, consistent with preserved DPOAEs (Fig. 5). Altogether, these data indicated a selective alteration of the stereocilia from IHCs.

**VGLUT3 is normally distributed in inner hair cells from VGLUT3A224V/A224V mice**

The variant strongly decreases (-70%) the expression of VGLUT3-p.A224V in the terminals of the striatum or hippocampus neurons, leaving its accumulation in the soma unaffected (Ramet et al. 2017). Thus, we wondered whether the p.A224V variant might alter the distribution of VGLUT3 within IHCs, especially in the basolateral area. Semi-quantitative analysis of the VGLUT3 distribution in IHCs from 1- to 2-month-old VGLUT3+/+ and VGLUT3A224V/A224V mice

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**Figure 2. Inner hair cell receptor potential is altered in VGLUT3A224V/A224V mouse**

A, mean compound action potential (CAP; N1-P1 amplitude) ± SD and summating potential (SP) ± SD, reflecting the auditory afferent fibre activation and inner hair cell receptor potential, respectively. CAP and SP were evoked by 16 kHz tone burst at 100 dB SPL in 1–2-month-old VGLUT3+/+ and VGLUT3A224V/A224V mice. n indicates the number of cochleae recorded. B and C, mean CAP amplitude ± SD (B) and SP amplitude ± SD (C) from 1–2-month-old VGLUT3+/+ and VGLUT3A224V/A224V mice. SP were measured at the black marker in (A). Level of significance: ** P < 0.01; *** P < 0.001, two-tailed Mann–Whitney Wilcoxon’s test. Symbols represent individual cochleae.
immunofluorescence throughout the cell, from the upper side of the nucleus to the basolateral side, showed similar targeting of VGLUT3-p.A224V between both genotypes (VGLUT3 immunofluorescence: 1710 ± 219 A.U. vs. 1678 ± 416 A.U. μm in IHCs from 4-month-old VGLUT3+/+ and VGLUT3A224V/A224V mice, respectively; P = 0.565; Fig. 7). Therefore, in contrast to what is observed in the central nervous system neurons (Ramet et al. 2017), the p.A224V variant does not disturb the VGLUT3 targeting within the hair cell cytoplasm.

