Developmental refinement of hair cell synapses tightens the coupling of Ca^{2+} influx to exocytosis

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Abstract

Cochlear inner hair cells (IHCs) develop from pre-sensory pace-maker to sound transducer. Here, we report that this involves changes in structure and function of the ribbon synapses between IHCs and spiral ganglion neurons (SGNs) around hearing onset in mice. As synapses matured they changed from holding *several* small presynaptic active zones (AZs) and apposed postsynaptic densities (PSDs) to *one* large AZ/PSD complex per SGN bouton. After the onset of hearing (i) IHCs had fewer and larger ribbons; (ii) $\text{Ca}_v1.3$ channels formed stripe-like clusters rather than the smaller and round clusters at immature AZs; (iii) extrasynaptic $\text{Ca}_v1.3$ -channels were selectively reduced, (iv) the intrinsic Ca^{2+} dependence of fast exocytosis probed by Ca^{2+} uncaging remained unchanged but (v) the apparent Ca^{2+} dependence of exocytosis linearized, when assessed by progressive dihydropyridine block of Ca^{2+} influx. Biophysical modeling of exocytosis at mature and immature AZ topographies suggests that Ca^{2+} influx through an individual channel dominates the $[\text{Ca}^{2+}]$ driving exocytosis at each mature release site. We conclude that IHC synapses undergo major developmental refinements, resulting in tighter spatial coupling between Ca^{2+} influx and exocytosis.

Keywords active zone; Ca^{2+} channel; Ca^{2+} dependence; exocytosis; vesicle

Subject Categories Neuroscience

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Introduction

Inner hair cells (IHC) ribbon synapses are molecularly and morphologically specialized to transmit acoustic information at hundreds of Hz with sub-millisecond precision over long periods of time (recently reviewed in Matthews & Fuchs, 2010; Rutherford & Pangršič, 2012; Pangršič *et al.*, 2012; Safieddine *et al.*, 2012). The IHC receptor potential is coupled to glutamate exocytosis by Ca^{2+} influx through $\text{Ca}_v1.3$ Ca^{2+} channels (Platzter *et al.*, 2000; Brandt *et al.*, 2003; Dou *et al.*, 2004). Tens of these channels are thought to cluster within the presynaptic density (Brandt *et al.*, 2005; Frank *et al.*, 2010; Zampini *et al.*, 2013). One to two dozen synaptic vesicles are tethered at the plasma membrane within nanometer distance of the Ca^{2+} channel cluster and are thought to represent the putative readily releasable pool (RRP) of vesicles for exocytosis (Moser & Beutner, 2000; Zenisek *et al.*, 2000; Lenzi *et al.*, 2002; Rutherford & Roberts, 2006; Frank *et al.*, 2010). Precisely how Ca^{2+} channels couple to exocytosis and which mechanism of exocytosis governs transmitter release is not well understood.

Exocytosis at individual IHC AZs has been suggested to be multi-vesicular (Glowatzki & Fuchs, 2002), whereby Ca^{2+} is thought to regulate the rate of release events without affecting the number of vesicles released per event. Ca^{2+} -nanodomain coupling, in which exocytosis is controlled by very few nearby Ca^{2+} channels (Augustine *et al.*, 1991; Stanley, 1993; Mintz *et al.*, 1995; Wang *et al.*, 2008; reviewed in Augustine *et al.*, 2003; Moser *et al.*, 2006; Neher & Sakaba, 2008; Eggermann *et al.*, 2012), has been put forward to explain the physiological coupling of Ca^{2+} influx to RRP exocytosis

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in mature IHCs (Brandt *et al*, 2005; Goutman & Glowatzki, 2007; Zampini *et al*, 2013). A Ca^{2+} nanodomain may synchronize multivesicular release (Jarsky *et al*, 2010; Graydon *et al*, 2011). Alternatively, Ca^{2+} microdomain coupling (Heil & Neubauer, 2010), in which exocytosis is controlled by many channels cooperatively, and a linear intrinsic Ca^{2+} dependence of fusion (Johnson *et al*, 2010) have been suggested to explain the coupling of Ca^{2+} influx to RRP exocytosis in mature IHCs and these concepts, too, may coexist with multivesicular release.

Before the onset of hearing, exocytosis is evoked by Ca^{2+} action potentials (Beutner & Moser, 2001; Glowatzki & Fuchs, 2002; Johnson *et al*, 2005) with low “ Ca^{2+} efficiency” (Beutner & Moser, 2001; Johnson *et al*, 2005). Non-mutually exclusive hypotheses for the increase in Ca^{2+} efficiency of exocytosis over development include (i) progressive confinement of Ca^{2+} influx to the AZ, (ii) establishment of nanodomain coupling between $\text{Ca}_v1.3$ channels and release sites, and (iii) a change in intrinsic Ca^{2+} dependence of the molecular exocytosis machinery. Upregulation of synaptotagmin IV during postnatal development has been proposed to “linearize” the intrinsic Ca^{2+} dependence of IHC exocytosis around the onset of hearing (Johnson *et al*, 2010), supporting hypothesis iii. However, this contrasts with Ca^{2+} uncaging experiments that found a supralinear intrinsic Ca^{2+} dependence of exocytosis in mature IHCs (Beutner *et al*, 2001). Moreover, in immature IHCs, a thorough biophysical analysis of the intrinsic Ca^{2+} dependence of exocytosis or of the spatial coupling between Ca^{2+} influx and exocytosis was lacking until now.

Here, we combined morphological approaches: transmission electron microscopy (TEM), immunohistochemistry, confocal and stimulated emission depletion (STED) microscopy; with physiological techniques: electrophysiology, confocal Ca^{2+} imaging, Ca^{2+} uncaging; and biophysical modeling of AZ function to study the IHC afferent synapse during postnatal maturation. We find the intrinsic Ca^{2+} dependence of fast exocytosis to be similar in immature and mature IHCs, arguing against hypothesis iii. On the other hand, we find a diminution of extrasynaptic Ca^{2+} channels (supporting hypothesis i) and indicate a developmental switch from Ca^{2+} microdomain to Ca^{2+} -nanodomain control of vesicle fusion around hearing onset (supporting hypothesis ii).

Results

Postnatal maturation of AZ ultrastructure in IHC afferent synapses

Afferent synaptic contacts between IHCs and type I SGNs are present in mice already at birth (Shnerson *et al*, 1981) and increase in number during the first postnatal week (Shnerson *et al*, 1981; Sobkowicz *et al*, 1982; recently reviewed in Bulankina & Moser, 2012). Around the onset of hearing (p12, Mikaelian *et al*, 1965), synaptic contacts mature (Sobkowicz *et al*, 1982) and the number of presynaptic ribbons and postsynaptic AMPA receptor clusters decreases (Huang *et al*, 2007, 2012; Sendin *et al*, 2007). By p21, IHC-afferent synapses are considered predominantly mature morphologically (Bulankina & Moser, 2012; Safieddine *et al*, 2012) and functionally (Khimich *et al*, 2005; Grant *et al*, 2010; Wong *et al*, 2013). We first investigated the molecular and structural changes

within individual IHC-SGN contacts during the 2-week period around hearing onset.

We triple-labeled afferent IHC synapses in whole-mounts of the organ of Corti for (i) RIBEYE/CtBP2 (the major component of the synaptic ribbon (Schmitz *et al*, 2000; Khimich *et al*, 2005)), (ii) GluA2/3 (AMPA receptor subunits of the SGN PSD (Matsubara *et al*, 1996; Khimich *et al*, 2005)), and (iii) a bassoon (AZ scaffold contributing to ribbon anchorage (tom Dieck *et al*, 1998; Khimich *et al*, 2005)), or (iiib) the Na^+/K^+ -ATPase $\alpha 3$ subunit marking the SGN membrane (McLean *et al*, 2009). Immature synapses displayed several small appositions of AZs and PSDs (p6, Fig 1A). These AZ/PSD complexes often appeared to form circular structures $>1\ \mu\text{m}$ in diameter (Fig 1B), larger than the ring-like GluA clusters of mature synapses (p20, Fig 1C; up to approximately 700 nm, Meyer *et al*, 2009). Na^+/K^+ -ATPase labeling of SGN indicated that these circular structures localized to the periphery of the contact between each individual immature postsynaptic bouton and an IHC (Fig 1C,D, p6). Some of the AZ/PSD complexes lacked detectable RIBEYE immunofluorescence and likely represented ribbonless AZs (quantified in Supplementary Fig S2). After the onset of hearing each IHC-SGN synapse displayed a single AZ/PSD complex (p20, Fig 1).

TEM confirmed the presence of extended immature synaptic contacts that included ribbon-occupied and ribbonless AZ/PSD appositions (Fig 2A, p6 & p9). An exemplary synapse reconstructed from serial ultrathin sections (Fig 2D, p6) illustrates three ribbon-occupied AZs and one ribbonless AZ (black arrowhead). Compared with more mature ribbons (Fig 2B, p15 & p20), immature ribbons assumed more spherical or oval shapes (Fig 2C), were smaller (Fig 2E), associated with fewer vesicles (Fig 2E and tomogram in Supplementary Fig S1), and were juxtaposed to shorter PSDs (Fig 2E). After hearing onset, droplet- or wedge-shaped ribbons became prevalent (Fig 2C). Different ribbon shapes were observed in the same IHCs but we did not analyze their subcellular distribution. A typical p20 ribbon spanned four 70-nm-thin sections, compared to 1.5 sections on average for p6 ribbons. Floating ribbons tethering vesicles were observed near membrane-anchored ribbons at p6 (Fig 2A, green arrowhead). All preparations for TEM were fixed and processed at around noon time, minimizing possible effects of circadian changes (Hull *et al*, 2006).

