

Glutamatergic Signaling at the Vestibular Hair Cell Calyx Synapse

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In the vestibular periphery a unique postsynaptic terminal, the calyx, completely covers the basolateral walls of type I hair cells and receives input from multiple ribbon synapses. To date, the functional role of this specialized synapse remains elusive. There is limited data supporting glutamatergic transmission, K^+ or H^+ accumulation in the synaptic cleft as mechanisms of transmission. Here the role of glutamatergic transmission at the calyx synapse is investigated. Whole-cell patch-clamp recordings from calyx endings were performed in an *in vitro* whole-tissue preparation of the rat vestibular crista, the sensory organ of the semicircular canals that sense head rotation. AMPA-mediated EPSCs showed an unusually wide range of decay time constants, from <5 to >500 ms. Decay time constants of EPSCs increased (or decreased) in the presence of a glutamate transporter blocker (or a competitive glutamate receptor blocker), suggesting a role for glutamate accumulation and spillover in synaptic transmission. Glutamate accumulation caused slow depolarizations of the postsynaptic membrane potentials, and thereby substantially increased calyx firing rates. Finally, antibody labelings showed that a high percentage of presynaptic ribbon release sites and postsynaptic glutamate receptors were not juxtaposed, favoring a role for spillover. These findings suggest a prominent role for glutamate spillover in integration of inputs and synaptic transmission in the vestibular periphery. We propose that similar to other brain areas, such as the cerebellum and hippocampus, glutamate spillover may play a role in gain control of calyx afferents and contribute to their high-pass properties.

Key words: calyx; glutamate receptor; hair cell; spillover; synaptic transmission; vestibular

Introduction

In the nervous system, most synaptic terminals form bouton endings. However, there are examples of much larger, specialized presynaptic and postsynaptic endings. For example, in the auditory brainstem, the calyx of Held and the endbulb of Held are two giant presynaptic endings that form hundreds of synapses with their postsynaptic partners to faithfully transmit sound information (Ryugo and Fekete, 1982; Sätzler et al., 2002). In the vestibular sensory epithelia, the basolateral walls of type I hair cell are completely covered by a giant postsynaptic calyx ending that has no fenestrations (Goldberg, 2000; Eatock and Songer, 2011). It is highly likely that the peculiar morphology of the hair cell/calyx

synapse has consequences for synaptic transmission. A few studies have investigated the mechanisms of synaptic transmission in the vestibular periphery. Type I hair cells contain ~ 7 – 22 individual ribbon synapses per hair cell (Lysakowski and Goldberg, 2008). There is evidence for glutamatergic synaptic transmission (Demêmes et al., 1995; Matsubara et al., 1999; Bonsacquet et al., 2006) and limited characterization of synaptic activity (Rennie and Streeter, 2006; Holt et al., 2007; Dulon et al., 2009; Contini et al., 2012; Songer and Eatock, 2013; Highstein et al., 2014). To date it remains unclear how exactly the hair cell receptor potential is converted into an afferent firing pattern at this unusual synapse.

In addition to quantal release at the type I hair cell/calyx synapse, a phenomenon termed “nonquantal” transmission occurs, in which slow changes in the calyx membrane potential mirror changes in hair cell receptor potential (Yamashita and Ohmori, 1990; Holt et al., 2007; Songer and Eatock, 2013; Highstein et al., 2014). Experimental evidence suggests both K^+ accumulation (Holt et al., 2007; Lim et al., 2011; Contini et al., 2012) and H^+ accumulation (Highstein et al., 2014) in the synaptic cleft as possible underlying mechanisms. Alternatively, it has been suggested, but not proven experimentally, that glutamate accumulation in the synaptic cleft might lead to nonquantal postsynaptic potential changes in the calyx (Yamashita and Ohmori, 1990; Goldberg, 1996).

Here, we investigate the properties of glutamatergic synaptic transmission at the type I hair cell/calyx synapse in the excised rat crista, the sensory organ of the semicircular canals. We show that

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EPSCs at this specialized synapse have an unusually wide range of time courses, including very slow events. We provide evidence that slow components of EPSCs are partially due to glutamate accumulation and spillover. Immunolabeling of synaptic ribbons and postsynaptic glutamate receptor patches reveals that these elements are often not directly juxtaposed, providing morphological evidence consistent with postsynaptic receptor activation via glutamate spillover. Finally, we show that glutamate accumulation causes slow depolarization of the postsynaptic membrane potential, bringing it closer to AP threshold. We propose that this effect contributes to the reported increase in sensitivity and signal-to-noise ratio (Fernández and Goldberg, 1971; Sadeghi et al., 2007) of the afferent response at higher frequencies of rotation (10–20 Hz for natural head movements), making these synapses ideal “high-frequency event” detectors.

Materials and Methods

Animals and preparation

Whole-cell patch-clamp recordings were performed from type I hair cells and calyx afferent endings in the central region of cristae of 12- to 28-day-old rats of either sex. In accordance with animal protocols approved by the Johns Hopkins University Animal Care and Use Committee, rats (Sprague-Dawley; Charles River Laboratories) were deeply anesthetized by isoflurane inhalation and decapitated. The inner ear tissue was removed from the temporal bone and placed into extracellular solution. The bony labyrinth was opened and part of the membranous labyrinth was dissected, including ampullae, utricle, the superior branch of the vestibular nerve that contains the afferent fibers and Scarpa's ganglion including its afferent fiber somata. The membranous labyrinth was then opened above the cristae and utricle and remaining cupulae and otoliths located on top of hair cells were removed. The preparation was secured on a coverslip under a pin, transferred to the recording chamber, and perfused with extracellular solution at a rate of 1.5 ml/min. The image was visualized with a 40× water-immersion objective, differential interference contrast (DIC) optics (Examiner D1 microscope, Zeiss), additional 4× magnification and viewed on a monitor via a NC70 Newvicon camera (Dage MTI). Type I hair cell recordings were performed from the lateral wall of the hair cell, after separating the calyx from the hair cell wall with positive pressure applied by the patch pipette. For calyx recordings, the pipette was positioned at the base of the calyx.

Data acquisition and analysis

Drugs and solutions. The intracellular solution for hair cell recordings contained the following (in mM): 135 KCl, 3.5 MgCl₂, 0.1 CaCl₂, 5 EGTA, 5 HEPES, 2.5 Na₂ATP, 290 mOsm, pH 7.2 (KOH). The intracellular solution for calyx afferent recordings contained the following (in mM): 20 KCl, 110 K-methanesulfonate, 0.1 CaCl₂, 5 EGTA, 5 HEPES, 5 Na₂ phosphocreatine, 4 MgATP, 0.3 Tris-GTP, 290 mOsm, pH 7.2 (KOH). The extracellular solution contained the following (in mM): 5.8 KCl, 144 NaCl, 0.9 MgCl₂, 1.3 CaCl₂, 0.7 NaH₂PO₄, 5.6 glucose, 10 HEPES, 300 mOsm, pH 7.4 (NaOH). To measure the EPSC reversal potential, to block large K⁺ currents at more positive holding potentials, in the intracellular solution KCl/K-methanesulfonate was replaced by CsCl, 280 mOsm, pH 7.2 (CsOH). In some experiments, to increase the extracellular K⁺ concentration to 40 mM, equimolar NaCl was replaced with KCl. Drugs were dissolved daily in the extracellular solution to their final concentrations from frozen stocks. Application of drug solutions was performed using a gravity-driven flow pipette (100 μm in diameter) placed near the recorded calyx, connected with a VC-6 channel valve controller (Warner Instruments). Glutamate transporter blocker DL-threo-β-Benzyloxyaspartic acid (DL-TBOA), competitive glutamate receptor blocker Kynurenic acid (KynA), HCN channel (Ih) blocker 4-Ethylphenylamino-1,2-dimethyl-6-methylaminopyrimidinium chloride (ZD7288), 4-aminopyridine (4-AP) (to block low-threshold K⁺ channels), NMDA receptor blocker (RS)-3-(2-Carboxypiperazin-4-yl)-propyl-1-phosphonic acid ((RS)-CPP), AMPA receptor blocker 2,3-dioxo-6-nitro-1,2,3,4-tetrahydrobenzo[f]quinoxaline-7-sulfonamide (NBQX) and cyclothiazide (CTZ) (to remove desensitization from AMPA receptors)

were purchased from Tocris Bioscience. Na⁺ channel blocker tetrodotoxin (TTX), Na₂ATP, Tetraethylammonium (TEA; to block high-threshold K⁺ channels), KCNQ channel blocker XE-991, and BaCl₂ (to block inward rectifier K⁺ channels) were purchased from Sigma-Aldrich.

As the basolateral walls of type I hair cells are covered by afferent calyx terminals, the question arose whether the perfused extracellular solutions did reach the synaptic cleft. When the preparation was perfused with NBQX, an AMPA receptor blocker, synaptic activity was abolished, demonstrating that the synaptic cleft between type I hair cell and calyx afferent was accessible for perfusion.

Data acquisition. Recording pipettes were fabricated from 1 mm borosilicate glass (WPI). Pipettes were pulled with a multistep horizontal puller (Sutter), fire polished, and coated with Sylgard (Dow Corning). Pipette resistances were 7–10 MΩ for calyces and 3–5 MΩ for hair cells. Recordings were performed either at 25–27°C (room temperature) or at 35–37°C (close to body temperature). All measurements were acquired using pCLAMP 10.2 software in conjunction with a Multiclamp 700B amplifier (Molecular Devices), digitized at 50 kHz with a Digidata 1322A, and low-pass filtered at 10 kHz. In current-clamp mode, bridge balance and pipette capacitance neutralization were adjusted. In both voltage-clamp and current-clamp, voltages were corrected off-line for the measured liquid junction potential of –9 mV for calyx recordings and of –4 mV for hair cell recordings.

Cell membrane and recording parameters. Membrane capacitance (C_m), membrane resistance (R_m), and series resistance (R_s) were calculated from the average current response (50 repetitions) to 10 mV hyperpolarizing steps from –79 to –89 mV (V_{step}). C_m was calculated as $C_m = Q/V_{step}$, where Q is the amount of charge stored on the membrane capacitor calculated as the area under the transient current response to the voltage step. R_m was calculated from the steady-state current response (I_{steady}) as $R_m = V_{step}/I_{steady}$. With $R_m \gg R_s$, R_s could be calculated as $R_s = V_{step}/I_{peak}$, where I_{peak} is the peak current of the transient current response to the voltage step. C_m was 28.1 ± 5 pF, R_m was 102.5 ± 7.6 MΩ, and R_s was 26 ± 2 MΩ ($n = 33$). Recordings with $R_s > 40$ MΩ were discarded.

To test for the effects of R_s errors in the recording configuration, in eight calyx afferent recordings, EPSCs were recorded before and after R_s compensation (85–95%) in 5.8 mM K⁺ extracellular solution at a holding potential of –103 mV. The large negative holding potential resulted in larger EPSC amplitudes and improved signal-to-noise ratio. For this dataset, with an average holding current of 337 pA, the estimated voltage error changed from 9.3 to 1.3 mV after R_s compensation. The pooled EPSC amplitude distributions for the eight recordings showed an extended tail with larger EPSC amplitudes after R_s compensation. The average EPSC amplitude doubled with R_s compensation, from 26.4 ± 4.9 to 58.4 ± 12 pA ($n = 8$, paired t test, $p = 0.02$).