**Alteration of the synaptic ribbon size in VGLUT3A224V/A224V mouse inner hair cells**

Given the role of VGLUT3 in the synaptic transfer between the IHC and the afferent nerve fibre, we addressed the question of whether additional consequences might occur in the pre-synaptic compartment and specifically in the synaptic ribbon. Figure 8 shows transmission electron microscopy examples of plasma-membrane anchored synaptic ribbons with a halo of synaptic vesicles. Multiple ribbons facing a single afferent terminal were observed in the IHCs from VGLUT3A224V/A224V mice but the low number of samples in our study precluded any quantitative analysis. Indeed, multiple ribbons are also found in a fraction of synapses in adults (Merchan-Perez & Liberman, 1996; Michanski et al. 2019; Payne et al. 2021). Therefore, we used immunolabelling to measure the number and size of synaptic ribbons. From 2 to 6 months, the number of ribbons immunolabelled with anti-Ctbp2 antiserum was comparable between wild-type mice and VGLUT3A224V/A224V mice (Fig. 9A–E). In addition, we did not notice any difference in the ribbon synapse number corresponding to the juxtaposed immunostainings of Ctbp2 and post-synaptic scaffolding protein Homer (at 6 months: 10.5 ± 1.9 synapses per IHC vs. 11 ± 0.7 synapses per IHC in VGLUT3+/+ mice and in VGLUT3A224V/A224V mice, respectively, P = 0.642; Fig. 9F). However, synaptic ribbons became abnormally elongated in knock-in mice (Fig. 9B, D). Using STED super-resolution microscopy in 2-month-old mice, Ctbp2-immunolabelled signals looked like ovoid spots. Synaptic ribbons in VGLUT3A224V/A224V mice were slightly but significantly larger (long object axis: 440.3 ± 125 nm vs. 458.2 ± 114 nm in VGLUT3+/+ mice and VGLUT3A224V/A224V mice, respectively, P = 0.213; short object axis: 287.4 ± 48 nm in VGLUT3+/+ and 302.4 ± 50 nm VGLUT3A224V/A224V mice, respectively, P = 0.009, Student’s t test; Fig. 10A–C). In contrast, we could quantify a drastic elongation of ribbons from IHCs in VGLUT3A224V/A224V mice at 4 and 6 months (at 6 months: long object axis: 473.7 ± 142 nm vs. 602.2 ± 285 nm in VGLUT3+/+ and VGLUT3A224V/A224V mice, respectively, P = 2.10-20; short object axis: 300.1 ± 61 nm in VGLUT3+/+ and 344.4 ± 111 nm VGLUT3A224V/A224V mice, respectively, P = 1.10-15, Student’s t test; Fig. 10D–I). Consistently, mean elliptical size calculated from long and short ribbon axes were comparable at 2 months of age between the genotypes but significantly increased at
4 and 6 months in IHCs from VGLUT3<sup>A224V/A224V</sup> mice (elliptical size: 0.102 ± 0.04 μm<sup>2</sup> vs. 0.111 ± 0.04 μm<sup>2</sup> in 2-month-old VGLUT3<sup>+/+</sup> and VGLUT3<sup>A224V/A224V</sup> mice, respectively and 0.116 ± 0.05 μm<sup>2</sup> vs. 0.18 ± 0.15 μm<sup>2</sup> in 6-month-old VGLUT3<sup>+/+</sup> and VGLUT3<sup>A224V/A224V</sup> mice, respectively, P = 3.10^-4, Student’s t test; Fig. 11A). Size distributions were skewed for 4- and 6-month-old VGLUT3<sup>A224V/A224V</sup> mice (Fig. 11B). The cumulative size histogram showed that IHCs from 6-month-old VGLUT3<sup>A224V/A224V</sup> mice contain ribbons resembling those found at 2 months as well as ribbons for which size increased up to 100 % (ribbon size up to 0.064 μm<sup>2</sup> and 0.065 μm<sup>2</sup> within the 10th percentile against 0.17 μm<sup>2</sup> and 0.35 μm<sup>2</sup> within the 90th percentile from 6 month IHCs in VGLUT3<sup>+/+</sup> mice and VGLUT3<sup>A224V/A224V</sup> mice, respectively; Fig. 11C). These results indicated that the p.A22AV variant did not affect ribbons in the same manner. Small synaptic bodies remained small, while large pre-synaptic ribbons became larger in VGLUT3<sup>A224V/A224V</sup> mice.