Comparisons of confocal RIBEYE immunofluorescence between p6 and p20 revealed an approximate halving of ribbon number per IHC and an approximate doubling of the integrated pixel intensities per ribbon (Fig 2F,G and Supplementary Fig S2). An increase in full-width-at-half-maximum (FWHM, Fig 2F) of these RIBEYE spots is consistent with the increase in ribbon size observed in TEM. For GluA2/3, both the reduction in puncta number and the increase in integrated intensity per punctum were larger than for ribbons (Supplementary Fig S2C,E). We note that confocal measurements provide only a semi-quantitative comparison because immunolabeling may not be linear or equally efficient amongst puncta and because some ribbons or GluA2/3 puncta can be smaller than the resolution limit of confocal microscopy, especially in immature IHCs. Nevertheless, the confocal images indicate an increase in the ratio of ribbon and GluA2/3 puncta number from approximately 0.6 at p6 to nearly 1 for p20 IHCs (Supplementary Fig S2D), consistent with our observations of ribbonless AZs in immature IHCs and with the idea of a 1:1 correspondence between ribbons and receptor clusters per SGN bouton upon maturation (shown in Fig 1C,D, p20).

Figure 1. Developmental changes in morphology of hair cell afferent synapses.

- A Projection of confocal images showing GluA2/3 (green), RIBEYE (magenta) across the basal region of a row of IHCs at p6 (left) and p20 (right). Scale bar, 2 μ m.
- B Single confocal sections. Upper left: At p6, the basal pole of each IHC was contacted by several patches of postsynaptic glutamate receptors (GluA2/3, blue) forming circular structures (highlighted by dashed lines in upper right and lower left). Upper right: Each GluA2/3 spot was juxtaposed to an AZ spot, marked by protein bassoon (green) and in some cases by a ribbon (RIBEYE, red). Two ribbonless AZs marked by colocalizing bassoon and GluA2/3 puncta were highlighted by arrowheads. Individual fluorescence channels are presented separately in gray scale below. Scale bars, 2 μ m.
- C Schematized distribution of ribbon, bassoon, Ca^{2+} channels and GluA2/3-containing receptors on immature (p6, upper) and mature (p20, lower) IHC-SGN contacts. Immature IHC-SGN contacts hold multiple AZ/PSD complexes. Mature IHC-SGN contacts show a single and larger AZ/PSD complex.
- D Projection of confocal images showing GluA2/3 (blue), RIBEYE (red), and $\text{Na}^{+}/\text{K}^{+}$ -ATPase $\alpha 3$ subunit (green) at p6 (left) and p20 (right). The latter antibody labels the membranes of SGNs, each of which terminates onto the IHC with a bouton-type contact. In p6 preparations, the large circular structures formed by GluA spots partially encompass the perimeter of the bouton contacts. At p20, each postsynaptic bouton contains only a single condensed glutamate receptor cluster that is juxtaposed to a single presynaptic ribbon. Scale bars, 2 μ m.

Immature ribbons were anchored to the AZ membrane via up to two rootlets (presynaptic densities, Figs 2A and 3A,B). In contrast, a single continuous density attached the entire base of the mature ribbon to the AZ (Fig 3C,D). Using anti-bassoon immuno-EM (Fig 3F) we indicate that bassoon localizes to the presynaptic density underneath the ribbon, where the majority of immuno-gold was found (on average 1.4 ± 0.3 particles at the base and 0.3 ± 0.1 elsewhere at the ribbon, $P < 0.001$, $n = 21$ labeled synapses). This is consistent with a previous immuno-EM study on photoreceptor ribbon synapses (Dick *et al*, 2001) and also supported by present STED microscopy of hair cell synapses (see Fig 3G and below). The notion of ribbon-anchor consolidation was supported by the observation of

fewer electron-dense connections between membrane and ribbon in random sections of ribbon-occupied AZs from p6 to p20 (Supplementary Fig S3).

Next, we studied the topography of bassoon and $\text{Ca}_v1.3$ at the mature AZ, employing dual-color STED microscopy of immunolabeled IHCs in whole-mounts. Bassoon and $\text{Ca}_v1.3$ immunofluorescence often formed elongated stripes that aligned intimately with each other (Fig 3G). Most p19 synapses had one Ca^{2+} -channel stripe, some had more than one stripe in parallel, and others had round or more complex configurations (Fig 4B). Before the onset of hearing we observed only spot-like Ca^{2+} -channel clusters (Fig 4A). These spots are reminiscent of the small presynaptic densities seen

in immature AZs (Fig 2A), suggesting that Ca^{2+} channels cluster within the presynaptic density of immature IHCs, as they do in the stripe-like densities of mature IHCs (Frank *et al*, 2010). As depicted in Fig 3E, TEM analysis of membrane-proximal vesicles and presynaptic densities, and STED imaging of $\text{Ca}_v1.3$ channels suggest that synapse maturation leads to a more ordered topology of Ca^{2+} channels and membrane-proximal vesicles. In summary IHC-SGN synapses underwent major structural changes around the onset of hearing characterized by larger ribbons, continuously anchored to a single elongated presynaptic density, juxtaposed to a single postsynaptic receptor cluster.

Progressive confinement of $\text{Ca}_v1.3$ and Ca^{2+} influx to AZs over development

$\text{Ca}_v1.3$ channels conduct >90% of the voltage-gated Ca^{2+} influx in IHCs before and after the onset of hearing (Platzer *et al*, 2000; Brandt *et al*, 2003). Over development, we tracked changes in abundance and location of $\text{Ca}_v1.3$ with immunofluorescence and confocal Ca^{2+} imaging (Fig 5), in experiments run in parallel under identical conditions for the different age groups. Using $\text{Ca}_v1.3$ -knockout mice as a negative control we observed non-specific immunoreactivity only near the cuticular plate at the apical pole of p9 IHCs (Supplementary Fig S4), as previously shown for IHCs of hearing mice (Brandt *et al*, 2005).

$\text{Ca}_v1.3$ immunofluorescence near ribbons was detectable already at p6 (arrowheads in Fig 5A) and increased until p14 (mean intensity around ribbons at p6: 6494 ± 366 a.u. versus p14: 7721 ± 265 a.u., $P = 0.007$). Thereafter, we did not find significant changes of synaptic $\text{Ca}_v1.3$ immunofluorescent spots except for a subtle increase in the long axis (p14: 0.336 ± 0.004 versus p20: 0.350 ± 0.003 μm , $P = 0.005$). Consistent with a previous report (Zampini *et al*, 2010) we also noted $\text{Ca}_v1.3$ immunofluorescence away from ribbons, appearing spot-like or diffuse along the entire basolateral plasma membrane of immature IHCs. Upon the onset of hearing, $\text{Ca}_v1.3$ immunoreactivity became largely confined to ribbon-occupied AZs (Fig 5A, lower panels). We quantified the density of $\text{Ca}_v1.3$ immunofluorescence immediately around ribbons (synaptic) versus distant from ribbons (extrasynaptic, see Materials and Methods). At p6, the density of extrasynaptic fluorescence was around 43% of that in synaptic regions, while by p20, it had decreased to only 9%.

Next, we studied changes in intracellular Ca^{2+} concentration ($[\text{Ca}^{2+}]_i$) elicited by step depolarizations (to -7 mV for 200 or 254 ms) in IHCs using fast confocal Ca^{2+} imaging before and after the onset of hearing (Fig 5B). To preferentially visualize sites of Ca^{2+} influx, we used the low-affinity Ca^{2+} indicator Fluo-5N (0.4 mM) in conjunction with the slow Ca^{2+} chelator EGTA (2 mM; Frank *et al*, 2009), and a fluorescently tagged RIBEYE-binding peptide (Francis *et al*, 2011) for identification of ribbon-occupied AZs. Different from the synapse-confined Ca^{2+} signals of mature IHCs (Fig 5B, p16), in immature IHCs we found a spatially extended rise of $[\text{Ca}^{2+}]_i$ (Fig 5B, p10) consistent with substantial extrasynaptic Ca^{2+} influx and with immunohistochemistry (Fig 5A), and also compatible with potential Ca^{2+} -induced Ca^{2+} release (Kennedy & Meech, 2002). Occasionally (two out of seven IHCs) we found wide-spread “global” Ca^{2+} signals (two uppermost rows in Fig 5B, and Supplementary Fig S5). In the other five immature IHCs the peak Ca^{2+} signal was “spot-like”

around ribbons (third row in Fig 5B, and Supplementary Fig S5), further suggesting that $\text{Ca}_v1.3$ clusters at AZs already before the onset of hearing. We approximated the strength of the extrasynaptic Ca^{2+} signal as the ratio of ΔF away from the ribbon over ΔF at the ribbon, which was greatest in immature IHCs with “global” signals, smaller in immature IHCs with “spot-like” signals, and even smaller in IHCs after the onset of hearing (Supplementary Fig S5).