Regarding EPSC kinetics, for the pooled EPSCs from eight calyces, no significant difference was observed before and after R_s compensation, with the average 10–90% rise times of 2.2 ± 0.9 ms ($n = 1397$ EPSCs) and 1.5 ± 0.4 ms ($n = 1347$ EPSCs; paired t test, $p = 0.4$) and EPSC decay time constants of 28.3 ± 9.4 ms and 22.5 ± 8.7 ms (paired t test, $p = 0.4$), respectively. Note that the distributions of 10–90% rise time and decay time constant showed a slight shift to the left after R_s compensation. For decay times, mainly the number of EPSCs with decay times <2 ms increased. This was expected; the time constant of the voltage-clamp was 0.7 ms ($C_m \times R_s$) without R_s compensation and 0.1 ms with R_s compensation, suggesting that only values $> \sim 1.4$ ms could be measured correctly without R_s compensation and $> \sim 0.2$ ms with R_s compensation.

Accordingly, for studying EPSC amplitudes, only recordings with R_s compensation were used, whereas for investigating decay time constants, recordings without R_s compensation were included, as the effects regarding spillover investigated in this study concerned mainly the slow components of EPSCs, with values much slower than the estimated clamp speed, even without R_s compensation.

Estimating the hair cell membrane potential. Efflux of K⁺ from the hair cells and/or the calyx ending could additionally affect [K⁺] in the synaptic cleft and subsequently affect hair cell and/or calyx membrane potential (Lim et al., 2011). For this study, calyx membrane potentials were

recorded with the synaptic cleft intact. However, for recording hair cell membrane potentials, the synaptic cleft was opened to allow for the patch pipette to reach the hair cell lateral wall, possibly resulting in a loss of K^+ accumulation due to K^+ efflux from the hair cell and/or calyx. Second, *in vivo*, the stereocilia bundles are exposed to endolymph (high in $[K^+]$) and the hair cell bodies are exposed to perilymph (high in $[Na^+]$) whereas *in vitro*, hair cell bodies and stereocilia are bathed in perilymph-like solution. Regarding mechano-transduction, Na^+ -influx should substitute for K^+ -influx *in vitro*. However, fewer transduction channels may be open at rest, due to a higher $[Ca^{2+}]$ in the perilymph-like solution compared with endolymph, resulting in a more negative resting membrane potential *in vitro* compared with *in vivo* (Farris et al., 2006). The “endovestibular” potential, as measured between endolymph and perilymph space *in vivo* is small, (2–5 mV) compared with the large endocochlear potential (~80 mV; Smith et al., 1958; Schmidt, 1963) and therefore its possible loss *in vitro* is of minor concern.

Holding potential. At a fairly negative holding potential of -103 mV, the holding current was typically ~ 300 pA and holding currents up to ~ 500 pA were considered acceptable. To assure that a large component of the holding current was due to active ion channels and not to unspecific leak, a mixture of blockers was applied to test the possible contribution of different ion channels (100 μM XE-991, 10–30 mM TEA, 2–4 mM 4-AP, 2 mM $BaCl_2$, 50 μM ZD7288 to block KCNQ channels, potassium channels, inward rectifiers, and I_h). Such a mixture reduced holding currents by $>50\%$ ($n = 9$, control: 433 ± 33 pA, with blockers: 205 ± 35 pA, paired t test, $p < 0.0002$).

Second, if large holding currents were indicative of large unspecific leak currents, one would predict that in “leaky recordings” the measured resting membrane potentials were less negative. Resting membrane potentials were compared for calyces with holding currents between 0 to -500 pA in 100 pA bins. Average resting membrane potentials were -69.8 ± 2.1 mV ($n = 23$), -70.7 ± 1.9 mV ($n = 27$), -69.1 ± 1.2 mV ($n = 34$), -67.6 ± 0.8 mV ($n = 43$), and -66.2 ± 1.8 mV ($n = 46$), respectively, and were not different from each other (ANOVA, $p = 0.07$), suggesting that recordings with large holding currents were qualitatively as good as recordings with smaller holding currents.

Data analysis. The EPSC decay time was either fit with one or two exponentials. For the fit with two exponentials, the following equation was used:

$$f(t) = A_1 \times [\exp(-t/\tau_1)] + A_2 \times [\exp(-t/\tau_2)],$$

where τ_1 and τ_2 are the “faster” and “slower” τ_{decays} , A_1 and A_2 are the respective amplitudes, and t is time.

Data were analyzed using Clampfit (Molecular Devices), MiniAnalysis (Synaptosoft), and MATLAB (MathWorks) software. EPSCs were detected using a routine in MiniAnalysis, with a threshold set at three times the value of the root mean square of the baseline noise. Events that showed summation were identified by eye and not included in the analysis. For τ_{decay} , 10–90% of the decay times were fit. Data are reported as mean \pm SE. Significance was measured by paired or unpaired t test for comparison of two parameters. For comparisons of more than two conditions, one-way ANOVA with Tukey’s HSD *post hoc* test was used. Level of statistical significance was set at $\alpha = 0.05$.

Labeling of fibers with biocytin

For calyx recordings, 0.3% biocytin was added to the intracellular solution. After recording, tissue was fixed in 4% paraformaldehyde overnight at 4°C and then kept in PBS. The tissue was quenched in 1% hydrogen peroxide for 10 min, permeabilized in 2% Triton X-100 in PBA for 1 h, incubated for 2 h using VECTASTAIN ABC kit (Vector Laboratories), and finally in 0.1% OsO_4 for 30 s. In between steps, the tissue was rinsed three times (10–30 min each) in PBS. After a final PBS rinse (3×15 min), the tissue was mounted on a slide with Mowiol mounting solution. A Nikon E600 compound microscope with 40 \times magnification was used to inspect the tissue for the number of labeled calyces.

Immunofluorescence and microscopy

Excised whole-tissue preparations of horizontal and anterior cristae of 17- to 21-d-old rats were immediately transferred from PBS to 4% para-

formaldehyde in PBS, fixed for 50 min, and then rinsed three times in PBS. Cochlear organs of Corti from equivalently aged rats were excised and fixed as described previously (McLean et al., 2009). For both vestibular and cochlear preparations, samples were first incubated in blocking buffer (PBS with 5% normal goat serum, 4% Triton X-100, and 1% saponin) for 1–2 h. Preparations were then incubated in primary antibody diluted in blocking buffer overnight. Samples were rinsed three times 10 min each in PBS with 0.2% Triton X-100 (PBT) before incubation in the appropriate secondary antibody diluted 1:500 in blocking buffer for 2 h. Immunofluorescent staining was performed using the following antibodies: mouse monoclonal (IgG1) anti-CTBP2 (612044, 1:300; BD Bioscience), rabbit polyclonal anti-GluR2/3 (AB1506; 1:100; Millipore), mouse monoclonal (IgG2A) anti-PSD95 (75–028, 1:300; NeuroMab), mouse monoclonal (IgG2A) anti-TuJ (neuronal class III β -tubulin; MMS-435P, 1:300, Covance), and mouse monoclonal (IgG1) anti-TuJ (MAB1637, 1:300, Millipore). Secondary antibodies, AlexaFluor 488 goat anti-mouse IgG1 (A21121), AlexaFluor 488 goat anti-mouse IgG2_A (A21131), AlexaFluor 647 goat anti-mouse IgG1 (A21240), AlexaFluor 647 goat anti-mouse IgG2_A (A21241), and AlexaFluor 568 goat anti-rabbit IgG (A11011) were purchased from Molecular Probes (Life Technologies). Preparations were again rinsed three times 10 min each in PBT, and then once for 10 min in PBS before mounting on glass slides in Vectashield mounting medium (Vector Laboratories). All incubation and rinses were performed on a rocking table at room temperature. The antibodies used in this study were chosen because their specificity has been characterized previously, and in all cases, experiments performed in the absence of primary antibody showed no immunoreactivity.

Fluorescence images were acquired using an Olympus Fluoview FV1000 confocal microscope with a 60 \times Olympus PlanoApo oil-immersion lens (NA 1.42) under the control of the Olympus Fluoview FV1000 v2.1 software. z -Stacks (46–100 optical sections) through the majority of the preparation were collected in 0.44 μm steps. The step size (optical section thickness) was determined by stepping at half the distance of the theoretical z -axis resolution (the Nyquist sampling frequency). Images were acquired in a 1024 \times 1024 raster ($x = y = 0.207$ μm /pixel) at subsaturating laser intensities for each channel. Images are presented as z -projections through a subset of the collected optical stack. All quantitative image analysis was performed on the raw image stacks, without deconvolution, filtering, or gamma correction.

To determine their total number and relative location, immunolabels were detected automatically using the Spots function in the Imaris 6.4 three-dimensional (3D) image visualization and analysis software (Bitplane). Image files were resampled upon opening in Imaris to include a specific 3D ROI that included the calyces to be analyzed. Automatically detected immunolabels were verified by eye by rotating the 3D image, and immunolabels not contained within the inner face of calyces were manually removed. Distances of CTBP2, GluR2/3, and PSD95 labels were calculated from the x , y , and z coordinates of immunolabels.

Results

Recordings from calyx afferent terminals in the crista of semicircular canals

To study the properties of synaptic transmission between type I vestibular hair cells and their afferent calyx endings, a method was established to dissect the rat inner ear and excise the cristae, the sensory organs of the semicircular canals, along with the utricle, nerve fibers and Scarpa’s ganglion (Fig. 1A1). Recordings were performed from the central zone of the crista at postnatal days (P)12–P28, because type I hair cells have acquired their characteristic morphological and electrophysiological properties by P12 (Rüsch et al., 1998; Hurley et al., 2006). Furthermore, the percentage of type I hair cells with morphologically fully developed calyces increases during development (Favre and Sans, 1979) and reaches a plateau at \sim P10 in the central zones of vestibular end organs (Rüsch et al., 1998), which contain higher numbers of type I hair cells and calyx afferent endings compared