**Calcium-triggered exocytosis in VGLUT3<sup>A224V/A224V</sup> mouse inner hair cells**

To determine whether synaptic release was affected in IHCs of VGLUT3<sup>A224V/A224V</sup> mice, we measured the membrane capacitance as a proxy of calcium-triggered exocytosis using perforated patch-clamp (Fig. 12). In hair cells, voltage steps of short duration (under 20 ms) recruit the readily releasable pool (RRP) that corresponds to the release of synaptic vesicles in the vicinity of the calcium channels, while longer duration of the stimuli (beyond 20 ms) mobilizes the SRP which reflects the exocytosis of remote vesicles and/or the resupply of synaptic vesicles.
to the release site (Moser & Beutner, 2000). In our experiments, secretion by IHCs was tested at two different time points: that is, before (1–2 months of age) and after (6 months) the elongation of synaptic ribbons. As shown in Fig. 12A, depolarizing voltage steps evoked calcium influx, which in turn triggered the fusion of synaptic vesicles to the plasma membrane, mirrored by jumps in the membrane capacitance. Between 1 and 2 months, the calcium current–voltage relationships were comparable (calcium peak current $I_{Ca}$: $-155.2 \pm 34.9$ pA vs. $-159.1 \pm 35.2$ pA for VGLUT3$^{+/+}$ mice and VGLUT3$^{A224V/A224V}$, respectively, $P = 0.817$; Fig. 12B1). Furthermore, the exocytic membrane capacitance was of indistinguishable amplitude between genotypes for any stimulus duration (ΔCm$_{20ms}$: $12.6 \pm 5.8$ fF vs. $14.6 \pm 6.9$ fF for VGLUT3$^{+/+}$ mice and VGLUT3$^{A224V/A224V}$, respectively, $P = 0.297$; ΔCm$_{100ms}$: $38.9 \pm 14.6$ fF vs. $35.3 \pm 12.7$ fF for VGLUT3$^{+/+}$ and VGLUT3$^{A224V/A224V}$, respectively, $P = 0.489$; Fig. 12C1). In addition, the calcium influx was of similar amplitude (Q$_{Ca}^{2+}_{20ms}$: $2.9 \pm 0.6$ pC vs. $3 \pm 0.6$ pC for VGLUT3$^{+/+}$ mice and VGLUT3$^{A224V/A224V}$ mice, respectively, $P = 0.578$; Q$_{Ca}^{2+}_{100ms}$: $13.8 \pm 3.1$ pC vs. $14.3 \pm 3.2$ pC for VGLUT3$^{+/+}$ mice and VGLUT3$^{A224V/A224V}$ mice, respectively, $P = 0.456$; Fig. 12C1). Thus, the exocytosis jumps plotted against the corresponding calcium influx overlapped (Fig. 12D1) and the release efficiency, that is, the exocytosis per unit of incoming calcium, was not different between wild-type mice and homozygous mice for short and long depolarizing voltage steps (ΔCm/Q$_{Ca}^{2+}_{20ms}$: $4.4 \pm 2.3$ fF/pC vs. $4.8 \pm 2$ fF/pC, $P = 0.309$ and ΔCm/Q$_{Ca}^{2+}_{100ms}$: $2.8 \pm 0.8$ fF/pC vs. $2.5 \pm 0.7$ fF/pC, $P = 0.249$ for VGLUT3$^{+/+}$ mice and VGLUT3$^{A224V/A224V}$ mice, respectively; Fig. 12E1). At 6

Figure 5. Stereociliary bundle morphology in outer hair cells

A–L, SEM of the stereocilia bundle from outer hair cells (OHCs) at mid-apical turns in 1–6-month-old VGLUT3$^{+/+}$, VGLUT3$^{+/A224V}$ and VGLUT3$^{A224V/A224V}$ mice. Scale bars: 2 μM. M–P, fraction of OHCs harbouring normal or altered stereocilia (ST) and of missing OHCs. # Indicates the animal’s number. Percentages are shown with their SD. Total numbers of OHCs examined are indicated in the histograms. Symbols represent individual cochleae. Level of significance: * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$, two-sample test for equality of proportions.
months of age, the calcium current–voltage relationship still did not substantially differ between genotypes (calcium peak current $I_{Ca}$: $-127.9 \pm 23$ pA vs. $-147.7 \pm 49.5$ pA for VGLUT3$^{+/+}$ mice and VGLUT3$^{A224V/A224V}$ mice, respectively, $P = 0.231$; Fig. 12B2). Although the exocytosis in IHCs evoked by short time stimulus duration (RRP, under 20 ms) remained identical ($\Delta Cm_{20ms}$ 8.7 ± 5.5 fF vs. 11.9 ± 6.6 fF for VGLUT3$^{+/+}$ mice and VGLUT3$^{A224V/A224V}$ mice, respectively, $P = 0.270$; Fig. 12C2), we observed a larger increase of exocytosis from IHC VGLUT3$^{A224V/A224V}$ mice in response to long depolarizing voltage steps ($\Delta Cm_{100ms}$: 32.7 ± 17.8 fF vs. 55.6 ± 25.1 fF for VGLUT3$^{+/+}$ and VGLUT3$^{A224V/A224V}$, respectively; $P = 0.025$, two-tailed Mann–Whitney