The intense, broadly distributed $\text{Ca}_v1.3$ immunofluorescence in immature IHCs suggests that the reduction of the whole-cell Ca^{2+} current around the onset of hearing reflects a reduction in the number of Ca^{2+} channels, primarily in the extrasynaptic membrane. We studied the IHC Ca^{2+} current before and after the onset of hearing using whole-cell patch-clamp recordings. We did not detect significant differences in the voltage dependence of activation (Supplementary Fig S6A,B) or in the extent or kinetics of inactivation (Supplementary Fig S6C) between p10 and p14 IHCs. Non-stationary fluctuation analysis of Ca^{2+} -tail currents (Roberts *et al*, 1990; Frank *et al*, 2010) indicated that the smaller number of Ca^{2+} channels almost completely accounted for the reduction of the whole-cell current ($P < 0.001$, ANOVA, Supplementary Fig S6D,E). The single channel current (i_{Ca}) estimate did not change significantly ($P = 0.1$, ANOVA, Supplementary Fig S6F) and the estimated maximal open probability ($P_{\text{open,max}}$) decreased slightly ($P < 0.001$, ANOVA; Supplementary Fig S6F), whereas single channel recordings indicated a reduction in i_{Ca} and an increase in P_{open} over development (Zampini *et al*, 2010, 2013). While both approaches agree on the notion of much higher total Ca^{2+} channel numbers in immature cells, they differ in their estimates of i_{Ca} and P_{open} and also report opposite trends over development likely for technical reasons (see text accompanying Supplementary Fig S6 for discussion). In summary, immature IHCs contain thousands of synaptic and extrasynaptic $\text{Ca}_v1.3$ channels that together support presensory Ca^{2+} action potentials. Upon maturation extrasynaptic $\text{Ca}_v1.3$ Ca^{2+} channels are downregulated and Ca^{2+} influx becomes confined to AZs, supporting hypothesis 1 in introduction.

Maturation of the coupling between Ca^{2+} influx and exocytosis

The developmental confinement of Ca^{2+} influx to AZs together with the changes in ultrastructure and molecular nanoanatomy of the AZ prompted us to test for parallel changes in the coupling of Ca^{2+} influx to exocytosis. In particular, we tested hypothesis ii: tighter functional coupling between Ca^{2+} channels and the RRP, and hypothesis iii: a change in the intrinsic (biochemical) Ca^{2+} dependence of the molecular exocytosis machinery.

No change in the intrinsic Ca^{2+} dependence of fast exocytosis

First, we addressed the intrinsic Ca^{2+} dependence of IHC exocytosis before the onset of hearing, which had not been done previously. UV-laser photolysis of caged Ca^{2+} was used to elevate $[\text{Ca}^{2+}]$ homogeneously throughout the cytosol in p6–8 IHCs and in p14–18 IHCs for comparison. $[\text{Ca}^{2+}]$ was monitored with ratiometric imaging and related to the resulting exocytic increase of membrane capacitance (ΔC_m). Approximately half (22 out of 40) of p6–8 IHCs and nearly all (32 out of 33) of p14–18 IHCs responded with a C_m rise that comprised two kinetic components with time constants of a few milliseconds and tens of milliseconds, respectively (Fig 6A,C). The total

amplitude of the C_m rise was largely independent of $[Ca^{2+}]_i$ at both developmental stages (Fig 6D), as previously reported for p14–25 IHCs (Beutner *et al*, 2001). However, it was approximately 3.6 times smaller in p6–8 IHCs than in p14–18 IHCs (324 ± 24 fF; $n = 40$ versus 1161 ± 104 fF; $n = 31$; $P = 1.5 \times 10^{-13}$, Wilcoxon Rank Test), potentially relating to the lower number of synaptic vesicles found at the AZs of p6–7 IHCs (Fig 2E). Moreover, when both kinetic components were present in p6–8 IHCs we found a greater contribution of the slow one (69%), while the two components contributed

equally in p14–18 IHCs (Supplementary Fig S7). In the other p6–8 IHCs the C_m rise was best approximated by a single exponential function with relatively slow time constants within the range of those of the slow component of the C_m rise in IHCs with bi-exponential responses (Fig 6C). Focusing on the fast component of exocytosis, we probed the intrinsic Ca^{2+} dependence of exocytosis by measuring the delay and rate constant of the C_m rise for a range of $[Ca^{2+}]_i$ (Fig 6C,E). We found a similar supralinear intrinsic Ca^{2+} dependence in p6–8 and p14–18 IHCs, indicating that the Ca^{2+} -

Figure 2. Maturation of ribbon synapse ultrastructure.

- A, B Representative electron micrographs of IHC ribbon synapses in pre-hearing (A) and hearing (B) mice. Before the onset of hearing (A), at p6 and p9, multiple appositions of small discontinuous pre- and postsynaptic densities were found (magenta arrowheads) typically accompanied by one or more round-shaped ribbon(s) anchored close to the presynaptic membrane via two rootlets. In some cases ribbons were floating in the cytoplasm at a greater distance from the membrane (green arrowhead). After the onset of hearing (B), at p15 and p20, typically one ribbon occupied the presynaptic cytoplasm at the AZ. In comparison to before the onset of hearing, the mature pre- and postsynaptic densities were relatively extended and continuous. The sections were cut parallel to the ribbons short axes, the perspective from which ribbon-shape was measured. Scale bar of 200 nm in (A) applies also to (B).
- C Through development the ribbon-shape changed from predominantly round to oval, droplet-like, or wedge-like shapes, as shown in (A) and (B).
- D 3D reconstructions from serial sections before (left, p6) and after (right, p20) the onset of hearing. Presynaptic densities connect the ribbon to the plasma membrane.
- E Quantification of synaptic parameters from random EM sections. Number of ribbon-associated vesicles, size of synaptic ribbon as well as the extent of postsynaptic density (PSD) increase with age (student's *t*-test). Only sections parallel to the ribbons' short axes with sharply delineated PSD were analyzed. Ribbon size was measured as the length of the longest extent across the synaptic ribbon.
- F Increase of ribbon size/content during development was corroborated by the increase in integrated puncta intensity from quantification of RIBEYE immunofluorescence. A developmental increase in full-width-of-half-maximum (FWHM) of Gaussian fits to puncta was also observed, Wilcoxon rank test.
- G Decrease of number of synaptic ribbon during development estimated by immunohistochemistry (immuno) and live imaging (peptide), student's *t*-test.

Data information: * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$. n.s., no significant difference.

binding properties of the molecule(s) mediating fast exocytosis do not change upon the onset of hearing. Because of the steep Ca^{2+} dependence we applied statistical comparison among p6–8 and p14–17 IHCs for the rate constants of the fast component and the exocytic delays within a narrow $[\text{Ca}^{2+}]$ range (15–25 μM , for which we found the best comparable representation of $[\text{Ca}^{2+}]$ changes). Neither the rate constants ($P = 0.7$, Student's *t*-test) nor the delays ($P = 0.86$, Student's *t*-test) were significantly different. The rate constants were larger and the delays shorter for the fast component in the present study (Fig 6C,E) than those observed previously at comparable $[\text{Ca}^{2+}]$, which we attribute to the use of shorter stimuli (100 μs laser stimulation versus 1 ms long arc lamp flashes) that achieved a faster rise of $[\text{Ca}^{2+}]$.

Linearization of the apparent Ca^{2+} dependence of exocytosis is limited to changes in the number of open Ca^{2+} channels

Next, we addressed the apparent Ca^{2+} dependence of exocytosis (Matveev *et al*, 2011) by studying the coupling between Ca^{2+} influx and RRP exocytosis in p6, p9–10, and p14–17 IHCs. We used strong (–17 mV) and brief (20 ms) depolarizations to evoke exocytosis of the RRP (Moser & Beutner, 2000; Beutner & Moser, 2001) in perforated-patch recordings. Following the framework of our previous analysis (Brandt *et al*, 2005) we studied the relationship between RRP exocytosis ($\Delta C_{m,20\text{ ms}}$) and the integrated Ca^{2+} influx (Q_{Ca}) while either varying the single Ca^{2+} channel current (i_{Ca}) or changing the number of open $\text{Ca}_v1.3$ channels ($N_{\text{Ca}} \times P_{\text{open}}$). The apparent Ca^{2+} cooperativity m was obtained by fitting the exocytosis- Q_{Ca} relationship for each cell with a power function: $\Delta C_m = A(Q_{\text{Ca}})^m$ (Augustine *et al*, 1991), limited to the Q_{Ca} range in which no obvious saturation of the exocytosis was observed. For Ca^{2+} -microdomain control (see Introduction), m should be similar to the intrinsic Ca^{2+} cooperativity (4–5, Beutner *et al*, 2001), regardless of how the Ca^{2+} influx was manipulated. For Ca^{2+} -nanodomain control (see Introduction), the m should differ between the two manipulations, because there, the number of released RRP vesicles is linearly determined by $N_{\text{Ca}} \times P_{\text{open}}$ ($m \sim 1$, Augustine *et al*, 1991, see our biophysical model below).