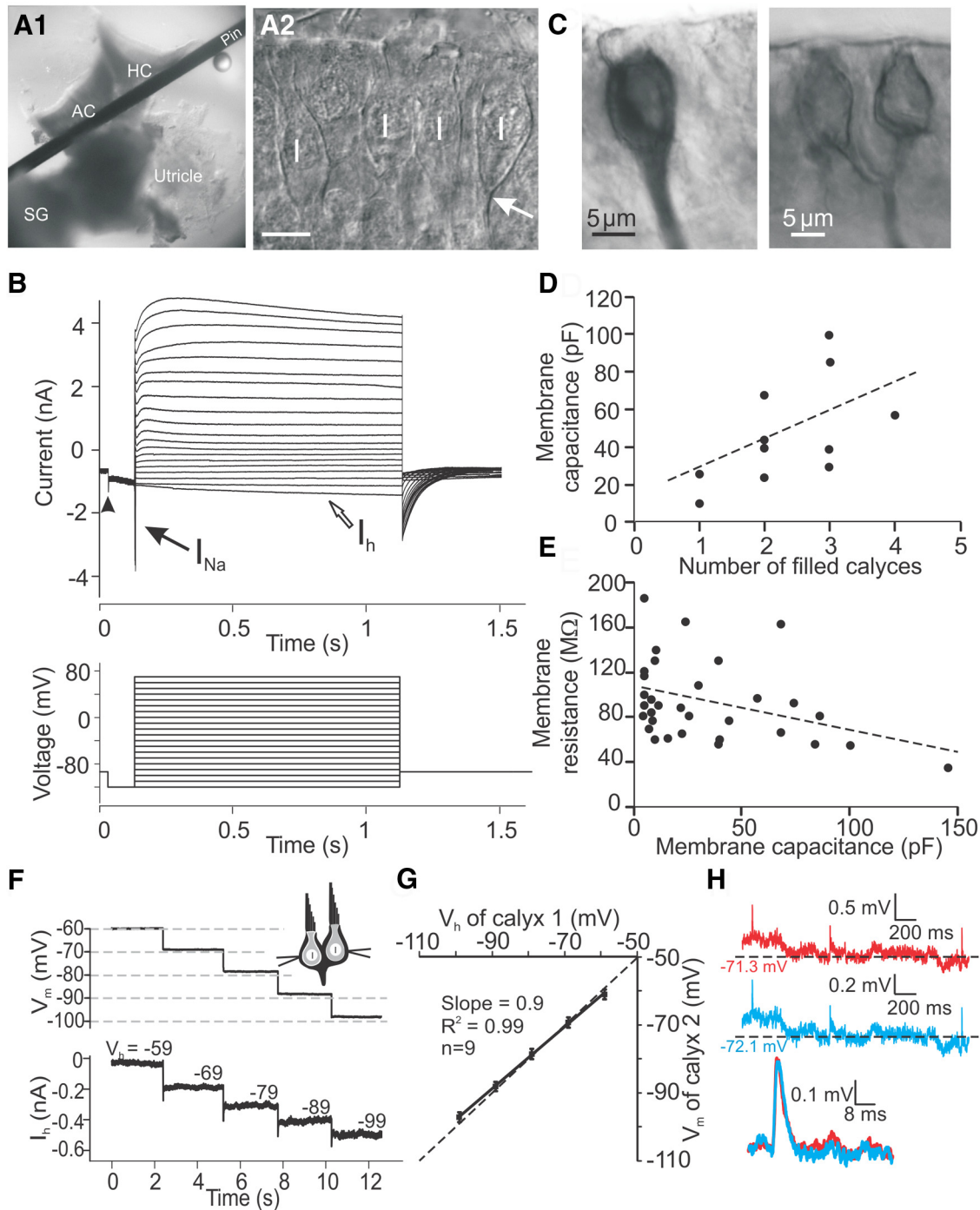


Figure 1. Calyx afferents visualized and recorded from the rat vestibular periphery. **A**, Whole-tissue preparation of anterior and horizontal canal cristae (AC and HC) held under a pin, along with the utricle and Scarpa's ganglion (SG; **A1**). Using DIC optics, calyces in the central zone were visualized as bright thick lines around type I hair cells (marked by I in **A2**). Recordings were made at the base of the calyx (arrow). Scale bar, 10 μm . **B**, Response of a calyx to the voltage step protocol (initial holding voltage of -103 and 10 mV steps between -129 and 61 mV). Calyx recordings exhibited large Na^+ inward currents (black arrow), delayed rectifier type outward currents during depolarizing voltage steps, and slowly deactivating currents during hyperpolarizing steps (arrowhead) and hyperpolarization-activated currents (I_h ; open arrow). **C**, Biocytin fills of example calyx recordings. **D**, Calyces had a wide range of C_m that increased with the number of labeled calyces. **E**, Calyces showed a wide range of R_m that decreased with increasing C_m . Thirty-three calyces, including 11 calyces from **D**. **F–H**, Double-recordings from complex calyces. **F**, In one calyx, the membrane was clamped to a holding potential between -59 and -99 mV in 10 mV steps, while in the other calyx the membrane potential was measured in current-clamp. **G**, In nine double-recordings from complex calyces, the difference between holding potential (V_h) of calyx 1 and the measured membrane potential of calyx 2 (V_m) was $<4\%$. **H**, Example double-recording from two calyces within a complex calyx in current-clamp. Membrane potentials of both calyces were similar and EPSPs were similarly shaped (bottom).

with peripheral zones (Lysakowski and Goldberg, 2008). When viewed at larger magnification with DIC optics, calyces were visible as thickenings around type I hair cells, which were identified by their typical morphology with a smaller apical diameter (the so-called “neck”) of the hair cell (Fig. 1A2). Recordings were

performed at the base of the calyx, close to where the calyx transitions into the vestibular nerve fiber (Fig. 1A2, arrow). During whole-cell recordings, the average resting membrane potential of calyces was -68.1 ± 0.5 mV ($n = 173$). Voltage step protocols were used to distinguish calyx recordings from type I hair cell

recordings (Fig. 1B). Calyx recordings exhibited large Na^+ inward currents with amplitudes of several nanoamperes, whereas hair cells showed smaller or no Na^+ currents (Wooltorton et al., 2007). Calyx recordings displayed delayed rectifier type outward currents during depolarizing voltage steps and slowly deactivating currents during hyperpolarizing steps from -103 to -129 mV (Fig. 1B, arrowhead), most likely due to KCNQ and erg channel activation (Hurley et al., 2006). Similar to auditory afferent fibers (Yi et al., 2010), a hyperpolarization-activated current (I_h) was also present in calyx afferent fibers (Meredith et al., 2012; Horwitz et al., 2014).

Afferent fibers with calyx endings can innervate an individual type I hair cell or form a “complex calyx” that contacts a group of hair cells with several calyces (Fernández et al., 1988). Additionally, “dimorphic” afferent fibers form calyx endings and also send branches off to form bouton endings onto type II hair cells. To investigate their morphology, some afferent fibers were filled with biocytin during recordings. In 13 such recordings, 11 showed calyx endings (Fig. 1C) and two showed calyx endings and additional branches, most likely ending in boutons. One to four calyces were counted to connect to individual fibers. Membrane capacitance (C_m) generally increased as the number of filled calyces per fiber increased (Fig. 1D). By dividing C_m by the number of filled calyces per fiber, C_m per individual calyx was estimated at 19 ± 3 pF for these neurons ($n = 11$). For 33 afferent calyx recordings, the average C_m was 28.1 ± 5 pF (range: 5–145 pF) and the average R_m was 102.5 ± 7.6 M Ω (range: 35–185 M Ω). The large ranges of C_m and R_m were due to the varying number of calyces (Fig. 1D). As expected, C_m and R_m were inversely related (Fig. 1E).

Because the recording electrode was positioned at the base of a calyx, in the presence of complex calyces, it most likely measures events generated from all calyces that contact an individual fiber. To investigate whether individual calyces within a complex calyx are exposed to the same voltage-clamp conditions, or whether “space clamp” problems occur, recordings were performed simultaneously from the lateral membranes of two calyces that were visually identified as part of a complex calyx. At one electrode, the holding potential was stepped from -59 to -99 mV in 10 mV steps, while at the same time at the second electrode, voltage changes in the neighboring calyx were monitored (Fig. 1F). In nine paired recordings, the difference between command and measured potential was $<4\%$, confirming that space clamp errors had minimal effects on recordings from complex calyces (Fig. 1G).

To investigate whether passive properties in complex calyces affected synaptic events traveling along the calyx membrane to the recording site, EPSPs recorded by two electrodes positioned at different calyces belonging to a complex calyx were compared (in six double recordings; Fig. 1H), similar to studies performed in neural dendrites (Magee and Cook, 2000). Membrane potentials recorded at the two electrodes were similar, differing by <0.02 mV, and EPSPs were close to identical, suggesting that synaptic events were not substantially distorted by passive cable properties when traveling to the recording site.

In summary, control experiments suggest that the waveforms of synaptic events recorded from a single or complex calyx afferent are not subject to substantial errors. However, the possibility still remains that a small percentage of synaptic events might be generated in distant branches of the afferent fiber that were unaccounted for and the waveform of such events might be affected by space clamp errors and passive membrane properties.

Amplitude distributions of AMPA receptor-mediated EPSCs in calyx afferent fibers

EPSCs were observed in 27% (46 of 173) of calyx afferent recordings. The large percentage of “silent synapses” is not surprising, as the average hair cell membrane potential recorded under similar conditions was found to be quite negative, -70.5 ± 1.7 mV ($n = 10$), suggesting a low probability of release. The recorded hair cell membrane potential reflects the experimental conditions (see Materials and Methods for further explanation) and the high spontaneous firing in type I afferent fibers that has been found *in vivo* (Yang and Hullar, 2007; Lasker et al., 2008) may be due to a more positive hair cell membrane potential under *in vivo* conditions. As expected, when the hair cell was depolarized, with elevated $[\text{K}^+]$ or ATP in the extracellular solution, EPSCs were activated in most silent calyx recordings. Similarly, a recent study reported that in utricular calyx recordings some afferents did not show synaptic activity, but exhibited EPSCs when the hair bundle of the presynaptic hair cell was deflected (Songer and Eatock, 2013).

EPSCs were completely ($n = 11$) and reversibly ($n = 4$) blocked by the specific AMPA receptor blocker NBQX (10 μM ; Fig. 2A). The current voltage relation of average EPSCs was close to linear and reversed at ~ 0 mV (4 calyx recordings; Fig. 2B, C). EPSCs often showed decay times with a faster and a slower component and therefore, were tested for a possible contribution from NMDA receptors. Experiments were performed in 0 Mg^{2+} extracellular solution, to relieve Mg^{2+} block of NMDA receptors, and with added 40 μM D-serine, a cofactor of NMDA receptors. Under these conditions, application of the NMDA receptor blocker CPP (20 μM) had no effect on the average EPSC waveform, neither at a negative (-80 mV) nor at a positive ($+60$ mV) holding potential (Fig. 2D). When the EPSC decay time was fit with two exponentials, neither of the decay time constants, $\tau_{1\text{decay}}$ or $\tau_{2\text{decay}}$, were affected by CPP (Table 1). Together, these findings suggest that AMPA receptors mediate the observed EPSCs at type I hair cell/calyx synapses.

EPSC amplitude distributions showed some variability between recordings. Figure 2E shows two representative EPSC amplitude distributions that span over the range of values found in 15 recordings (Fig. 2F). Both distributions were well fit with a Gaussian and additionally exhibited a small tail at larger amplitudes. Peak amplitudes for these two distributions were 20 pA (black) and 93 pA (green), with coefficients of variation of 0.44 and 0.31 (holding potential, -103 mV). For 15 recordings, on average $>90\%$ of EPSCs were smaller than 150 pA (Fig. 2F, dashed trace) and the mean and median EPSC amplitudes were 80.8 ± 1.0 and 74.3 pA, and the resulting mean and median conductances were 784 and 721 pS ($n = 1865$ EPSCs, pooled from 15 recordings).