Figure 6. Time course of the stereocilia bundle alteration
Percentage of normal (empty circle), altered (filled square) and missing (empty triangle) IHC (A) and outer hair cell (B) plots as a function of the age in the VGLUT3$^{+/+}$ (blue), VGLUT3$^{+/+}$ (green) and VGLUT3$^{A224V/A224V}$ (red) mice. Percentages are shown with their confidence interval. Stars indicate significant difference between heterozygotes (green star) or homozygotes (red star) vs. wild type. Symbols represent individual cochleae. Level of significance: * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$, two-sample test for equality of proportions.

Figure 7. VGLUT3 distribution in inner hair cells from VGLUT3$^{+/+}$ and VGLUT3$^{A224V/A224V}$ mice
A and B, maximum projection average of confocal sections of 4-month-old VGLUT3$^{+/+}$ (A) and VGLUT3$^{A224V/A224V}$ (B) mid-apical turns IHCs immunolabelled against VGLUT3 (green) processed and imaged under identical conditions. Left panel: IHCs have been centred in respect to the centroid of the nucleus; right panel: average of the maximum projection fluorescence ± SD measured through the vertical central axis crossing the IHCs (dashed white line). $n$ indicates the number of IHCs immunolabelled from four cochleae for each genotype. C, semi-quantitative analysis of VGLUT3 immunofluorescence. Bar histograms represent the integral of the maximum projection fluorescence ± SD measured through the vertical central axis crossing the IHCs of 4-month-old VGLUT3$^{+/+}$ (blue) and VGLUT3$^{A224V/A224V}$ (red) corresponding to (A and B), respectively. Symbols represent individual hair cells.
Wilcoxon’s test; Fig. 12C2). Consequently, the rate of the sustained exocytosis measured by the slope of the line fit to the exocytic capacitance between 20 and 100 ms increased in homozygous mice (ΔCm20–100 ms = 300.1 ± 174 fF.s⁻¹ vs. 545.4 ± 278.5 fF.s⁻¹ for VGLUT3⁺/⁺ mice and VGLUT3A224V/A224V mice, respectively; P = 0.021, two-tailed Mann–Whitney Wilcoxon’s test). To determine whether the larger secretion was due to an increase in the corresponding calcium influx, we first measured the calcium influx but we did not find a significant difference between the mouse lines (QCa²⁺ 100 ms = 11.2 ± 2.5 pC vs. 13.6 ± 5.1 pC for VGLUT3⁺/⁺ mice and VGLUT3A224V/A224V mice, respectively; P = 0.425; Fig. 12C2). Next, we plotted the membrane capacitance changes against the corresponding calcium entry. Still, we found an increase in release for overlapping range of incoming calcium (Fig. 12D2). Finally, the release efficiency was comparable between wild-type mice and homozygous mice for short depolarizing steps (Cm/ QCa²⁺ 20 ms: 3.5 ± 2.1 fF/pC vs. 4.2 ± 1.8 fF/pC for VGLUT3⁺/⁺ mice and VGLUT3A224V/A224V mice, respectively; P = 0.242; Fig. 12E2) but larger in the VGLUT3A224V/A224V homozygous mice for longer duration (Cm/ QCa²⁺ 100 ms: 2.8 ± 1.4 fF/pC and 4.3 ± 1.7 fF/pC for VGLUT3⁺/⁺ mice and VGLUT3A224V/A224V mice, respectively; P = 0.036, two-tailed Mann–Whitney Wilcoxon’s test; Fig. 12E2). Thus, the p.A224V variant of the SLC17A8 gene is associated with an increase in the SRP leaving the calcium influx unaffected.