To vary i_{Ca} , we slowly changed the extracellular $[\text{Ca}^{2+}]$ (Fig 7A,C) in both directions while supplementing divalent ions (Mg^{2+}) at low Ca^{2+} concentration ($< 1.3\text{ mM}$ $[\text{Ca}^{2+}]_e$). We observed a supralinear exocytosis- Q_{Ca} relationship in the range of low Q_{Ca} (Fig 7C, average

of the maximal Q_{Ca} used for fitting indicated by dashed vertical lines) before and after the onset of hearing (supralinear apparent Ca^{2+} dependence, p6–8: $m = 3.10 \pm 0.39$, $n = 6$ IHCs; p14–17: $m = 2.91 \pm 0.38$, $n = 7$ IHCs). Next, to gradually reduce $N_{\text{Ca}} \times P_{\text{open}}$ (Fig 7D) while maintaining i_{Ca} (Hess *et al*, 1984), we slowly bath-perfused preparations with the antagonistic dihydropyridine isradipine (10 μM). In immature IHCs, we found a supralinear apparent Ca^{2+} dependence ($m = 2.35 \pm 0.18$, $n = 7$ IHCs) consistent with control of any given release site by several channels (“ Ca^{2+} microdomain-like” control of exocytosis). The exponent m declined with maturation to a quasi-linear apparent Ca^{2+} dependence after the onset of hearing (p14–p17: 1.42 ± 0.13 ; $n = 7$ IHCs, see also Supplementary Fig S8 for corroborating data), suggesting a change towards “ Ca^{2+} nanodomain-like” control of exocytosis.

Linearization of the apparent Ca^{2+} dependence of exocytosis for $N_{\text{Ca}} \times P_{\text{open}}$ changes indicates tightening of Ca^{2+} influx-exocytosis coupling: insights from modeling

How precisely Ca^{2+} influx through nearby Ca^{2+} channels regulates the exocytosis of a given RRP vesicle depends on several parameters, including the number, distance and open probability of Ca^{2+} channels, the concentration and binding kinetics of Ca^{2+} buffers and the Ca^{2+} -binding properties of the Ca^{2+} sensor of exocytosis (Moser *et al*, 2006; Matveev *et al*, 2011). Given this complexity we turned to biophysical modeling in order to reconcile the topography of membrane-proximal vesicles (here assumed to represent the RRP) and Ca^{2+} channels at the AZ with the apparent Ca^{2+} dependence of exocytosis observed in IHCs during changes of the number of open channels ($N_{\text{Ca}} \times P_{\text{open}}$) or i_{Ca} before and after the onset of hearing. We based the models on morphological and biophysical information (e.g. data from EM, STED, Ca^{2+} channel analysis, Ca^{2+} uncaging of this study and published work: for parameters and computational methods see Supplementary Material S9). We then studied paradigmatic topographies of Ca^{2+} channels and Ca^{2+} sensors for mature AZ (scenarios “M1–3”, Fig 8A and Supplementary Material S9) and immature AZs (scenarios “IM”, Fig 8B and Supplementary Material S9).

For the mature AZ, 14 vesicles (40-nm-diameter red disks in Fig 8A,B) were randomly distributed at the longer edges of an $80 \times 420\text{ nm}$ presynaptic density (seven vesicles per one side), with

Figure 3. Nanoanatomy of IHC AZs.

- A TEM of an immature IHC ribbon synapse using high-pressure freezing and freeze substitution (HPF/FS).
- B Corresponding serial 3D reconstruction of (A), showing the immature ribbon connected via two rootlets to the presynaptic membrane. For color code see (D).
- C Representative TEM of a mature ribbon, connected with one continuous presynaptic density to the membrane, as demonstrated in the 3D reconstruction in (D).
- D 3D reconstruction: ribbon: red; AZ membrane: blue; rootlet (immature)/presynaptic density (mature): magenta; membrane-proximal SVs: orange; ribbon-associated SVs: yellow.
- E Schematics of immature (top) and mature (bottom) AZs inspired by exemplary 3D reconstructions and STED imaging of $Ca_v1.3$ immunofluorescence. Ca^{2+} channels formed small clusters and vesicles were less organized at the immature AZ. Stripe-like Ca^{2+} -channel cluster with vesicles positioned regularly along each side at the mature AZ.
- F Pre-embedding immunogold labeling for bassoon (arrowhead) was detected at the ribbon-anchoring presynaptic density of mature ribbons. Two primary antibodies specific to the N-terminal or C-terminal of bassoon, respectively, were used (see Materials and Methods). Scale bar, 200 nm.
- G STED image of immunolabeled bassoon (magenta) and $Ca_v1.3$ channel clusters (green) in mature IHCs: stripe-like morphology and closely aligned immunofluorescence of bassoon and $Ca_v1.3$ reminiscent of the presynaptic density reconstructed from TEM (D). The image is a maximum intensity projection after smoothing with a Gaussian ($\sigma = 1$ pixel) in ImageJ. The color lookup was chosen to optimize contrast and visibility of dimmer structures. Pixel size, 20 nm. Scale bar, 2 μ m. Insets, 700 $\times$ 700 nm.

Figure 4. Development of Ca_v1.3 cluster morphology.

- A** Projections of STED sections of immunolabeled Ca_v1.3 clusters in the organ of Corti whole-mount at p10 (0.6 μ m step size, total z covered: 4 μ m). Ca_v1.3 clusters in pre-hearing IHCs appeared spot-like (A), in contrast to the stripe-like appearance in mature IHCs (B). Images were processed as in Fig 3G. Pixel size, 20 nm, 700 $\times$ 700 nm regions-of-interest centered at individual Ca_v1.3 clusters.
- B** Projections of STED sections of immunolabeled Ca_v1.3 clusters in the organ of Corti whole-mount at p19 (0.6 μ m step size, total z covered: 4 μ m). Ca_v1.3 clusters in pre-hearing IHCs appeared spot-like (A), in contrast to the stripe-like appearance in mature IHCs (B). Images were processed as in Fig 3G. Pixel size, 20 nm, 700 $\times$ 700 nm regions-of-interest centered at individual Ca_v1.3 clusters.

the vesicular Ca²⁺ sensor (black spots in Fig 8A,B) in contact with the density in the plane of the plasma membrane. The size of the presynaptic density (grey area in Fig 8A,B) was chosen based on exemplary synapses reconstructed from serial EM sections (e.g. Fig 3D), and is consistent with Ca_v1.3 immunofluorescence in STED microscopy (Figs 4G and 5B). We placed 14–90 channels within the presynaptic density, representing different scenarios of channel-vesicle coupling. In scenario M1 (Fig 8A, upper panel), 36 channels were distributed randomly in the presynaptic density area. In scenario M2 (Fig 8A, middle panel), in addition to the 36 randomly distributed ones, 14 channels were placed in physical contact with the vesicle Ca²⁺ sensors, simulating molecular coupling (review in Eggermann *et al*, 2012) and nicknamed hereafter as “private channel” to highlight its dominance on the Ca²⁺ signal of a given vesicle. In scenario M3, only those 14 channels in physical contact with the vesicles were implemented (Fig 8A, lower panel). For each scenario, 100 different realizations of random channel and vesicle positions were considered. Analogous to experimental analysis, we compared the slope m of vesicle release against Ca²⁺ influx in double-logarithmic plots among scenarios of different degree of coupling. Scenarios M1–3 resulted in different exponents m for $N_{Ca} \times P_{open}$ change, all below 2: 1.8 for M1, 1.2 for M2 and 1.1 for M3 (Fig 8E), while m estimates for i_{Ca} change were all approximately 4 (Fig 8F). As another measure of Ca²⁺ influx to exocytosis coupling, we studied

the effective number of contributing channels n_{ch} (Fig 8D for scenario M2), which is the ratio of the total mean steady state [Ca²⁺] (Fig 8C) to the mean steady state [Ca²⁺] elicited by the channel that contributed the most. With increasing dominance of the nearest channel, the effective number n_{ch} of channels contributing to [Ca²⁺] at the Ca²⁺ sensor went down from 4.4 for M1 to 2.1 for M2 and 1.2 for M3. Specifically, $n_{ch} = 2.3$ (1.2) in scenario M2 (M3) indicates that each “private channel” on average contributes 44% (83%) of [Ca²⁺] at the corresponding Ca²⁺ sensor, whereas $n_{ch} = 4.4$ in scenario M1 implies a contribution of only 23% by the nearest channel. Figure 8D illustrates for the case of scenario M2 how n_{ch} depends on the spatial position of channels and Ca²⁺ sensors in the plane of the plasma membrane. Simply moving the sensor 20 nm away from the “private channel” towards the vesicle center in M2 (scenario M2d, Supplementary Figs S9 and S14), increased n_{ch} to 5.8 and m to 2.5 for $N_{Ca} \times P_{open}$ changes, highlighting the strong effect of spatial proximity for the relevance of the nearest channel in controlling release of a given vesicle. In conclusion, modeling confirmed that the experimentally tractable Ca²⁺ cooperativity m during $N_{Ca} \times P_{open}$ changes is diagnostic for the relevance of the nearest channel in controlling fusion and support the notion of a Ca²⁺-nanodomain control of exocytosis in mature IHCs.