Variability between EPSC amplitude distributions of individual calyces could be caused by a number of mechanisms, for example by variations in hair cell membrane potential and subsequent changes in calcium influx, affecting the level of multivesicular release (Wadiche and Jahr, 2001) as it has been shown at other ribbon synapses (Singer et al., 2004; Li et al., 2009). To test whether a similar mechanism might apply here, EPSC amplitudes were compared at two different levels of hair cell depolarization, in extracellular solution with either 5.8 or 40 mM K^+ . In type I hair cell recordings, switching from 5.8 to 40 mM K^+ external solution, depolarized the hair cell membrane potential by ~ 40 mV, from -70.5 ± 1.7 mV to -29.6 ± 1.5 mV ($n = 10$). In afferent fiber recordings, when changing from 5.8 to 40 mM K^+ external solution, EPSC rates increased from 1.2 ± 0.5 EPSCs/s to

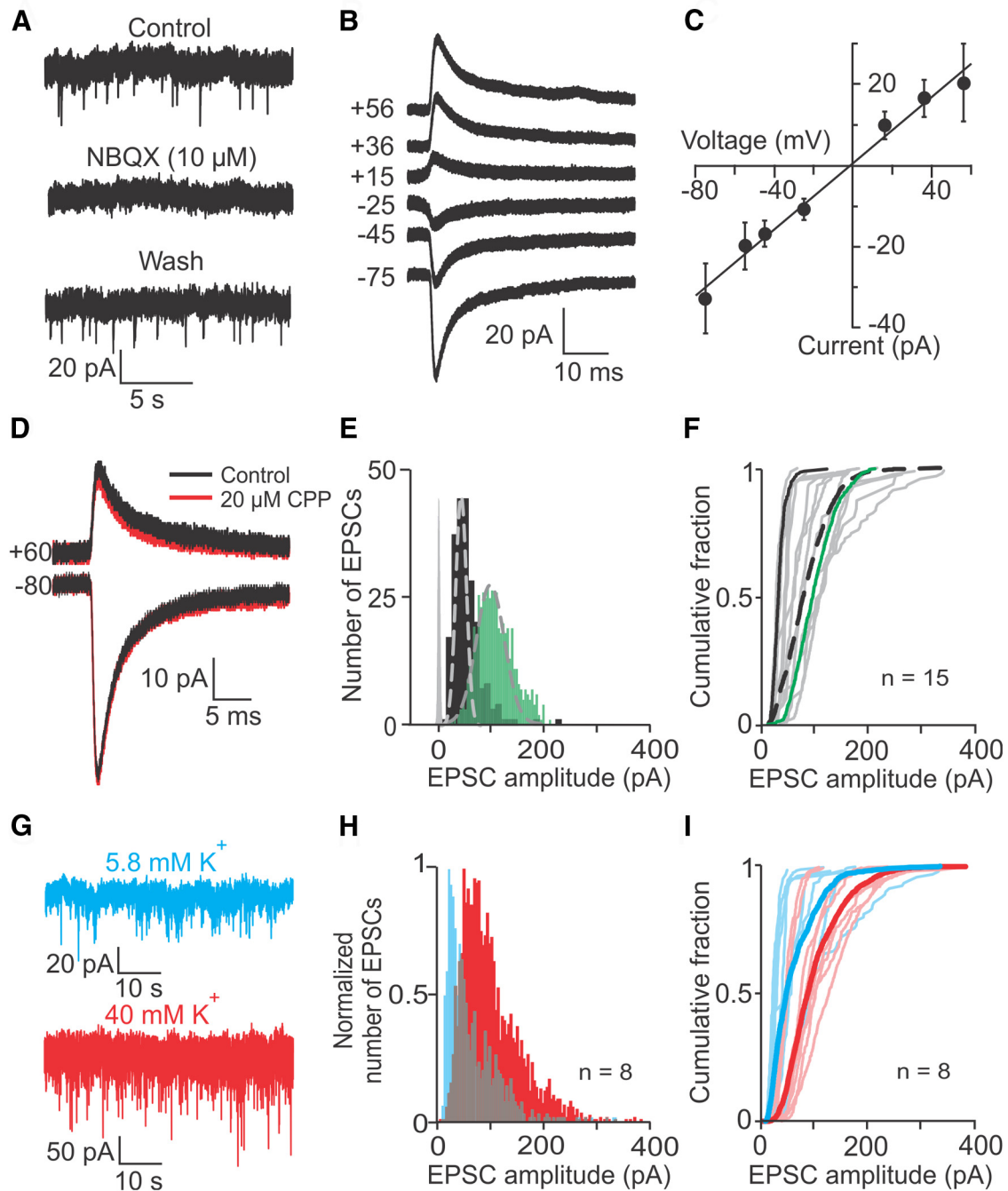


Figure 2. EPSC amplitude distributions recorded from vestibular calyces. **A**, EPSCs are mediated by AMPA receptors, as shown by complete and reversible block by the AMPA receptor blocker NBQX ($10 \mu\text{M}$). External solution with 5.8 mM K^+ ; holding potential of -103 mV . **B**, Calyx recording with average EPSCs at different holding potentials (indicated on the trace in mV). **C**, Current voltage relation of average EPSCs (4 recordings) is close to linear and reverses at $\sim 0 \text{ mV}$. **D**, The NMDA receptor blocker CPP ($20 \mu\text{M}$) had no effect on the EPSC waveform at positive or negative holding potentials (indicated on traces). **E**, Two example EPSC amplitude distributions were close to Gaussian (dashed lines), with small tails toward larger values. Baseline noise distribution in gray. **F**, Cumulative fraction plots of EPSC amplitudes for 15 calyces (gray) and their average (dashed black). Black and green lines correspond to calyces shown in **E**. **G**, EPSC amplitudes and rate increased during application of 40 mM K^+ external solution. **H**, Normalized cumulative fraction plots of EPSC amplitudes pooled from 8 calyces in 5.8 mM K^+ (blue) and in 40 mM K^+ external solution (red). **I**, Cumulative fraction plots of EPSC amplitudes for eight calyces in 5.8 mM K^+ (individual recordings: light blue; average distribution: blue) shifted to the right in 40 mM K^+ external solution (individual recordings: light red; average distribution: red).

$15.3 \pm 3.3 \text{ EPSCs/s}$ (8 calyx recordings, paired t test, $p = 0.0005$; Fig. 2G). EPSCs recorded in 40 mM K^+ were completely ($n = 7$) and reversibly ($n = 3$) blocked by $10 \mu\text{M}$ NBQX, assuring that all events were mediated by AMPA receptors, as in control conditions. In 40 mM K^+ , the peak of the pooled amplitude distribution doubled (48.5 pA , $n = 2902 \text{ EPSCs}$, 8 calyx recordings, Fig. 2H, red) compared with 5.8 mM K^+ (20.2 pA , $n = 843 \text{ EPSCs}$, 8

calyx recordings; Fig. 2H, blue). The distribution also widened, exhibiting a longer “tail.” For the population of recorded afferents, the average cumulative fraction plot showed a significant rightward shift (Fig. 2I, blue and red bold traces), with mean amplitudes increasing from $63.3 \pm 1.6 \text{ pA}$ to $101.3 \pm 1.0 \text{ pA}$ (t test, $p < 0.0001$) and the median amplitude increasing from 48.5 to 88.9 pA . This observed increase in EPSC amplitudes with hair

Table 1. EPSC kinetics before and after NMDA receptor blocker (20 μ M CPP) application in four calyces

	Condition	$V_h = -80$ mV ^a	<i>p</i> value	$V_h = 60$ mV ^b	<i>p</i> value
τ_1 (ms)	Control	2.1 \pm 0.16	0.3	4.97 \pm 0.69	0.96
	CPP	2.5 \pm 0.71		4.93 \pm 0.99	
τ_2 (ms)	Control	10.76 \pm 1.83	0.73	12.77 \pm 2.3	0.83
	CPP	10.14 \pm 0.86		13.03 \pm 2.5	
$A_2/(A_1 + A_2)$	Control	0.69 \pm 0.09	0.09	0.71 \pm 0.1	0.22
	CPP	0.54 \pm 0.1		0.66 \pm 0.08	

p values for paired *t* test. $A_2/(A_1 + A_2)$ shows relative contributions of $\tau_{1\text{decay}}$ and $\tau_{2\text{decay}}$ (see Materials and Methods).

^aControl: *n* = 984 EPSCs; CPP: *n* = 1039 EPSCs.

^bControl: *n* = 356 EPSCs; CPP: *n* = 410 EPSCs.

cell depolarization could be due to increased multivesicular release and could have important functional consequences as larger EPSC amplitudes may result in larger EPSPs, increasing the likelihood of AP generation in the calyx.

AMPA-mediated EPSCs in calyx afferent fibers have mixed fast and slow kinetics

In afferents with either single or complex calyx terminals, EPSCs showed a wide range of waveforms in 5.8 mM K^+ , with the hair cell at rest. Figure 3A shows three example EPSCs: EPSCs in Figure 3A1 and A2 are best fit with a single exponential, Figure 3A1 showing a faster event with a time constant of decay (τ_{decay}) of 0.54 ms, and Figure 3A2 showing a slower event with a τ_{decay} of \sim 6 ms. Figure 3A3 shows an EPSC with both faster and slower components, best fit with two exponentials with a τ_1 of 0.6 ms and τ_2 of 3.5 ms. Summated events made up $<$ 10% of all EPSCs and were excluded from waveform analysis (data not shown). For the population of 46 recorded calyces, τ_1 was 2.7 \pm 0.4 ms (range: 0.6–11.5 ms for individual recordings) and τ_2 was 16.1 \pm 1.9 ms (range: 4.3–101.7 ms). The contribution of the slow component to the total amplitude, measured as $A_2/(A_1 + A_2)$, was on average 48.5 \pm 0.02% (range: 13.4–71.2%), indicating large variability between recordings regarding the contribution of faster and slower components. Half of the recordings had average EPSCs with decay times best fit by two exponentials (23 of 46 cells), whereas the other half had decay times best fit with one exponential. Forty-four percent of this latter group (10 of 23 cells) had slow average $\tau_{\text{decays}} >$ 10 ms. Thus, in a large percentage of recordings (72%), a slow process was involved in the generation of EPSCs, either alone or in combination with a faster process. In summary, EPSC waveforms were variable within and between recordings and best fit with either one or two exponentials. Therefore, as both fitting methods included some level of inaccuracy and depending on the goal of the analysis, either method was applied.

To provide an overview over the range of waveform kinetics within and between recordings, distributions of τ_{decays} were assembled based on single exponential fits of EPSC decay times. The distribution of τ_{decays} for a single recording is shown in Figure 3B, with a peak $<$ 5 ms and a long tail reaching values up to 120 ms. For this recording, the mean τ_{decay} was 13.4 \pm 1.8 ms and the median was 6.8 ms (*n* = 126 EPSCs for one example calyx recording). Similarly, for the population of recorded calyces, the distribution of τ_{decays} had a peak $<$ 5 ms and showed a long tail toward larger values, with mean and median values of 18.7 \pm 2.8 ms and 3.4 ms (*n* = 4633 EPSCs, 46 calyx recordings; Fig. 3C). The slowest τ_{decays} reached values of $>$ 500 ms (Fig. 3C, inset), leading to unusually slow EPSCs. Faster EPSCs ($\tau_{\text{decay}} <$ 10 ms) spanned over the entire range of EPSC amplitudes, whereas most of the slower events ($\tau_{\text{decay}} >$ 10 ms) had smaller amplitudes ($<$ 50 pA; Fig. 3D, gray shaded area).

To test whether EPSCs with unusually slow τ_{decays} also appear at body temperature, EPSC waveforms were measured at both room temperature and body temperature in individual calyx recordings (25–27°C vs 35–37°C). Pooled distributions of τ_{decays} at both temperatures show that unusually slow EPSCs with τ_{decays} reaching values up to 273 ms were maintained at body temperature (Fig. 3E). Average decay time constants were not significantly different at both temperatures (mean τ_{decays} of 10.7 \pm 5.6 and 7.4 \pm 2.4 at 25–27°C and 35–37°C, respectively, *n* = 6 calyx recordings, paired *t* test, *p* = 0.3). Similarly, analysis of the EPSCs using double exponential fits did not show any significant difference in the average τ_{decays} (25–27°C: $\tau_{\text{fast}} = 4.3 \pm 0.9$ ms, $\tau_{\text{slow}} = 25.2 \pm 5.3$ ms; 35–37°C: $\tau_{\text{fast}} = 4.5 \pm 1.0$ ms, $\tau_{\text{slow}} = 16.2 \pm 2.5$ ms; *n* = 6; paired *t* test, *p* $>$ 0.7 for both conditions). The only difference between the two temperatures was observed for EPSCs with $\tau_{\text{decays}} <$ 10 ms. For these faster EPSCs, the mean τ_{decay} decreased at body temperature compared with room temperature (mean τ_{decay} of 3.4 \pm 0.1 ms and 3.0 \pm 0.1 ms at 25–27°C and 35–37°C, respectively, 6 calyx recordings, paired *t* test, *p* $<$ 0.001). Cumulative plots of the τ_{decays} for 46 individual recordings again illustrate the wide range of distributions between recordings at room temperature (Fig. 3F). For the average cumulative fraction plot (black line), \sim 75% of events had a $\tau_{\text{decay}} <$ 10 ms and 25% of EPSCs showed larger τ_{decays} . Cumulative fraction plots of τ_{decay} of 10 recordings at body temperature (red lines) show that their values lie within the range of recordings performed at room temperature. In summary, these findings suggest that although temperature does affect faster τ_{decays} , the underlying process responsible for the unusually slow τ_{decays} is still functional at body temperature. As such, the remaining data presented in this paper were collected at room temperature.