Discussion

In this study, we showed that the mutant mouse carrying the p.A224V variant of the SLC17A8 gene develops a progressive hearing loss, phenocopying human auditory neuropathy. The mutation appears to primarily alter the stereociliary bundle structure of IHCs followed by a change in ribbon size and exocytosis (Fig. 13).

The VGLUT3A224V/A224V mouse as a model of progressive auditory neuropathy

Only a few mutations that cause a progressive and early-onset form of human deafness DFNA25 have been described (Ruel et al. 2008; Ryu et al. 2016). Among them, the c.616dupA variant of SLC17A8 introduces a stop codon and hence leads to a truncated protein (Ryu et al. 2016; Qi et al. 2021). The p.A211V allele of SLC17A8 is an additional variant that has not yet been fully characterized (Ruel et al. 2008). Here, the mouse line carrying two copies of the VGLUT3-p.A224V allele shows increasing threshold shift with age. Thus, this result further validates the p.A211V as one of the mutations responsible of DFNA25 (Ruel et al. 2008). However, the mild hearing impairment in the VGLUT3⁺/A224V heterozygotes contrasts with the dominant transmission in DFNA25 (Greene et al. 2001; Thirlwall et al. 2003). This might be explained by the slow rate in the stereocilia alteration with respect to the short mouse lifespan. Decent DPOAEs were recorded in the knock-in mutant mouse. In contrast, otoacoustic emissions were absent in patients harbouring the p.A211V allele (Thirlwall et al. 2003). Thus, DFNA25 is more likely to be a progressive hearing loss in human rather than an auditory neuropathy (i.e. absent or desynchronized ABR with intact DPOAEs). Recent work has shown that VGLUT3 is expressed in OHCs to signal to the type II auditory nerve fibres (Weisz et al. 2021). However, robust DPOAEs measured in the VGLUT3-null mice do not support a role of VGLUT3 in cochlear amplification (Ruel et al. 2008; Seal et al. 2008; Kim et al. 2019). The lack of otoacoustic emissions in DFNA25 patients might be due to additional factors encountered during lifetime such as noise (Thirlwall et al. 2003). Consistently, the vulnerability to noise in mice carrying one or two variant alleles must be probed to determine whether acoustic injury accounts for the above discrepancies. Altogether, the mouse model harbouring the VGLUT3-p.A224V variant mimics a progressive hearing loss with auditory neuropathy features, but might be arguable so far as to consider it as a faithful model of DFNA25.
Collapse of the stereocilia bundle in VGLUT3A224V/A224V

In the VGLUT3A224V/A224V homozygous mice, the major alteration lies in the stereocilia architecture from IHCs. This is somehow surprising since VGLUT3 is absent from IHC stereocilia and the loss of VGLUT3 does not affect the transducer activity, as shown by intact hair cell receptor potential in the absence of VGLUT3 (Obholzer et al. 2008; Ruel et al. 2008). The p.A224V variant may result in a protein misfolding that leads to its aberrant aggregation and to distal adverse effects. A massive protein accumulation could create a traffic jam at the upper

Figure 9. Ribbons and synapses count in inner hair cells

A–D, synaptic ribbons and post-synaptic densities immunostained with antibodies against Ctbp2 (green) and Homer (red), respectively, in inner hair cells (IHCs) from 6-months VGLUT3+/+ (A, C) and VGLUT3A224V/A224V (B, D) mice. Scale bar in (A and B): 5 μm. C and D show high magnification of the white dashed squares from (A) and (B), respectively. Scale bar in (C, D): 1 μm. E, mean ± SD number of ribbons per IHC given by the immunostaining of Ctbp2 from 2-, 4- and 6-month-old VGLUT3+/+ and VGLUT3A224V/A224V mice. F, mean ± SD number of synapses per IHC given by the juxtaposed immunostaining of Ctbp2 and Homer from IHCs of 6-month-old VGLUT3+/+ and VGLUT3A224V/A224V mice. White numbers indicate the number of IHCs analysed. Symbols represent average for individual cochlea. For each age and genotype, four cochleae have been used except for 6-month quantification in E (n = five cochleae).