Despite the fact that implementing molecular coupling (M2–3) was sufficient to reproduce m near unity during $N_{Ca} \times P_{open}$ changes, this may not be the case if the channel density at the AZ is very high. Indeed, when we placed 40 additional randomly distributed Ca²⁺ channels in scenario M2 (a total of 90 channels, scenario M2b in Supplementary Fig S9), m increased from 1.2 to 2.0 and n_{ch} from 2.3 to 3.5. One possibility to preserve the dominance of the “private channels” is to implement an exclusion zone around them where no other channels can be located. Scenario M2c (Supplementary Figs S9 and S14, same density of channels outside the exclusion zone as in M2b) assumed circular exclusion zones with the width of a channel diameter and resulted in estimates of $m = 1.2$ and $n_{ch} = 2.1$ (very similar to those of scenario M2). On the other hand, if the density of the channels in the presynaptic density was low, m near unity could be reproduced even with two “private channels”. For example, addition of one more “private channel” per vesicle in M3 (scenario M3b in Supplementary Figs S9 and S14) led to $m = 1.2$ ($N_{Ca} \times P_{open}$ change) and $n_{ch} = 2.7$, close to the values obtained for scenario M2. Placing additional Ca²⁺ channels outside the presynaptic density (10/ μ m², likely a strong overestimate of the average extrasynaptic density, see Brandt *et al*, 2005; Zampini *et al*, 2013) had negligible impact. In all above-mentioned scenarios m stayed high, approximately 4, when manipulating i_{Ca} .

Modelling the apparent Ca²⁺ dependence of immature IHCs based on AZ topography is more challenging because (i) of the large fraction of extrasynaptic Ca²⁺ channels (possibly driving extrasynaptic exocytosis) and (ii) the fact that half of the IHCs lacked the fast component of exocytosis elicited by Ca²⁺ uncaging (which was used for modelling). However, higher m predictions for $N_{Ca} \times P_{open}$ changes than found for the mature scenarios M2–3 were readily obtained with models based on realistic AZ topography (IM1), such as that depicted in Fig 8B. There, a presynaptic density area similar to that of M1–3 was distributed within 1 μ m diameter disk in the form of separate patches corresponding to either pairs of smaller or single bigger densities (consistent with EM reconstructions), wherein 60 Ca²⁺ channels were distributed with similar average

Figure 5. Maturation of hair cell Ca^{2+} signaling.

- A Organs of Corti immunolabeled for $\text{Ca}_v1.3$ (green) and RIBEYE/CtBP2 (magenta). At p6 (top), a single section shows both ribbon-occupied (arrowheads) and ribbonless $\text{Ca}_v1.3$ spots as well as diffuse membrane staining. At p11 (middle, projection) puncta of $\text{Ca}_v1.3$ were observed throughout the basolateral membrane, especially the basal pole. By p14 the clusters of $\text{Ca}_v1.3$ were concentrated in the basal poles near synaptic ribbons, while the entire inner hair cell membrane remained diffusely labeled for $\text{Ca}_v1.3$. At p20 there was very little extrasynaptic labeling and the cell membrane was no longer diffusely immunoreactive for $\text{Ca}_v1.3$. Each synaptic ribbon was accompanied by a cluster of $\text{Ca}_v1.3$ and only few $\text{Ca}_v1.3$ clusters remained ribbonless. Scale bars, 10 μm .
- B Examples of XY scans across the basolateral portion of IHCs before (upper, p10) and after (lower, p14) the onset of hearing. Left and right panels show the increase in Fluo-5N fluorescence during a step depolarization to -7 mV for 254 ms (2 mM [EGTA]_i) and peptide-labeled synaptic ribbons (position marked with arrowheads). Note the relatively widespread submembraneous Ca^{2+} signal in immature IHCs as compared to the local Ca^{2+} rise in mature IHCs. Still, the highest [Ca^{2+}] was usually observed at the ribbons-occupied AZs. Scale bar, 3 μm .

density as in M2. Scenario IM1 did not force physical contact of Ca^{2+} channel and vesicular Ca^{2+} sensor, which was placed in shortest distance to the nearest presynaptic density. In addition, 60 channels were distributed randomly outside the presynaptic density patches. The predicted m ($N_{\text{Ca}} \times P_{\text{open}}$ change) was 2.1, which is similar to the experimental observation (2.3) and considerably higher than the m ($N_{\text{Ca}} \times P_{\text{open}}$ change) estimates of M2 or M3 (Fig 8E,F). On the other hand, m (i_{Ca} change) was 4.1, which is very similar to those of M1–3: $m = 3.9$ –4.1 (Fig 8F). However, m considerably decreased for $N_{\text{Ca}} \times P_{\text{open}}$ ($m = 1.1$) but not for i_{Ca} changes ($m = 4.0$) when implementing “private channels” as in scenarios M2 and M3 (scenario IM1b, Supplementary Material S9, Supplementary Figs S9 and S14).

In our model, we realized that RRP depletion (fusion exceeding vesicle replenishment, see Supplementary Material S9) could cause exocytosis to saturate with increasing Ca^{2+} influx depending on the model parameters (Figs 8D,E and Supplementary Fig S10). Saturation would lead to underestimation of m if data of this Q_{Ca} range was included into the fit. Release leveled-off more in the case of i_{Ca} variation, where, different from $N_{\text{Ca}} \times P_{\text{open}}$ changes (Fig 7D, see also Supplementary Fig S8), saturation was also observed experimentally and fitting was restricted to the non-saturating range (Fig 7C). Our treatment of this “release saturation” is described in Supplementary Material S9. Using simulations, we also explored the Q_{Ca} range in which the number of open channels per cluster is very low, which is difficult to achieve experimentally because of the

Figure 6. Intrinsic Ca^{2+} dependence of exocytosis in pre-hearing IHCs.

A Representative examples of UV-laser evoked capacitance rise (ΔC_m) of IHCs from mice of the specified ages. Upper: Intracellular $[Ca^{2+}]_i$ of a p14 (black), a p7 (magenta) and a p6 (blue) IHCs after UV-flash, measured ratiometrically with mag-fura-2. Lower: ΔC_m of the corresponding IHCs (solid lines). Lighter dashed lines represent bi-exponential fits to the data and lighter solid and dotted lines represent the fast and slow components respectively.

B The average ($\pm$ SEM) ΔC_m of p14–18 (black) and p6–8 IHCs (magenta: all cells; grey: only cells with bi-exponential ΔC_m).

C Rate constants from exponential fits to the ΔC_m upon flash photolysis plotted against post-flash $[Ca^{2+}]_i$. Circles mark the rate constants for first (fast) component and rhombi mark those for the second (slow) component of the bi-exponential fits. Rate constants of responses in p6–8 IHCs, which were best fitted with a single exponential are also marked with rhombi. Note the overlap between pre-hearing (p6–8) and hearing (p14+) age groups. Black (p14–18) and magenta (p6–8) symbols are newly acquired data, while open circles represent previously published data on mature IHCs (Beutner et al, 2001; Pangršič et al, 2010).

D Total response amplitudes (sum of first and second component, in case of double exponential fits) were largely independent of post-flash $[Ca^{2+}]_i$.

E Delay of responses plotted against post-flash $[Ca^{2+}]_i$.

Data information: Solid lines in (C) and (E) are model fits from a previous publication (Beutner et al, 2001).

approximately 20% residual current with isradipine block. As expected, m approached unity regardless of the topography, because release becomes dependent on Ca^{2+} influx through single openings

of the few available channels. In contrast, for i_{Ca} changes m values remained near constant even at considerably lowered Q_{Ca} outside the saturation region. In conclusion, biophysical modeling supports

Figure 7. Maturation of IHCs Ca^{2+} influx-exocytosis coupling: experimental data.

A Depolarizations (20 ms to -17 mV) elicited inward Ca^{2+} current (I_{Ca} , lower panel) and triggered ΔC_m (upper panel) in a representative pre-hearing (p7) IHC. Increasing $[\text{Ca}^{2+}]_e$ enlarged both I_{Ca} and ΔC_m .
 B Time course for ΔC_m and integrated Ca^{2+} influx (Q_{Ca}) of the same IHC in (A) for repetitive depolarizations (20 ms to -17 mV) at 60 s interval. Experiment started at 5 mM $[\text{Ca}^{2+}]_e$ and periods where bath solution was slowly perfused with a different $[\text{Ca}^{2+}]_e$ are marked by horizontal bars.
 C Plotting ΔC_m , 20 ms against Q_{Ca} reveals a supralinear apparent Ca^{2+} dependence of exocytosis for low Q_{Ca} . Solid lines represent best fit power functions ($\Delta C_m = A(Q_{Ca})^m$) to the whole data set in this range. Each small symbol represents an individual response to depolarization (p15-16, grey, $n = 7$ IHCs; p6, magenta, $n = 7$ IHCs). Dashed vertical lines indicate the average maximal Q_{Ca} limiting the Q_{Ca} range used for fitting.
 D Decreasing apparent Ca^{2+} cooperativity of exocytosis through blocking L-type Ca^{2+} channels with isradipine. Each small symbol represents an individual response to depolarization (p14-15, grey, $n = 6$ IHCs; p6-8, magenta, $n = 7$ IHCs).

Data information: In (C) and (D), solid lines show best fit power functions to the whole data set. Average values of the exponent m are indicated on top.

the experimentally derived notion of a maturational tightening of Ca^{2+} channel-exocytosis coupling.