Glutamate spillover contributes to the slow kinetics of EPSCs at the type I hair cell/calyx synapse

One explanation for the occurrence of small and slow EPSCs is that glutamate spillover to glutamate receptors located further away from the release site occurs, as observed in other systems (DiGregorio et al., 2007). The presence of such a phenomenon at the calyx afferent seems likely because of the closed synaptic space. To test whether the slow kinetics of EPSCs could be due to glutamate spillover, three experimental approaches were used. As a first approach to induce accumulation of glutamate in the synaptic cleft the probability of glutamate release was increased by depolarizing the hair cell by raising the external $[K^+]$ from 5.8 to 40 mM. The expectation was that increased glutamate accumulation and possible spillover would cause an increase in τ_{decay} . Figure 4A1 shows the normalized average EPSC in 5.8 mM K^+ (black) versus 40 mM K^+ (gray) in an example calyx recording. As expected, the average τ_{decay} increased from 2.3 \pm 0.17 ms in 5.8 mM K^+ (*n* = 125 EPSCs) to 11.6 \pm 0.75 ms in 40 mM K^+ (*n* = 777 EPSCs; *t* test, *p* $<$ 0.0001). From this example recording, the distribution of τ_{decays} shows an increase in the number of slower decay time constants in 40 mM K^+ (Fig. 4A2). Figure 4A3 compares EPSC decay times for 17 recordings in 5.8 and 40 mM external K^+ , with increased τ_{decays} plotted above the unity line. In 9 of 17 calyces with average EPSC decay time constants of $<$ 10 ms in 5.8 mM K^+ , τ_{decay} increased in 40 mM K^+ , whereas eight recordings did not show a significant change. When all 17 recordings were considered together, the average τ_{decay} showed a significant increase of \sim 70%, from 4.9 \pm 0.6 ms in 5.8 mM K^+ to 8.5 \pm 1.7 ms in 40 mM K^+ (*n* = 17, paired *t* test, *p* = 0.03; Fig. 4F).

In a second set of experiments, glutamate accumulation was increased by application of the general glutamate transporter

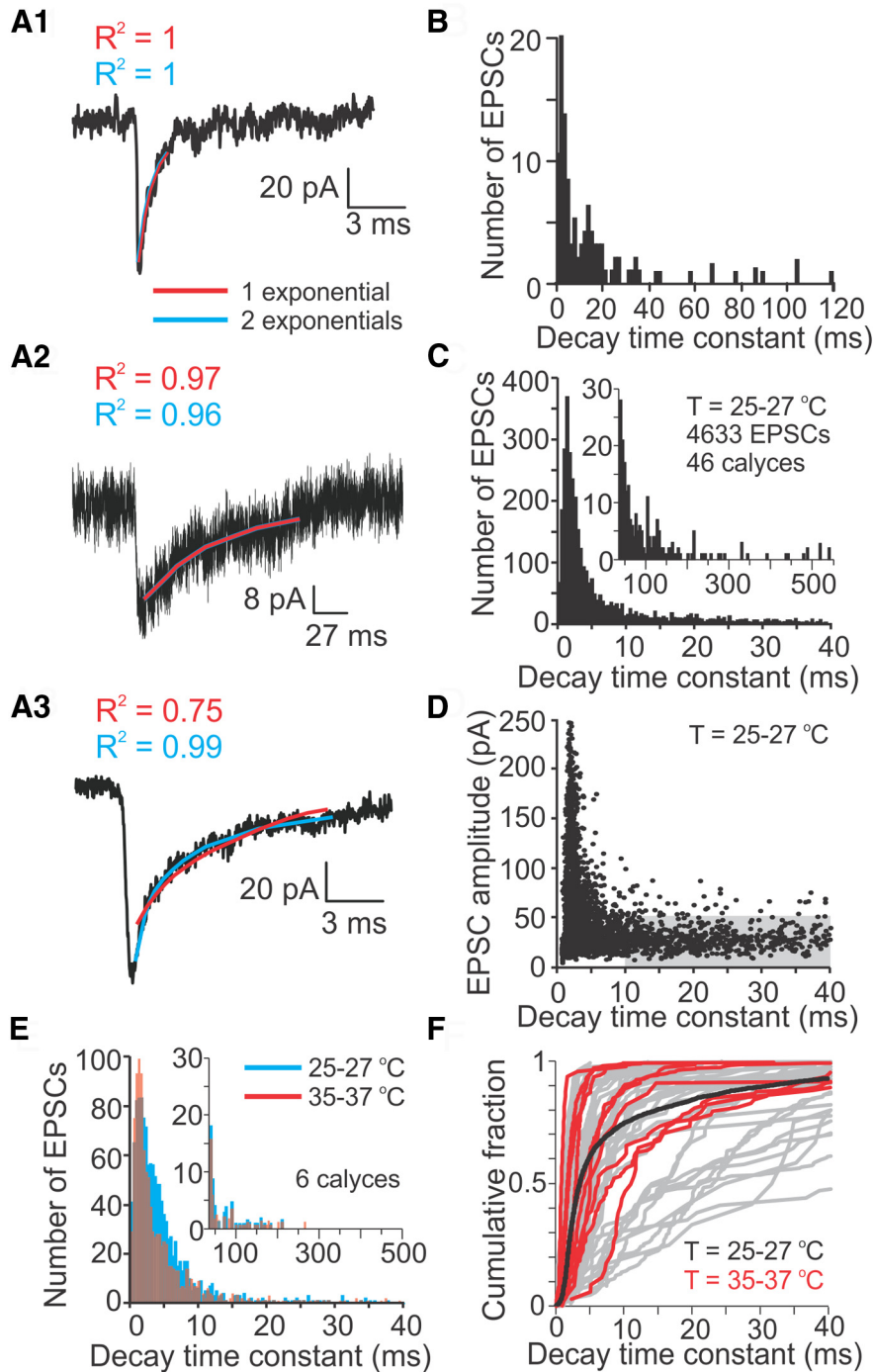


Figure 3. EPSC time course of decay shows fast and also unusually slow components. **A**, Example EPSCs from an individual calyx recording at room temperature show a faster EPSC (**A1**) and a slower EPSC (**A2**), both with decay times best fit with a single exponential and an EPSC with a decay time best fit with two exponentials (**A3**). **B**, The distribution of τ_{decays} (single exponential fit) for the calyx recording in **A** exhibits a tail with unusually slow $\tau_{decays} > 10$ ms. **C**, Distribution of τ_{decays} for pooled EPSCs from 46 calyx recordings at room temperature shows a tail with unusually slow $\tau_{decays} > 10$ ms. Inset, Distribution for values > 40 ms. **D**, EPSC amplitudes as a function of τ_{decay} , for individual EPSCs recorded from 46 calyces at room temperature. The gray box marks EPSCs that were considered small (< 50 pA) and slow ($\tau_{decay} > 10$ ms). **E**, Comparison of EPSC τ_{decays} recorded at room and body temperature. Distributions pooled from six recordings. At body temperature, slow events persist. **F**, Cumulative fraction plots of τ_{decays} for individual calyces recorded at room temperature (gray) and their average (black), and at body temperature (red).

blocker TBOA to block uptake of synaptic glutamate. To raise the baseline release rate, ATP was applied during control conditions, drug application, and washout. ATP activates P2X receptors on type I hair cells and has a depolarizing effect (Rennie and Ashmore, 1993). In addition, ATP also induced an inward current in

the calyx (data not shown). This inward current was present in all conditions and did not affect the shape of the EPSCs. During application of 200 μ M TBOA, the average EPSC slowed down reversibly (Fig. 4B1) and the distribution of τ_{decays} showed an increase in slower events (Fig. 4B2). For the example recording shown, the average τ_{decay} increased during TBOA application from 5.7 ± 1.2 ms ($n = 105$ EPSCs) to 10.1 ± 1.7 ms ($n = 111$ EPSCs) and decreased after washout to 6.7 ± 0.9 ms ($n = 102$ EPSCs; one-way ANOVA with *post hoc* Tukey test, $p < 0.03$ for TBOA compared with control and washout; Fig. 4B2, inset). For the population of recorded calyces, the average τ_{decay} increased by $\sim 40\%$ from 4.7 ± 0.6 ms to 6.6 ± 1.0 ms ($n = 11$, paired *t* test, $p = 0.007$; Fig. 4F).

Finally, a competitive blocker of glutamate receptors, kynurenic acid (KynA), was used to block the receptors that were reached by a lower glutamate concentration, thereby blocking the effect of spillover (Cadetti et al., 2008). Again, ATP was applied at all times to raise the baseline release rate of glutamate. During application of 250 μ M KynA the average EPSC showed faster τ_{decays} (Fig. 4C1) and the distribution of τ_{decays} showed fewer slower events (Fig. 4C2) compared with control. For the example recording shown, the average τ_{decay} decreased from 9.8 ± 0.9 ms ($n = 390$ EPSC) to 6.3 ± 0.7 ms ($n = 164$ EPSC) and returned to control values after washout (8.9 ± 0.7 ms, $n = 201$ EPSCs; one-way ANOVA with *post hoc* Tukey test, $p < 0.004$ for KynA compared with control and washout; Fig. 4C2, inset). For the population of recorded calyces, the average τ_{decay} of EPSCs decreased by $\sim 40\%$ from 5.9 ± 0.9 ms to 3.6 ± 0.7 ms in KynA ($n = 8$, paired *t* test, $p = 0.0001$; Fig. 4F).

Direct comparison of normalized average EPSCs recorded from the same calyx during TBOA or KynA application showed that τ_{decay} of the average EPSC during TBOA application could be best fit by two exponentials (τ_{decays} of 1.4 ms and 11.9 ms, $R^2 = 0.99$ vs 0.90, for a single exponential fit), whereas τ_{decay} of the average EPSC during KynA application could be best fit by a single exponential (τ_{decay} of 2.5 ms, $R^2 = 0.99$ for both fits; Fig. 4D). When plotting τ_{decay} with drug versus τ_{decay} in control for all experiments, results recorded in TBOA and KynA were separated by the unity line (Fig. 4E).