Figure 10. Synaptic ribbon morphology in inner hair cells

A–B, D–E and G–H, representative STED images of fluorescently labelled synaptic ribbons of 2-month (A, B), 4-month (D, E) and 6-month-old (G, H) inner hair cells from VGLUT3+/+ (A, D and G) and VGLUT3A224V/A224V (B, E and H) mice. Scale bar: 1 μm. C, F and I, long vs. short axes for VGLUT3+/+ (blue) and VGLUT3A224V/A224V (red) ribbons at 2 months (C), 4 months (F) and 6 months (I). n indicates the number of ribbons examined, displayed as small dots. Insets show the mean values and SD for long vs. short axes. ** P < 0.01 and *** P < 0.001, Student’s test for independent samples.
side of the nucleus and destabilize the route for protein turnover towards the stereocilia machinery. The failure in proteolysis has been shown to eventually provoke a disoriented stereocilia bundle (Freeman et al. 2019). In the human auditory neuropathy AUNA1, the over-expression of the diap3 protein leads to microtubule accumulation within the cuticular plate followed by the collapse of the stereocilia (Surel et al. 2016). However, in the striatum and hippocampal neurons, the point mutation leads to a reduced expression in the neuron terminals leaving the amount of VGLUT3 in the soma unaltered (Ramet et al. 2017). This argues against an abnormal accumulation of the p.A224V variant in the soma compartment preventing its distribution towards the terminals. In addition, in a virtual 3D model, replacing alanine in position 211 (224 in mouse) by a valine does not substantially modify the VGLUT3 structure as alanine and valine amino acid are closely related (Ramet et al. 2017). Finally, we did not observe an increase in the VGLUT3 immunofluorescence at the apical side of IHCs, although this relies on a semi-quantitative approach. The mechanisms underlying the homogeneous distribution of mutated VGLUT3 in hair cells vs. the differential expression in the central nervous system neurons are not known but hair cell can be assimilated to a single compartment without ‘neurites’. Further experiments are therefore required to determine whether unfolded protein response operates in VGLUT3^{A224V/A224V} homozygous mice and whether the cytoskeleton is disrupted at the IHC apical side. Other mechanisms may contribute to the progressive destruction of the stereocilia. For instance, the genetic background of the mouse line used in our study is known to be homozygous for the defective Cdh23ahl allele of the gene encoding cadherin 23, which is a constituent of hair cell stereocilia (Noben-Trauth et al. 2003; Siemens et al. 2004; Kazmierczak et al. 2007). Although VGLUT3 is not localized at the stereocilia bundle level, we cannot exclude the possibility that the Cdh23ahl allele potentiates, in any way, the stereocilia distortion. Large calcium influx in the C57Bl/6J has also been proposed to contribute to high-frequency hearing loss (Liu et al. 2019). Calcium overflow may exacerbate the phenotype in the VGLUT3^{A224V/A224V} mice but sound encoding is not altered at 3–4 months of age in the triple knock-out mice for endogenous intracellular calcium buffers, making a potential calcium overload that would deteriorate the hair cells unlikely (Pangrišić et al. 2015).

### Synaptic plasticity in VGLUT3^{A224V/A224V} mice

Akin to VGLUT3^{−−} mice (Seal et al. 2008; Akil et al. 2012; Kim et al. 2019), we observe that the anatomy of synaptic ribbons changed at later stages. It has been shown that the synaptic ribbon size varies with the calcium influx amplitude (Martinez-Dunst et al. 1997; Schnee et al. 2005; Frank et al. 2009; Regus-Leidig & Specht, 2010; Sheets et al. 2012; Ohn et al. 2016). However, the lack of change in the calcium current amplitude in the IHCs of VGLUT3^{A224V/A224V} mice excludes such a mechanism. The p.A224V mutation occurs within a cytoplasmic loop that faces the pore of the transporter and thus may impede the loading of intra-vesicular glutamate. In this model, less glutamate could be released into the synaptic cleft leading to a reduced activity of the post-synaptic receptors and to poor recruitment of the