Discussion

Here we studied the maturation of afferent IHC synapses around the onset of hearing. We found a transformation from an immature synaptic contact containing multiple small and peripheral spot-like AZ/PSD complexes into a consolidated contact with a single large AZ/PSD complex per SGN bouton. Ca^{2+} channels and Ca^{2+} influx became progressively confined to synaptic AZs over development. $\text{Ca}_v1.3$ channels first formed spot-like clusters whereas, following AZ remodeling, they formed an extended stripe alongside bassoon protein within the presynaptic density. Synaptic ribbons were larger in size and more flat in shape after the onset of hearing. Ca^{2+} uncaging triggered a fast component of exocytosis in only half of the immature IHCs, where it was smaller than in mature IHCs but had a

similar intrinsic Ca^{2+} dependence. A “linearization” of the apparent Ca^{2+} dependence of exocytosis observed for $N_{Ca} \times P_{open}$ changes, together with computational modeling of immature and mature AZ topographies, indicated a developmental switch from more Ca^{2+} -microdomain-like to more Ca^{2+} -nanodomain-like coupling between $\text{Ca}_v1.3$ channels and vesicular release sites.

Maturation of molecular nanoanatomy of IHC afferent synapses

High-resolution immunofluorescence and electron microscopy jointly demonstrated major structural changes of IHC synapses around the onset of hearing, schematized in Figs 1C and 3E. We propose that this change to a more ordered stripe-like AZ topography, together with the dramatic reduction of extrasynaptic $\text{Ca}_v1.3$ channels, enhances the Ca^{2+} efficiency of exocytosis and tightens the spatial coupling between $\text{Ca}_v1.3$ channels and vesicular release sites (Figs 7 and 8). We consider two not mutually exclusive candidate mechanisms for the structural refinement of the synapse over

Figure 8. Maturation of IHCs Ca^{2+} influx-exocytosis coupling: modeling.

- A** Scenarios M1–3 of the mature AZ topography. Grey area: presynaptic density, red discs: RRP vesicles, green discs: Ca^{2+} channels, black spots: Ca^{2+} sensors of exocytosis. Scenario M1: random positioning of channels within the presynaptic density, Ca^{2+} sensor pointing towards the channel cluster. Scenario M2: introduction of private channels simulating molecular coupling of release site and channel. Scenario M3: only private channels, unrealistically low number of channels. Plots correspond to one out of a hundred realizations considered in the simulations.
- B** Scenario IM1 of the immature AZ topography. Legend as in (A). Note the more extended synaptic contact with several Ca^{2+} channel clusters, which were placed at the sites of presynaptic densities.
- C** Total mean steady state $[Ca^{2+}]$ profile at the AZ membrane for a particular realization of scenario M2.
- D** Effective number of channels contributing to total mean steady state $[Ca^{2+}]$ shown in (C).
- E** Average overall release (number of vesicles released) versus Ca^{2+} charge influx per synapse after the first 20 ms after the stimulus onset when $N_{Ca} \times P_{open}$ was changed by blocking channels (modeling the isradipine experiment). Dots correspond to the values averaged over different synapses and realizations as described in Supplementary Material S9. Straight lines are linear fits in log-log scale within the interval of fivefold decrease in Q_{Ca} . Numbers correspond to the slope of fits, representing the apparent Ca^{2+} cooperativity of release, m . Blue—mature scenario M1, black—mature scenario M2, green—mature scenario M3, magenta—immature scenario IM1.
- F** The same as (E) but Q_{Ca} was varied by manipulating i_{Ca} instead of $N_{Ca} \times P_{open}$.

development: (i) merging of the individual synaptic specializations within a contact and (ii) pruning of all but one AZ/PSD complex per contact.

Merging may be mediated by interactions of pre- and/or post-synaptic scaffold molecules and could involve trans-synaptic regulation. Such cluster-cluster interactions may join multiple small ribbons, Ca^{2+} -channel clusters, or GluA clusters into single large structures. Several lines of evidence indicate an important role of bassoon in this process. First, the involvement of cytomatrix of the AZ proteins in synaptogenesis was demonstrated for other synapses (Zhai *et al*, 2000, 2001; Shapira *et al*, 2003) whereby bassoon also recruits other proteins to the AZ (Shapira *et al*, 2003; Takao-Rikitsu *et al*, 2004; Maas *et al*, 2012). Second, bassoon resides in the stripe-like presynaptic density of the mature IHC synapse of wild-type mice (Fig 3) and also in the immature presynaptic density (Fig 1). Third, while synapses are still formed in IHCs lacking functional bassoon (Khimich *et al*, 2005), the presynaptic density and Ca^{2+} -channel clusters remain spot-like (Frank *et al*, 2010). Notably, around hearing onset bassoon mRNA is upregulated in the mouse organ of Corti under control of thyroid hormone (Sendin *et al*, 2007). Perhaps bassoon facilitates AZ merging via interactions with CAST (Takao-Rikitsu *et al*, 2004), RIBEYE (tom Dieck *et al*, 2005), piccolo (Wang *et al*, 2009), and other proteins. The fusion of neighboring AZ-anchored ribbons is plausible given the strong RIBEYE-RIBEYE interactions (Magupalli *et al*, 2008), reports of ribbon-ribbon fusion during retinal synaptogenesis (Regus-Leidig *et al*, 2009), their stimulus-dependent structural plasticity (Spiwoks-Becker *et al*, 2004), and our observation of floating ribbons near anchored ones at immature IHC AZs (Fig 2).

Alternatively, or in parallel with merging, all but one of the AZs may be degraded. Pruning is a well-established mechanism for synaptic refinement in general (Goda & Davis, 2003). In the organ of Corti, pruning is thought to refine SGN afferent connectivity at the level of fibers and contacts (Sobkowicz *et al*, 1982; Huang *et al*, 2007, 2012; Sendin *et al*, 2007) but its role in development within a synaptic contact is not clear. Molecularly, the pruning of AZs might involve protein degradation, which appears to be counteracted by bassoon and piccolo via negative regulation of ubiquitin-ligase activity (Waites *et al*, 2013), such that their greater abundance might protect the largest of the initially formed AZs. In addition, a mutual regulation of ribbon size and Ca^{2+} -channel clusters by Ca^{2+} influx and RIBEYE expression levels, respectively, was indicated in zebrafish lateral line hair cells during early development (Sheets *et al*, 2011, 2012). Super-resolution live-imaging of fluorescently-tagged ribbons and other AZ/PSD components in organotypic organ of Corti cultures will be required to directly evaluate the hypotheses of merging and pruning in future studies.

Maturation of Ca^{2+} influx-exocytosis coupling in IHCs

So far, neither the intrinsic Ca^{2+} dependence of exocytosis nor the Ca^{2+} influx-exocytosis coupling in immature IHCs was well understood. We characterized the intrinsic Ca^{2+} dependence of exocytosis and the spatial coupling between Ca^{2+} influx and exocytosis in IHCs before and after the onset of hearing. We note that the amplitude of the fast C_m rise evoked by Ca^{2+} uncaging, as used for studying the intrinsic Ca^{2+} dependence of exocytosis, was about 50 times larger

than that reflecting the exocytosis of the RRP, analyzed for apparent Ca^{2+} dependence. Moreover, the mediating vesicle population(s) remain to be elucidated and might involve parallel extrasynaptic fusion of vesicles or vesicle priming and subsequent fusion (Fuchs *et al*, 2003). We assume that fast C_m rise reports synaptic vesicle fusion and the Ca^{2+} dependence of transmitter release, based on the observations of (i) partial cross-depletion of exocytosis elicited by Ca^{2+} influx and uncaging (Beutner *et al*, 2001) and (ii) the abolition of fast exocytosis in the absence of the putative priming factor and Ca^{2+} sensor of hair cell exocytosis otoferlin (Roux *et al*, 2006; Pangršič *et al*, 2010). The Ca^{2+} dependent kinetics of the fast component were, when present, comparable between IHCs before and after the onset of hearing, suggesting a common Ca^{2+} sensor of exocytosis, most likely otoferlin. It will be interesting to study the molecular changes underlying the emergence of the fast component in immature IHCs in future experiments.

What then accounts for the major developmental increase in efficiency of whole-cell Ca^{2+} influx to evoke exocytosis? In agreement with a recent single-channel study (Zampini *et al*, 2010) we find that immature IHCs express several thousands of $\text{Ca}_v1.3$ channels. Upon maturation and the cessation of Ca^{2+} action potential firing (Kros *et al*, 1998; Brandt *et al*, 2007), Ca^{2+} influx was largely restricted to AZs (Fig 5B). This reduction of extrasynaptic Ca^{2+} channels likely accounts for much of the dramatic developmental increase in the efficiency of Ca^{2+} influx to drive exocytosis (Beutner & Moser, 2001; Johnson *et al*, 2005) and lowers the metabolic expenditure for clearing Ca^{2+} . Within synapses, Ca^{2+} channels appear to be retained and consolidated and, likely due to stabilizing interactions with scaffolding proteins.