When the analysis for all three experimental approaches (application of elevated $[K^+]$, TBOA, or KynA) was repeated with fitting EPSC decay times with two exponentials, results were consistent with the single exponential analysis. Interestingly, com-

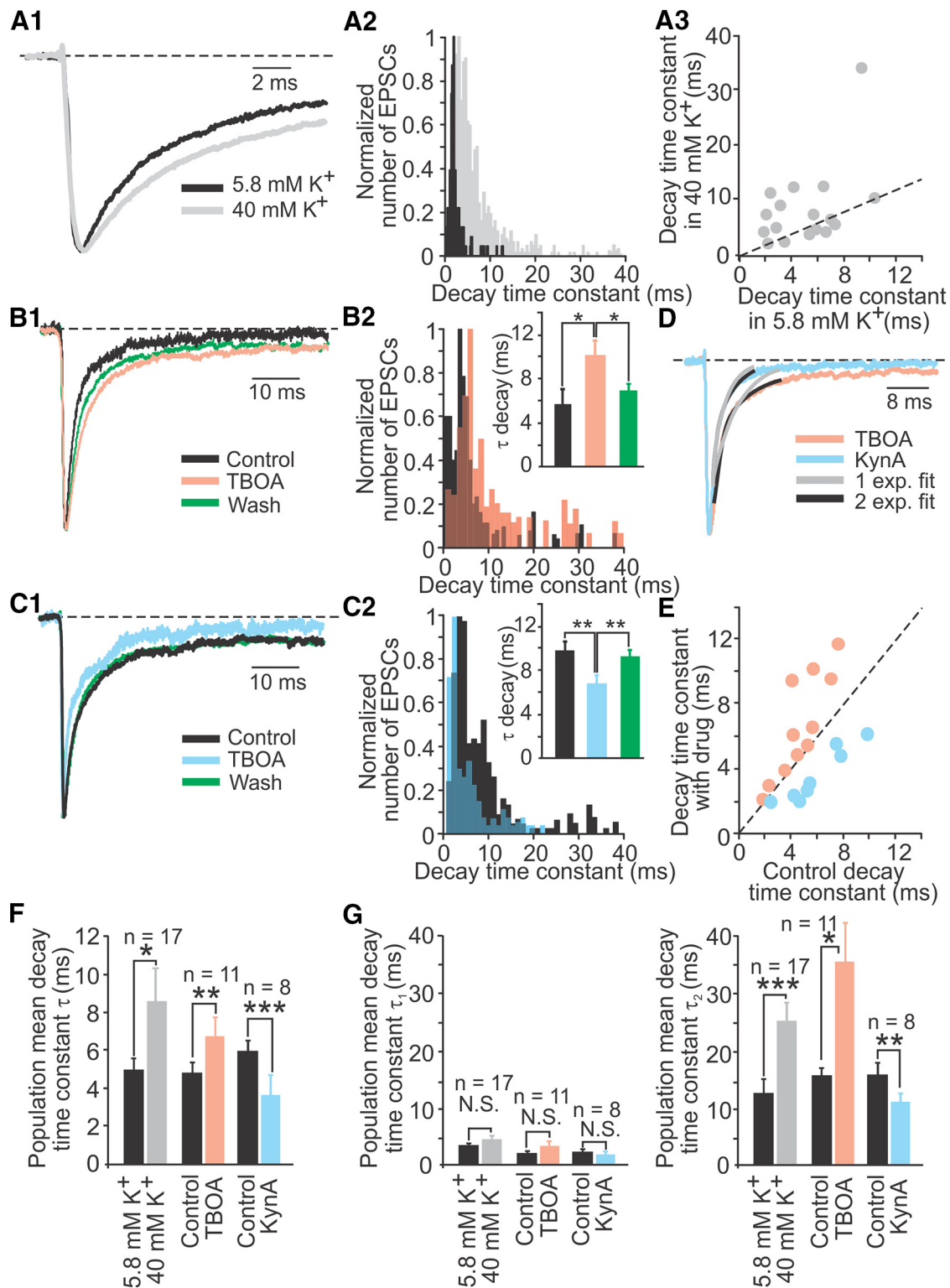


Figure 4. Glutamate accumulation and spillover contribute to slowing of EPSCs in calyx afferents. **A**, During application of 40 mM K^+ , the normalized average EPSC from an example calyx recording showed slower kinetics (**A1**) with an increase in τ_{decay} (**A2**). Nine of 17 calyces showed an increase in τ_{decay} (**A3**). The dashed line represents the unity line. **B**, In 200 μ M TBOA, the normalized average EPSC from an example calyx recording slowed down reversibly (**B1**) and the distribution of τ_{decays} (**B2**) showed an increase in larger τ_{decays} , resulting in a significant increase in the mean τ_{decay} (**B2**, inset). **C**, In 250 μ M KynA, the normalized average EPSC from an example calyx recording became reversibly faster (**C1**) and the distribution of τ_{decays} (**C2**) showed an increase in smaller τ_{decays} , resulting in a decrease in the mean τ_{decay} (**C2**, inset). **D**, An example recording showing a faster one exponential decay for the average EPSC during KynA application compared with a two exponential decay during TBOA application. **E**, Changes in τ_{decay} during TBOA or KynA application. **F**, Single exponential fits and (**G**) double-exponential fits of τ_{decays} for the population of recorded calyces show significant changes for all conditions applied. *n*, Number of recordings per condition; * $p < 0.05$, ** $p < 0.01$, *** $p = 0.0001$.

Table 2. EPSC kinetics before and during application of extracellular solution with 40 mM K⁺ (*n* = 17 recordings), 200 μM TBOA (*n* = 11 recordings), or 250 μM KynA (*n* = 8 recordings)

	Control	High [K ⁺]	<i>p</i> value	Control	TBOA	<i>p</i> value	Control	KynA	<i>p</i> value
τ ₁	3.2 ± 0.4	4.2 ± 0.6	0.08	1.9 ± 0.4	3.2 ± 0.8	0.07	2.2 ± 0.4	1.8 ± 0.4	0.15
τ ₂	12.2 ± 2.5	24.6 ± 3.2	0.0007*	15.3 ± 1.4	34.9 ± 6.7	0.02*	15.5 ± 1.8	10.9 ± 1.7	0.006*
A ₂ /(A ₁ + A ₂)	0.4 ± 0.06	0.4 ± 0.05	0.4	0.4 ± 0.05	0.4 ± 0.04	0.4	0.5 ± 0.06	0.4 ± 0.05	0.08

*Significant *p* values for paired *t* test.

A₂/(A₁ + A₂) shows relative contributions of τ_{1decay} and τ_{2decay} (see Materials and Methods).

pared with controls, significant changes were not found for τ_{1decay}, but for τ_{2decay}. τ_{1decay} stayed at 2–4 ms for all controls and drugs tested. However, τ_{2decay}, at 12–16 ms in control, increased in elevated [K⁺] and in TBOA by 100–130% and decreased in KynA by ~30% (Fig. 4G; Table 2).

During the experiments with elevated [K⁺] or TBOA, one factor that could have shaped the EPSC waveform is enhanced glutamate receptor desensitization due to glutamate accumulation (Trussell et al., 1993). Individual EPSCs in calyx afferents were affected by desensitization: in 5.8 mM K⁺ external solution, both τ_{1decay} and τ_{2decay} increased approximately three times in 100 μM cyclothiazide (CTZ), a drug that removes desensitization from AMPA receptors [Yamada and Rothman, 1992; control vs CTZ (*n* = 26 vs 47 EPSCs): τ₁ = 1.34 ± 0.14 vs 4.28 ± 0.37 ms (*t* test, *p* < 0.0001) and τ₂ = 9.32 ± 2.03 vs 33.42 ± 4.81 ms (*p* = 0.0005)]. However, in elevated [K⁺] or TBOA, no effect was found for τ_{1decay}, and a change in τ_{1decay} would have been expected if desensitization had caused EPSC waveform changes in these conditions. Together with the most direct evidence, a decreased τ_{2decay} in KynA, the most likely explanation for the results of all three experimental approaches is that glutamate accumulation and spillover occurred to shape EPSCs with slow-waveform components.

Not all glutamate receptor patches in the calyx are juxtaposed to presynaptic ribbons

To investigate whether the relative location of presynaptic release sites (hair cell ribbons) and postsynaptic AMPA receptor patches (on afferent calyces) somehow favors the occurrence of spillover, vestibular cristae from rats aged P17–P21 were isolated and labeled with antibodies against CTBP2 (a marker of presynaptic ribbons) and glutamate receptor subtypes GluA2/3 (a marker of postsynaptic AMPA receptors). Additionally, an antibody against Tubulin J (TuJ) was used to label the cytoplasm of afferent fibers and terminals to identify calyces by their typical shapes. Figure 5A shows a single optical section of four hair cells from the central zone of an anterior crista. Figure 5A, inset 1 shows immunolabeled puncta, CTBP2 (green), and GluA2/3 (red), at the inner face of a TuJ labeled calyx (blue). Between calyces, a type II hair cell can be identified by the presence of its nucleus without a surrounding calyx label and with CTBP2 and GluA2/3 immunolabeled puncta at its base, marking its bouton-type ribbon synapses (Fig. 5A, inset 2). For the present study, only the labels at the type I hair cell/calyx synapse, at the inner face of the TuJ labeled calyx, were quantified.

To quantitatively assess the relative position of presynaptic ribbons and postsynaptic AMPA receptor patches in type I hair cell/calyx synapses, first the total number and spatial locations of CTBP2 and GluA2/3 immunolabels were determined from 3D reconstructions from stacks of confocal micrographs from 17 counts of single or complex calyces (2–3 calyces each) located in the central zones of cristae of nine animals from three litters (4–7 calyces/animal giving a total of 36 calyces). The average number

of CTBP2 and GluA2/3 labels counted were not different (38.9 ± 3.7 and 32.5 ± 4.2, *n* = 17, *t* test, *p* = 0.25). The average number of CTBP2 labels per hair cell was calculated as 19.0 ± 1.3 and that of GluA2/3 labels per calyx was calculated as 15.5 ± 1.5. Inspection by eye revealed that CTBP2 and GluA2/3 labels were often not juxtaposed. As shown in Figure 5A and C, in some cases single GluA2/3 and CTBP2 immunolabels appeared juxtaposed (arrowhead, magnified inset 4), supporting fast activation of glutamate receptors. Sometimes GluA2/3 immunolabels were located in the vicinity of CTBP2 immunolabeled puncta (Figs. 5A, C, white arrows, magnified insets 1 and 6), potentially allowing glutamate receptor activation through spillover, with possible direct activation as well. Finally, some GluA2/3 immunolabels appeared alone with no closely related CTBP2 immunolabels or vice versa (blue arrows, magnified insets 1, 4, 5, and 6).

To quantify the juxtaposition of presynaptic and postsynaptic immunolabels objectively, the distance between the centers of immunolabels was calculated. These values were compared with equivalent measurements from immunolabeled cochlear inner hair cell (IHC) afferent synapses (Fig. 5B, inset 3), where CTBP2 and GluA2/3 immunolabels are known to be juxtaposed in a near 1:1 relationship (Khimich et al., 2005; Liberman et al., 2011), as it was also observed in the experiments presented here. Figure 5D (red) shows that in the IHC area ~90% of CTBP2 patches were <1 μm away from a GluA2/3 patch and vice versa. In contrast, <50% of GluA2/3 immunolabels were within 1 μm of a CTBP2 immunolabel and vice versa in the vestibular crista (Fig. 5D, blue). Note, that in control experiments in both vestibular crista and the organ of Corti ~90% of GluA2/3 antibodies were colocalized with PSD95 immunolabels (data not shown), a structural protein found in glutamatergic postsynaptic densities in general (Kennedy, 1997; Béique et al., 2006), suggesting that GluA2/3 detects most glutamatergic postsynaptic densities in calyx afferents. The calculated distances between CTBP2 and GluA2/3 immunolabels do not necessarily reflect absolute distances between presynaptic ribbons and postsynaptic AMPA receptor patches (with deviations likely resulting from the inherent limits of resolution imposed by immunofluorescence and confocal microscopy).