**Figure 11. Increase of the elliptic surface of ribbon’s section in inner hair cells**

A, mean ± SD elliptic surface calculated from the ribbon’s long and short axes lengths of 2-, 4- and 6-month-old IHCs from VGLUT3^{+/+} and VGLUT3^{A224V/A224V} mice. Symbols represent individual ribbons. B, distribution of the ribbons plots against the elliptical surface estimates of their cross-section derived from STED-imaging (bin width = 0.04 μm²). C, cumulative distribution of the ribbons as a function of the elliptical surface (bin width = 0.0001 μm²). Open symbols indicate the 10th (triangle) and 90th (diamond) percentile. Level of significance: *** P < 0.001, Student’s test for independent samples.

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afferent auditory nerve fibres. In hair cells, synapses with larger ribbons are associated with reduced activity (Merchan-Perez & Liberman, 1996; Sheets et al. 2012; Ohn et al. 2016; Kim et al. 2019). While high-spontaneous rate fibres with low-threshold activation project onto small synaptic ribbons on the pillar side, low-spontaneous rate (LSR) fibres with high-threshold activation connect to large ribbons on the modiolar side (Merchan-Perez & Liberman, 1996; Liberman et al. 2011; Ohn et al. 2016). The increase in the number of thin and elongated synaptic ribbons may reflect an increase in the LSR fibre pool connecting the IHCs in VGLUT3A224V/A224V mice. In this case, the hearing loss at late stages in VGLUT3A224V/A224V mice could not only be attributable to the stereocilary bundle defect but also to the delayed and high jitter of the first-spike latency within auditory nerve fibres.

Figure 12. Increase of the sustained releasable pool exocytosis in 6-month-old inner hair cells of VGLUT3A224V/A224V mice

A1, E1, perforated patch-clamp recordings of calcium-triggered exocytosis in 1/2- (A1, E1) and 6-month-old (A2, E2) inner hair cells (IHCs) of VGLUT3+/+ (blue) and VGLUT3A224V/A224V mice (red). A1, A2, inward calcium current (ICa2+), membrane capacitance jump (ΔCm), membrane resistance (Rm) and series resistance (Rs) evoked by 20, 50 and 200 ms long depolarizing voltage steps from holding potential of -87 mV to -27 mV. B1, B2, C2, exocytic ΔCm and calcium charge QCa2+ plotted against the duration of depolarization. Insets show the membrane capacitance increases in response to depolarizing voltage steps from 5 to 20 ms duration. Calcium-triggered exocytosis was evoked by depolarizations to -27 mV from holding potential of -87 mV. n: number of IHCs. D1, D2, exocytic ΔCm plotted against the corresponding calcium charge QCa2+ for 20, 50 and 100 ms step depolarization. * P < 0.05, two-tailed Mann–Whitney Wilcoxon’s test. Data are means ± SD. Thin lines and symbols represent individual cells.
Although we cannot rule out a reduction in the amount of glutamate released by the hair cells, the overexpression of VGLUT3-p.A224V variant in autaptic culture almost totally rescues the defect in exocytosis (Ramet et al. 2017). Consistently, the p.A224V mutation does not alter the vesicular uptake probed in heterologous system (Ramet et al. 2017). Thus, measurement of the afferent terminal activity is required to determine whether the p.A224V reduced the quantal sizes particularly at the hair cell ribbonsynapse (Glowatzki & Fuchs, 2002; Özcete & Moser, 2020).