Biophysical experiments and modeling indicated tighter spatial coupling between Ca^{2+} influx and RRP exocytosis after the onset of hearing. In agreement with our own and other published work, m was near unity for manipulation of $N_{\text{Ca}} \times P_{\text{open}}$ in mature IHCs (Fig 7D). This means that release scaled linearly with the number of open channels, suggesting either a Ca^{2+} -nanodomain-like coupling or a linear intrinsic Ca^{2+} -dependence of exocytosis. However, we found a larger m when manipulating i_{Ca} , consistent with a nonlinear Ca^{2+} -dependence also seen with Ca^{2+} uncaging. On the other hand, in immature IHCs we found a supralinear apparent Ca^{2+} dependence of exocytosis when manipulating i_{Ca} or $N_{\text{Ca}} \times P_{\text{open}}$, consistent with previous reports (Johnson *et al*, 2005) and indicative of a more Ca^{2+} -microdomain-like coupling with a cooperative Ca^{2+} sensor. Together, these findings argue against the hypothesis that the Ca^{2+} dependence of IHC exocytosis is “linearized” upon maturation as a result of a change in the Ca^{2+} sensor of fusion (Johnson *et al*, 2010). Moreover, the “linearization” of Ca^{2+} dependence of exocytosis observed in whole-cell C_m recordings is unlikely to result from summation of heterogeneously supralinear AZs (Heil & Neubauer, 2010) because in such a scenario the apparent Ca^{2+} dependence should be linear no matter how it was manipulated.

Does the experimentally observed “linearization” of apparent Ca^{2+} dependence during manipulation of $N_{\text{Ca}} \times P_{\text{open}}$ indeed reflect the establishment of a Ca^{2+} -nanodomain control of exocytosis in mature IHCs? After all, these protocols are thought to apply only to initial release at stimulus onset when pool depletion has not commenced (Augustine *et al*, 1991; Stanley, 1993; Mintz *et al*, 1995). Here, however, we had to accept considerable release of the RRP

when probing exocytosis with 20 ms depolarizations in elevated $[Ca^{2+}]_e$ to obtain a good signal to noise ratio. Therefore, we performed biophysical modeling to critically assess the consistency of our conclusions and also to gain further insights into the mechanism of IHC stimulus-secretion coupling. Combining data from a variety of different experimental approaches, we carefully investigated the effects of release saturation, resulting primarily from RRP depletion, on the predicted apparent Ca^{2+} dependence of exocytosis. Saturation was most pronounced when increasing i_{Ca} and less prevalent during manipulation of $N_{Ca} \times P_{open}$. Importantly, RRP depletion is counteracted by fast vesicle replenishment at the IHC AZ (Pangršič *et al*, 2010; Goutman 2012), and for experimental m estimation during i_{Ca} manipulation we restricted the Q_{Ca} range outside the saturating range. The m estimates derived from full and reduced Q_{Ca} range did not differ much for $N_{Ca} \times P_{open}$ (Supplementary Fig S8). Because m was much lower for manipulation of $N_{Ca} \times P_{open}$ than for i_{Ca} , opposite of the expected result if saturation impacted our analysis, we trust that it indicates Ca^{2+} -nanodomain control of exocytosis at mature AZs. The m estimate of approximately 1.4 placed strong constraints on the topography of the AZ models. Within our modeling framework and explored parameter space this was reproduced best when favoring control of exocytosis by “private channels”, which may indeed reflect molecular coupling (e.g. Liu *et al*, 2011).

The notion of a developmental tightening of the spatial Ca^{2+} channel release site coupling is further supported by a decrease of the EGTA-sensitivity of RRP exocytosis (6% ΔC_m reduction with 5 mM [EGTA]_i compared with perforated-patch in p14–25 IHCs versus 37% in p6 IHCs for 10 ms depolarization, Beutner & Moser, 2001). A similar conclusion was reached at the calyx of Held synapse (Wang *et al*, 2008), another auditory synapse with high rates of synaptic transmission and high fidelity. There, developmental downregulation of the AZ component septin 5 seems critically involved in the transition from Ca^{2+} -microdomain control before to Ca^{2+} -nanodomain control after the onset of hearing (Yang *et al*, 2010). The precise molecular mechanism governing the maturation of coupling between Ca^{2+} channels and release sites at the IHC AZ remains to be elucidated in future studies.

Materials and Methods

Animals

C57Bl/6 mice (aged 6–30 days) mice were used for experiments. All experiments complied with national animal care guidelines and were approved by the University of Göttingen Board for animal welfare and the animal welfare office of the state of Lower Saxony.

Patch-clamp and confocal Ca^{2+} imaging

IHCs from apical coils of freshly dissected organs of Corti were patch-clamped as described (Moser & Beutner, 2000). The standard pipette solution contained (in mM): 115 Cs-glutamate, 13 TEA-Cl, 20 CsOH-HEPES, 1 $MgCl_2$, 2 $MgATP$, 0.3 $NaGTP$, 10 EGTA, 0.4 Fluo-5N (Penta- K^+ salt, Invitrogen), and carboxytetramethyl-rhodamine (TAMRA)-conjugated RIBEYE-binding dimer peptide (2 μM , Francis *et al*, 2011) for Ca^{2+} imaging. The pipette solution for

perforated patch experiments contained (in mM): 135 Cs-gluconate, 10 TEA-Cl, 10 4-aminopyridine, 10 CsOH-HEPES, 1 $MgCl_2$, and 250 $\mu g/ml$ amphotericin. The extracellular solution contained (in mM): 104 NaCl, 35 TEA-Cl, 2.8 KCl, 5 $CaCl_2$, 1 $MgCl_2$, 10 NaOH-HEPES, 10 D-glucose, pH 7.3. Extracellular $[Ca^{2+}]$ was varied between nominally Ca^{2+} -free to 10 mM for experiments manipulating i_{Ca} . In all cases, NaCl concentration was adjusted for osmolality of solutions with different $CaCl_2$ concentrations. Currents were low-pass filtered at 2.9 kHz (8.5 kHz for fluctuation analysis) and sampled at 50 kHz (100 kHz for fluctuation analysis). Cells with holding current greater than -50 pA were discarded. An EPC-9 amplifier or an EPC-10 amplifier and “Patchmaster” software (HEKA Elektronik, Lambrrecht, Germany) was used for measurements. Whole-cell capacitance measurement was performed in perforated-patch configuration as previously described (Moser & Beutner, 2000), except that ΔC_m was estimated as the difference of the mean C_m over 100 ms after the end of the depolarization (the initial 40 ms were skipped). All voltages were corrected for liquid-junction potentials and voltage-drops across series resistance. Ca^{2+} currents were further isolated from background current using a P/n protocol. Confocal Ca^{2+} imaging was performed as described (Frank *et al*, 2009). In brief, presynaptic Ca^{2+} signal of IHCs were observed as changes of Ca^{2+} indicator fluorescence in XY scans using long (200–254 ms) step depolarizations to -7 mV.

UV-laser photolysis of caged calcium

UV-laser photolysis of caged calcium was performed as previously described (Nouvian *et al*, 2011). Briefly, to obtain step-wise increases in intracellular calcium, 100 μs of pulsed laser light from a DPSSL-355/1000 UV laser (Rapp OptoElectronic, Hamburg, Germany) were applied after achieving the whole-cell configuration. $[Ca^{2+}]_i$ was measured by ratiometric imaging using the calcium indicator dye mag-fura-2 (Invitrogen, Darmstadt, Germany). The dye was excited by a monochromatic light source alternating between 340 and 380 nm, and imaged using a CCD camera (TILL Photonics, Gräfelfing, Germany). $[Ca^{2+}]_i$ was determined as previously described (Beutner *et al*, 2001). The pipette solution for flash-photolysis contained (in mM): 83 Cs-gluconate, 16 TEA-Cl, 18 Cs-HEPES (pH 7.2), 0.3 mag-fura-2, 10 DM-nitrophen (gift of A. Leonov and C. Griesinger, Göttingen; or Calbiochem, Darmstadt, Germany), 5 1,3-diaminopropan-2-ol-tetraacetic acid and 10 $CaCl_2$. The extracellular solution for flash-photolysis contained (in mM): 97 NaCl, 35 TEA-Cl, 2.8 KCl, 10 $CaCl_2$, 1 $MgCl_2$, 10 Na-HEPES, 1 CsCl, 11.1 D-glucose (pH 7.2). A double exponential function with variable delay was used to fit the change in C_m , from which rate constants and response delay were extracted. In cases where the rate constants of the two components differed by less than a factor of four, the traces were re-fitted with a single exponential function. IHCs in which no C_m responses could be elicited were excluded from analysis.