In summary, the complex spatial relationship between presynaptic and postsynaptic immunolabels in the vestibular crista compared with the cochlea can explain the variable shapes of EPSCs with fast and/or slow components of decay. The histological data are consistent with the hypothesis that glutamate spillover plays an important role in activating EPSCs. The large variability in the relative localization of presynaptic and postsynaptic sites, within and between calyces, may account for the variability of EPSC waveforms within and between recordings.

Glutamate accumulation and spillover increases the firing rate of calyx afferents

To study the functional implications of glutamate accumulation and spillover at the type I hair cell/calyx synapse, calyx AP firing rates and membrane potentials were examined in current-clamp.

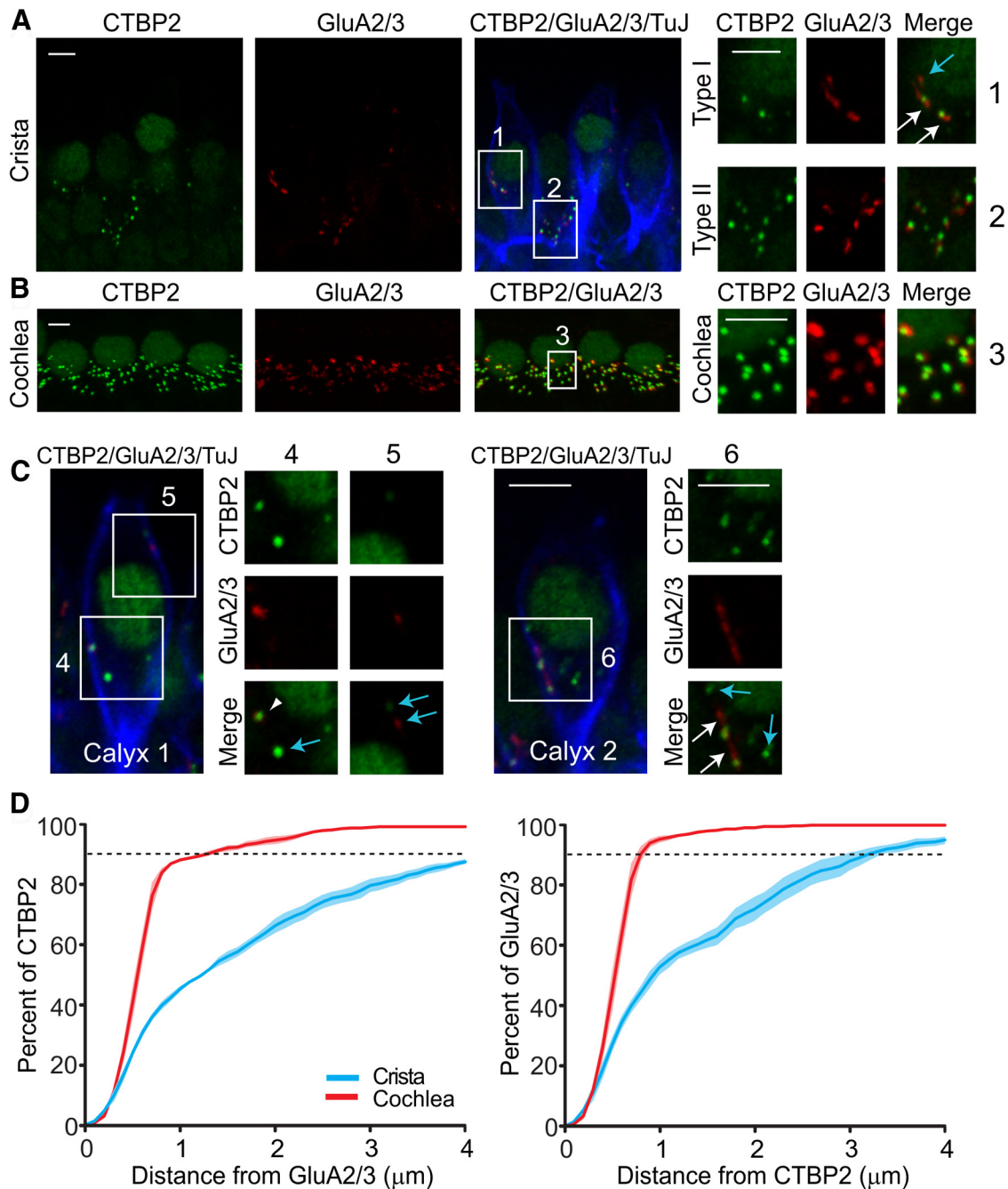


Figure 5. Structural evidence is consistent with glutamate spillover in the synaptic cleft. **A**, High-magnification micrographs of single optical sections from the central region of the horizontal crista show presynaptic ribbons (anti-CTBP2, green) and postsynaptic glutamate receptors (anti-GluA2/3, red) associated with afferent endings (anti-TuJ, blue). Scale bar, $10 \mu\text{m}$. Insets, Type I hair cell/calyx afferent synapses (inset 1) and type II hair cell/bouton afferent synapses (inset 2). White arrows, GluA2/3 close to CTBP2 immunolabeled puncta. Blue arrow, GluA2/3 immunolabels with no closely related CTBP2 immunolabels. Scale bar, $3 \mu\text{m}$. **B**, High-magnification micrographs of z-projections from the apical turn of the organ of Corti. Scale bar, $10 \mu\text{m}$. Inset 3 shows a close to 1:1 relationship between presynaptic and postsynaptic labels. Scale bar, $3 \mu\text{m}$. **C**, High-magnification micrographs of two example calyces. Some GluA2/3 immunolabels were juxtaposed to CTBP2 immunolabels (inset 4, arrowhead). Sometimes GluA2/3 immunolabels were located in the vicinity of CTBP2 immunolabeled puncta (inset 6, white arrows), whereas other GluA2/3 immunolabels appeared alone with no closely related CTBP2 immunolabels or vice versa (insets 4, 5, and 6, blue arrows). Scale bar, $3 \mu\text{m}$. **D**, Comparison of the distances between CTBP2 and GluA2/3 immunolabels. Light colors represent SE.

Again, for a higher baseline level of release, $100 \mu\text{M}$ ATP was applied throughout the experiment. Application of $200 \mu\text{M}$ TBOA repeatedly increased the firing rate of afferents from 8.2 ± 5.1 spikes/s to 30.1 ± 10.3 spikes/s ($n = 5$, paired t test, $p = 0.03$; Fig. 6A). The shape of the average AP in control (black) compared with TBOA (red) was not different (Fig. 6B). However, in TBOA, the membrane potential depolarized gradually over sec-

onds by a maximum value of ~ 5 mV, as shown in the example in Figure 6A.

To explore this phenomenon in a more controlled fashion, APs were blocked with $1\text{--}2 \mu\text{M}$ TTX. Then, either ATP alone, ATP and TBOA, or ATP and KynA were applied for ~ 100 s or until the response reached a plateau. Between drug applications, external solution (no ATP) was applied for washout. These

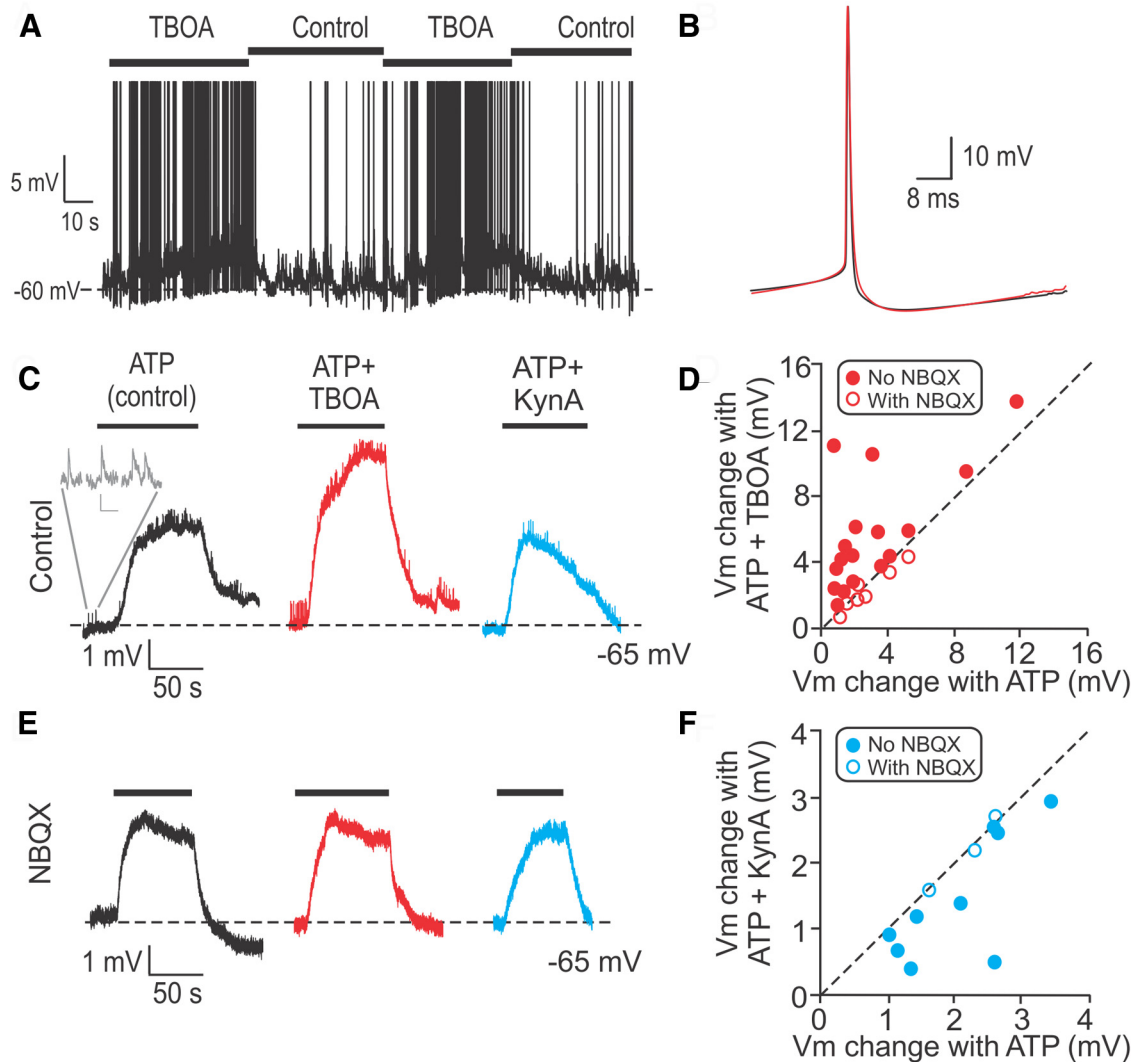


Figure 6. Glutamate accumulation and spillover increase the firing rate of calyces. **A**, In an example calyx recording, during application of 200 μM TBOA, the membrane potential depolarized and the firing rate increased from 2.9 ± 0.7 spikes/s to 12.1 ± 7.3 spikes/s. In TBOA the membrane potential depolarized by ~5 mV. **B**, The AP shape did not change during TBOA application (red). **C**, Comparison of changes in membrane potential during individual ATP applications (to increase release rate from hair cells) without (control) and with TBOA or KynA. Throughout the experiment, 2 μM TTX was applied to block AP generation. Inset (gray), Example EPSPs. Scale bar: 0.2 mV, 100 ms. In control, ATP causes depolarization mediated by P2X receptors on the calyx membrane. **D**, Most calyces were further depolarized during TBOA application compared with control (filled symbols above dashed unity line). Empty symbols: TBOA and NBQX application. **E**, During application of 10 μM NBQX, the effects of TBOA and KynA shown in **C** were blocked. Same calyx as in **C**. **F**, Most calyces showed less depolarization during KynA application compared with control (filled symbols below dashed unity line). Empty symbols: KynA and NBQX application.