The IHCs calcium-triggered exocytosis probed by short-time duration corresponds to the RRP of synaptic vesicle. In our experiments, the RRP is not affected in IHCs from VGLUT3A224V/A224V mice. In hair cells, calcium channels are densely packed underneath the pre-synaptic element and the RRP correlates to the membrane-proximal synaptic vesicles, that is, the vesicles tether to the synaptic body and docked to the plasma membrane (Khimich et al. 2005; Schnee et al. 2005; Rutherford & Roberts, 2006; Meyer et al. 2009; Frank et al. 2010; Pangršić et al. 2010; Graydon et al. 2011; Wong et al. 2014; Neef et al. 2018). Thus, the calcium current and the RRP size are likely to scale with the number of synapses (Meyer et al. 2009). The comparable calcium-current amplitude and similar amount of RRP between both genotypes are quite consistent with the preserved number of synapses.

The secretion evoked by longer time duration corresponds to the SRP of synaptic vesicles. In our experiments, the SRP is not affected in IHCs from VGLUT3A224V/A224V mice. In hair cells, calcium channels are densely packed underneath the pre-synaptic element and the RRP correlates to the membrane-proximal synaptic vesicles, that is, the vesicles tether to the synaptic body and docked to the plasma membrane (Khimich et al. 2005; Schnee et al. 2005; Rutherford & Roberts, 2006; Meyer et al. 2009; Frank et al. 2010; Pangršić et al. 2010; Schnee et al. et al. 2011; Moser et al. 2019). The secretion associated with the SRP is dramatically increased in mutant mice. This change may be due to a larger number of synaptic vesicles populating the synaptic ribbons as they become elongated in the mutant mouse (Kantarzhiieva et al. 2013; Payne et al. 2021). In this hypothesis, the synaptic vesicles attached to the ribbon and facing the cytoplasm, the ‘ribbon-associated SV’ might be the morphological correlate to the SRP (Hull et al. 2006; Frank et al. 2010; Snellman et al. 2011; Maxeiner et al. 2016). However, the mutant models in which the ribbons are missing do not support this hypothesis: the ablation of RIBEYE in mice or zebrafish hair cells leading to a ribbon-free active zone does not impair the secretion probed by capacitance measurements (Lv et al. 2016; Becker et al. 2018; Jean et al. 2018). In the triple knock-out mice for endogenous calcium buffers, a larger SRP has been measured using membrane capacitance patch-clamp recordings but the auditory thresholds and driven-rate of afferent fibres remained largely unaffected, suggesting extra-synaptic release (Pangršić et al. 2015). The increase in the SRP in the p.A224V homozygous mutant may be similar and reflect a form of pre-synaptic homeostatic plasticity, that is, release more glutamate to overcome a deficit in mechanotransduction with eventually a spiking rate reduction in the afferent auditory nerve fibre (Delvendahl et al. 2019).

In conclusion, these results highlight the critical role of VGLUT3 in maintaining the hair cell integrity to operate sound coding. The VGLUT3A224V/A224V mouse will be a valuable asset to fully elucidate molecular mechanisms underlying DFNA25 and to identify an effective treatment.

A, in response to incoming sound stimulation, stereocilia are deflected, leading to the mechanotransducer channel opening and the depolarization of the inner hair cell. The calcium influx through the voltage-gated calcium channel triggers the exocytosis of glutamate-laden synaptic vesicles. The release of glutamate in the synaptic cleft activates the auditory nerve fibre, which conveys the neural message to the brainstem. B, in the VGLUT3A224V/A224V mice, the collapse of the stereocilia impairs the mechanotransduction, degrading the hair cell receptor potential and most probably reducing the activity within the auditory nerve fibres. C, at later stage, the synaptic ribbons undergo a morphological change that may additionally downgrade the sound-stimulation encoding.
References


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Additional information

Data availability statement

The data that support the findings of this study are available from the corresponding authors upon reasonable request.

Competing interests

All authors declare no competing interests.

Author contributions


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Keywords

cochlea, deafness, exocytosis, sensory cells

Supporting information

Additional supporting information can be found online in the Supporting Information section at the end of the HTML view of the article. Supporting information files available:

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