Immunohistochemistry and immunofluorescence microscopy

Apical cochlear turns were fixed for 25 min in 99% methanol at $-20^\circ C$. Primary antibodies: mouse anti-CtBP2 (1:200; BD Biosciences), rabbit anti-GluA2/3 (1:200; Chemicon), rabbit anti- $Ca_v1.3$ (1:50; Alomone Labs), goat anti-CtBP2 (1:150; Santa Cruz Biotech), mouse anti-GluA2 (1:75 Chemicon), mouse anti-Sap7f407 to

bassoon (1:1,000; Abcam), mouse anti- Na^+/K^+ ATPase $\alpha 3$ subunit (1:200; Thermo Scientific). Secondary antibodies: AlexaFluor488, 594, and 647 (1:200; Molecular Probes). Images were acquired with an SP5 confocal microscope (Leica) with a 100 $\times$ oil-immersion objective (NA = 1.4). Experiments were repeated until all antigens were stained with relative uniformity in at least one preparation from each age group processed in batch. Each preparation yielded several images. Those with readily-apparent bleaching were discarded. Each image had 5–7 IHCs. Stacks from two preparations per age, stained and acquired in parallel across age groups, were analysed with Igor Pro 6 (Wavemetrics). Intensity per synapse was calculated in the optical section with the peak intensity. A region-of-interest (ROI) was determined by fitting a 2D Gaussian function on a 1 μm^2 region surrounding each CtBP2 or $\text{Ca}_v1.3$ immunofluorescent spot (Frank *et al*, 2010). Comparison of plasma membrane versus ribbon-associated $\text{Ca}_v1.3$ immunofluorescence was performed by first connecting all ribbons within 3 μm of each other in 3D, and then using the connecting lines to define the axes of cylinders of 800 nm diameter to demarcate a volume of basal plasma membrane. The average intensity of this basal plasma membrane was compared with that of the ribbon-proximal region, defined as 3D spheres of 800 nm diameter centered at all connected ribbons. Three IHCs per age group were analyzed like this. Average voxel intensity of the stack was subtracted as background.

Two-color STED images were acquired on a custom setup with a stage-scanner (Mad City Labs) using Atto590- and Atto647N-conjugated secondary antibodies (Atto-Tec) excited with pulsed diode lasers (PicoQuant) at 595 and 640 nm, temporally interleaved at 50 ns intervals. After excitation, the fluorescence of dyes in the periphery of the excited spot was suppressed through stimulated emission by the STED laser, emitting 1.2 ns pulses of 775 nm light at 20 MHz (IPG Photonics). The STED beam was guided through a polymeric phase plate (vortex pattern; RPC Photonics) and circularly-polarized before coupling into the objective. Because emitted fluorescence is confined to the zero intensity center of the STED beam, using one STED-beam for both dyes ensured that a small misalignment of the excitation beams would not show in the resulting images. The fluorescence was detected at 600–640 nm (Atto590) and 650–690 nm (Atto 647N) with SPCM-AQRH13 fiber-coupled photon-counting modules (Perkin Elmer). The fiber core acted as a confocal pinhole of 1.1 Airy disks.

Transmission electron microscopy

Cochleae were explanted around noon and perfusion-fixed on ice with 4% PFA and 0.5% glutaraldehyde in 1 $\times$ PBS, pH 7.2. After 1 h of incubation, the apical cochlear turns were explanted in 1 $\times$ PBS and fixed overnight on ice with secondary fixative comprising 2% glutaraldehyde in 0.1 M sodium cacodylate buffer, pH 7.2. The samples were washed in sodium cacodylate buffer and postfixed on ice for 1 h with 1% osmium tetroxide ((v/v) in 0.1 M sodium cacodylate buffer), followed by a 1 h washing step in sodium cacodylate buffer and three brief washing steps in distilled water. The samples were stained en bloc with 1% (v/v) uranyl acetate in distilled water for 1 h on ice. After a brief wash with distilled water, samples were dehydrated at room temperature in increasing ethanol concentrations, infiltrated in Epon resin (100% EtOH/Epon 1:1 (v/v), 30 and 90 min; 100% Epon, overnight), and embedded for 48 h at

70°C. Following conventional embedding 65–75 nm sections were obtained approaching from the anterior edge using an Ultracut E (Reichert-Jung). Slices were postfixed and -stained with uranyl acetate/lead citrate following standard protocols. Micrographs were taken with a JEOL electron microscope (JEM 1011) equipped with a Gatan Orius 1200A camera using the Digital Micrograph software package at an 8000-fold magnification.

Quantitative image analysis was performed as follows: For the size of ribbons, the longest axis of each ribbon in a section, excluding the membrane-bound rootlet region, was measured. For ribbon-associated synaptic vesicles the first row of vesicles around the ribbon were counted per section. For analysis, Student's *t*-test was used, if not otherwise indicated. Three-dimensional reconstruction was performed on 4–7 serial 70 nm sections with the free software Reconstruct (Fiala, 2005).

For electron-tomography, 250 nm conventionally embedded sections were applied to Formvar-coated 100 copper mesh grids and stained with 4% uranyl acetate and Reynold's lead citrate, subsequently 10 nm gold particles were applied to the grid. A single tilt series was acquired at a JEOL JEM 2100 electron microscope at 200 kV from -48 to $+60^\circ$ with 1° increment using Serial-EM software. The tomogram was generated using the IMOD package etomo and model was rendered using 3dmod (bio3d.colorado.edu/imod/).

High-pressure freezing

For high-pressure freezing organs of Corti were explanted as described above and placed in aluminium specimen carrier of 200 μm (type A) depth filled with inhibiting solution (Pangršič *et al*, 2010). The aluminium lid (type B) has been dipped in hexadecan (Sigma-Aldrich) before placing onto the sample. Samples were frozen immediately using the HPM010 (Bal-tec) and rapidly transferred into liquid nitrogen for storage. Freeze substitution was performed in an EM AFS2 (Leica) according to (Rostaing *et al*, 2006; Siksou *et al*, 2007). Briefly, samples were incubated for 4 days in 0.1% tannic acid in acetone at -90°C . Before raising the temperature to -20°C ($5^\circ\text{C}/\text{h}$) and three washing steps with acetone 2% osmium tetroxide in acetone has been added and incubated for 16 h. Finally temperature has been raised to 4°C ($10^\circ\text{C}/\text{h}$) and samples were washed in acetone and warmed to RT. Finally samples were infiltrated and embedded in Epon resin.

Immunogold pre-embedding

Immunogold pre-embedding labeling on p14 wild-type mice was performed according to (Nieratschker *et al*, 2009), using the guinea pig anti-bassoon antibody (anti-bassoon (Cterm) in Fig 3F; Synaptic Systems; 1:500), mouse anti-Sap7f407 antibody (anti-bassoon (Nterm) in Fig 3F, 1:400; Abcam) and goat anti-guinea pig or anti-mouse nanogold IgG from Nanoprobes. For silver enhancement the Nanoprobes HQ SilverTM enhancement kit (Nanoprobes) was used.

Data analysis and statistical tests

Data are presented as mean $\pm$ SEM, unless otherwise specified. Normality was assessed with the Jarque–Bera test. *F*-test was used to assess equality of variance in normally distributed data sets. Unpaired, two-tailed Wilcoxon rank test (also known as Mann–Whitney *U*-test)

was used to compare data of non-normal distribution or when variances of experimental groups were unequal. In case of normally-distributed equal-variance data, Student's unpaired two-tailed *t*-test was used to compare two samples (* indicates $P < 0.05$). Comparison of dispersion was performed with a modified Levene's test (Brown–Forsythe test, Brown & Forsythe, 1974), using median instead of mean for improved robustness under non-normality. One-way ANOVA followed by Tukey's test was used to detect differences in multiple comparisons for FA.

Computational model

Supplementary Material S9 describes the model implementation in detail. In brief, the spatial positioning of RRP vesicles and Ca^{2+} channels (AZ topography) at mature and immature synapses was based on functional (fluctuation analysis, Ca^{2+} imaging, published single channel recording data) and morphological (EM and STED) estimates. Ca^{2+} channel gating was modelled as a three state Markov chain with gating rates based on the experimental data by (Neef et al, 2009). The spatial $[\text{Ca}^{2+}]$ profile was assumed to equilibrate instantaneously upon channel opening and closing. $[\text{Ca}^{2+}]$ levels were estimated by treating open channels as hemispherical Ca^{2+} sources and by using the linearized buffer approximation (Naraghi & Neher, 1997). Ca^{2+} diffusion was constrained by the reflective plasma membrane but considered unaffected by synaptic vesicles or the synaptic ribbon. Ca^{2+} profile was assumed to be shaped by the following mobile Ca^{2+} buffers: calretinin, calbindin, parvalbumin, and ATP. The concentrations of endogenous Ca^{2+} buffers were taken from (Hackney et al, 2005) and [ATP] was set to 2 mM. Ca^{2+} triggered fusion of the readily releasable pool vesicles followed the seven state Markov chain model proposed by (Beutner et al, 2001) for both mature and immature synapses. The refilling of the vesicle docking sites was treated as a single step process with a fixed rate. The value of the refilling rate was set based on the experimental data by (Pangršič et al, 2010). Channel gating and vesicle fusion-replenishment dynamics were propagated by using Gillespie's algorithm (Gillespie, 1977) and its extended version for time dependent reaction rates. All calculations were performed in MATLAB. To decrease the computation time to convenient level, 100 CPU cores were used for simulating a chosen synapse scenario.

Supplementary information for this article is available online: <http://emboj.embojpress.org>

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Author contributions

ABW, MAR, CW and TM designed the study. ABW performed electrophysiology, Ca^{2+} imaging and analysis of immunohistochemical data. ABW and TP performed calcium uncaging experiments. MAR performed immunohistochemistry, confocal and STED microscopy. MG and FW constructed the biophysical model with input from ABW, MAR, CW and TM. All numerical simulations were performed and analyzed by MG. FG built the two-channel STED microscope. TF performed non-stationary fluctuation analysis. CW performed electron microscopy, SM did part of the immuno-EM. ABW, MG, MAR, CW and TM prepared the manuscript.

Conflict of interest

The authors declare that they have no conflict of interest.

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