shorter instead of continuous ATP applications were used because prolonged ATP applications resulted in decreased rates of EPSPs. ATP application depolarized calyces due to P2X receptor activation (Fig. 6C, black trace). This depolarization served as the control value. Additionally, ATP depolarized the hair cell and increased its rate of release, resulting in higher EPSP rates in the calyx. Calyces showed an increased depolarization during application of TBOA and ATP compared with ATP alone (Fig. 6C). On average, responses increased by 5.0 ± 0.7 mV (range: 1.5–11.2 mV) compared with the average depolarization of 2.3 ± 0.6 mV (range: 0.8–8.8 mV) in ATP alone (paired *t* test, $p = 0.002$, $n = 17$; Fig. 6C). All of the 17 calyces tested showed further depolarization during application of TBOA and ATP, placing all data points above the unity line when plotting the change in membrane potential in TBOA and ATP compared with ATP alone (Fig. 6D, filled symbols). In contrast, most calyces showed a decreased depolarization during application of KynA and ATP compared with ATP alone (Fig. 6C). On average, responses de-

creased from 1.9 ± 0.3 mV to 0.8 ± 0.5 mV (paired *t* test, $p = 0.02$, $n = 9$), placing most data points below the unity line when plotting the change in membrane potential in KynA and ATP compared with ATP alone (Fig. 6F, filled symbols). Importantly, when these experiments were repeated in the presence of 10 μM NBQX (Fig. 6E), depolarizations in ATP alone compared with TBOA and ATP (2.8 ± 0.5 mV vs 2.4 ± 0.5 mV, paired *t* test, $p = 0.08$, $n = 7$), as well as depolarizations in ATP alone compared with KynA and ATP (2.2 ± 0.3 vs 2.2 ± 0.4 mV, $n = 3$, paired *t* test, $p = 0.8$) were not significantly different (Fig. 6D,F, open symbols). Together, these results suggest that the changes in membrane depolarization in response to TBOA or KynA application were mediated by AMPA receptors in response to glutamate accumulation and spillover.

Discussion

The present study characterizes AMPA receptor-mediated synaptic transmission at the highly specialized synapse between the

vestibular type I hair cell and its afferent calyx ending. Results suggest that glutamate accumulation and spillover in the synaptic cleft lead to a wide range of EPSC kinetics, including unusually slow synaptic events. These effects induce slow depolarizations of the postsynaptic membrane resulting in changes in firing rate.

Comparison of EPSC amplitudes at the vestibular hair cell/calyx synapse and other ribbon synapses

Mean EPSC amplitudes of 60–80 pA at a holding potential of -103 mV were similar to measurements reported previously in the calyx (Rennie and Streeter, 2006; Dulon et al., 2009; Eatock and Songer, 2011), taking into account different recording conditions. EPSC amplitude distributions were well fit by a Gaussian, with a peak at ~ 20 pA, most likely representing the preferred release of individual vesicles, in contrast to the preferred release of multiple vesicles as seen at other hair cell ribbon synapses (Glowatzki and Fuchs, 2002; Goutman and Glowatzki, 2007; Li et al., 2009; Schnee et al., 2013). EPSC amplitudes increased with hair cell depolarization. This phenomenon could be explained by enhanced multivesicular release (Wadiche and Jahr, 2001) similar to changes in EPSC amplitude that occur with depolarization at ribbon synapses in the frog inner ear (Li et al., 2009) and the retina (Singer et al., 2004). Larger EPSC amplitudes may result in larger EPSPs and may increase the likelihood of AP generation in the calyx with hair cell depolarization.

Glutamate accumulation and spillover result in slow EPSC kinetics at the hair cell/calyx synapse

EPSC waveforms recorded in calyx afferents had an unusually wide range of time courses, with τ_{decays} ranging over two orders of magnitude, from <5 to >500 ms, as determined by single exponential fits. For a subset of EPSCs, decay times were best fit with two exponentials, suggesting that more than one mechanism might be at play. When decay times were analyzed with two exponential fits, a faster average $\tau_{1\text{decay}}$ in the millisecond range and a slower average $\tau_{2\text{decay}}$ in the tens of millisecond range were found. At body temperature, the slower components of the EPSC waveforms remained, suggesting that this phenomenon also occurs under physiological conditions.

Several lines of evidence support the hypothesis that glutamate accumulation and spillover highly contribute to slowing EPSC kinetics. First, when the glutamate concentration was increased in the synaptic cleft by either increasing the hair cell release rate via increasing external $[K^+]$ or by decreasing glutamate uptake via blocking glutamate transporters with TBOA, the number of slower EPSCs and the average τ_{decay} increased. Second, a competitive blocker of glutamate receptors, KynA that blocks receptors activated by spillover, decreased the number of slower events and the average τ_{decay} . When the analysis for the experimental approaches with elevated $[K^+]$, TBOA, or KynA was repeated with two exponential fitting of the EPSC decay times, only the slower $\tau_{2\text{decay}}$ but not the faster $\tau_{1\text{decay}}$ was found to be affected. Therefore, it is unlikely that mechanisms like glutamate receptor desensitization (Trussell et al., 1993) or a changed level of multivesicular release (Yamada and Rothman, 1992; Wadiche and Jahr, 2001; Singer et al., 2004) underlie the described changes in elevated $[K^+]$, TBOA, or KynA, as for these mechanisms a change in the faster $\tau_{1\text{decay}}$ in the millisecond range would have been expected. Third, presynaptic ribbons and postsynaptic glutamate receptor patches were only partially juxtaposed at the type I hair cell/calyx synapse, favoring scenarios where glutamate might reach postsynaptic receptors by spillover.

Previous studies in other brain areas have shown that spillover can activate extrasynaptic NMDA, AMPA, and metabotropic receptors (Kullmann et al., 1996; Barbour and Häusser, 1997; Scanziani et al., 1997; Oliet et al., 2001; Szapiro and Barbour, 2007; Hires et al., 2008) and can play a role as an adjunct to conventional synaptic transmission. However, spillover has also been described as the main mode of synaptic transmission between the climbing fiber and molecular layer interneurons in the cerebellar cortex (Szapiro and Barbour, 2007) and between mitral cells in the olfactory bulb (Isaacson, 1999). Similarly, data presented here suggest a prominent role of spillover at the vestibular hair cell/calyx synapse during its normal mode of operation, which includes high rates of release, even at rest (Yang and Hullar, 2007; Lasker et al., 2008).

The slow kinetics of EPSCs at the hair cell/calyx synapse with a $\tau_{1\text{decay}}$ of 2–4 ms and a $\tau_{2\text{decay}}$ of 12–16 ms is in strong contrast to the fast kinetics of EPSCs at the IHC afferent synapse in the auditory pathway, with a mean τ_{decay} of 0.5 ms and no second, slower component (Grant et al., 2010). At the IHC afferent synapse, fast events are critical for conserving timing and precision to allow for coding of a wide range of frequencies (up to 111 kHz in bats; Kössl and Vater, 1985) and of small interaural time differences for directional hearing (Carr and Konishi, 1990). For this purpose, an afferent ending receives input from a single ribbon synapse (Liberman, 1980) and individual large multivesicular EPSCs activate individual APs (Yi et al., 2010; Rutherford et al., 2012), avoiding any form of summation. In contrast, the semicircular canals convey head movements <20 Hz and the type I hair cell/calyx synapse seems to be designed to rather integrate by receiving inputs from multiple ribbons (Lysakowski and Goldberg, 2008). Due to spillover, slow events, and summation, slow depolarizations occur postsynaptically and this integrated signal changes AP firing rates.

Other possible contributors to EPSC kinetics at the calyx

Additional mechanisms may contribute to slow depolarizations of the calyx afferent. Several studies have provided evidence for K^+ accumulation in the synaptic cleft between type I hair cell and calyx (Holt et al., 2007; Lim et al., 2011; Contini et al., 2012). K^+ accumulation could induce calyx depolarization via hair cell depolarization and enhanced transmitter release and/or by directly inducing calyx depolarization. Additionally, buildup of H^+ in the synaptic cleft of the turtle hair cell/calyx synapse has been shown to cause slow changes in postsynaptic currents (Highstein et al., 2014). In the present study, the effects of glutamate accumulation and spillover on the calyx membrane potential, as tested by application of TBOA and KynA, were completely blocked by the AMPA receptor blocker NBQX, suggesting that these effects were mediated by glutamate, but not K^+ or H^+ accumulation. Further investigations are needed to characterize possible relative contributions of glutamate, K^+ or H^+ accumulation in signaling at hair cell/calyx synapses in different species and vestibular endorgans.

Experimental conditions for the present study were specifically chosen so that the effects of glutamate accumulation could be measured. The hair cell was depolarized with ATP, for enough glutamate release to occur to induce spillover; on the other hand, release rates had to be kept low enough, so that individual EPSC waveforms were not obscured by summation. With TBOA, glutamate accumulation in the synaptic cleft was enhanced artificially. *In vivo*, at body temperature, the players involved may work together differently compared with the experiments *in vitro*. For example, firing rates of afferents *in vivo* are higher (~ 50

spikes/s in mice; Yang and Hullar, 2007; Lasker et al., 2008) compared with *in vitro* conditions used here (~25 spikes/s). In fact, the high firing rates at rest found *in vivo* may be partially due to depolarization based on a steady-state current activated by desensitized receptors that are constantly exposed to a low glutamate concentration due to spillover (Trussell et al., 1993; Otis et al., 1996).

Functional significance of glutamate accumulation and spillover at the calyx

Studies in different species have shown that vestibular afferents with calyx endings show an increase in their sensitivity with increasing frequency over the range of natural stimuli (<20 Hz; Fernández and Goldberg, 1971; Ramachandran and Lisberger, 2006), and therefore most likely function as “high-frequency” event detectors (Sadeghi et al., 2007). The increase over the low to mid frequency range of stimuli (<1 Hz) has been explained by the mechanics of the end organ, described by a torsion pendulum model (Fernández and Goldberg, 1971). For higher frequencies, an additional mechanism that acts between hair cell mechanotransduction and the spike generator in the calyx afferent needs to be introduced to explain the measured increase in sensitivity. We propose that such a mechanism may include increased glutamate release at higher frequencies of stimulation resulting in pronounced accumulation and spillover and further increase in firing rate. Indeed, a frequency dependent modulation of spillover has been observed in the hippocampus, where at high-frequencies of stimulation, submicromolar glutamate concentrations can persist along the surface of dendrites for hundreds of milliseconds (Hires et al., 2008).

Conclusions

Our results provide structural and functional evidence that glutamate spillover plays a prominent role in signal transmission at the hair cell/calyx synapse. In other brain areas, this mechanism is used to provide high pass properties (DiGregorio et al., 2002), synchrony (Isaacson, 1999), or gain compression (Oikonomou et al., 2012) of neuronal responses. We propose that spillover in the vestibular periphery may play a role in modifying afferent response sensitivity. Future experiments with better control of hair cell depolarization levels are required to directly evaluate the effect of glutamate accumulation and spillover on the sensitivity of afferents at different amplitudes and frequencies of stimulation.

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