Striated organelle, a cytoskeletal structure positioned to modulate hair-cell transduction

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The striated organelle (SO), a cytoskeletal structure located in the apical region of cochlear and vestibular hair cells, consists of alternating, cross-linked, thick and thin filamentous bundles. In the vestibular periphery, the SO is well developed in both type I and type II hair cells. We studied the 3D structure of the SO with intermediate-voltage electron microscopy and electron microscope tomography. We also used antibodies to α-2 spectrin, one protein component, to trace development of the SO in vestibular hair cells over the first postnatal week. In type I cells, the SO forms an inverted open-ended cone attached to the cell membrane along both its upper and lower circumferences and separated from the cuticular plate by a dense cluster of exceptionally large mitochondria. In addition to contacts with the membrane and adjacent mitochondria, the SO is connected both directly and indirectly, via microtubules, to some stereociliary rootlets. The overall architecture of the apical region in type I hair cells—a striated structure restricting a cluster of large mitochondria between its filaments, the cuticular plate, and plasma membrane—suggests that the SO might serve two functions: to maintain hair-cell shape and to alter transduction by changing the geometry and mechanical properties of hair bundles.

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Air-cell stereocilia are important for mechanotransduction. Much is known about their structure, their mechanical properties, and their geometric arrangement (1–3). These cells are embedded in the cuticular plate, a dense structure composed of an actin gel (4) located at the apical end of the hair cell, which is thought to act as a rigid platform, helping stereocilia return to their resting positions after stimulus-evoked displacements (5, 6). What has never been investigated is how the cuticular plate might be stabilized by structures underneath it. Do these structures provide a foundation for the cuticular plate and the hair bundle? One such structure is the striated organelle (SO), also known as a laminated (or Friedman’s) body, which is a cytoskeletal lattice underlying the apicolateral hair-cell membrane and consisting of alternating thick and thin filament bundles. When first described in vestibular hair cells in the 1960s, this structure was thought to be a pathological feature (7, 8). It was later found to be a normal component of mammalian vestibular and cochlear inner, but not outer, hair cells (9–11). A striated structure, similar in position but differing in morphological details, has been described in other vertebrates (12–14). Slepecky et al. provided a description of the SO (10, 15, 16), including the periodicity of its filaments, their radial direction, attachment to the plasma membrane, and association with microtubules. Previous studies, even those based on conventional transmission electron microscopy (TEM) and deep-etch freeze fracture (9, 10, 15, 17, 18), did not provide a picture of the 3D structure of the organelle. Electron-microscopic tomography (EMT) provides a promising approach to study 3D structures. The approach has been used to study other structural features of hair cells (19, 20) and is used here to study the SO’s 3D architecture, localization, shape, size, prevalence, and its connections with neighboring structures. We also used immunocytochemistry to identify and localize one protein component and to trace the development of the SO over the first postnatal week. Our findings on SO structure and 3D architecture of the apical region in vestibular hair cells reveal previously undescribed aspects of hair-cell cytoskeletal organization and suggest new hypotheses about the hair bundle and mechanoelectric transduction.

Results

Material was examined with both conventional TEM and intermediate voltage electron microscopy (IVEM, 400 KeV). Results were similar in both cases. From these data and EMT software, we produced 3D reconstructions of the SO and the apical portion of three type I hair cells (cells 1–3) and one type II cell (cell 4) from the utricular macula. Extraintracellular hair cells were preferred because they are smaller than those from the striola, which allowed us to use a magnification high enough (5,000×) to see details, but low enough to get the whole apical hair cell in the field of view. Three reconstructions (cells 1, 3, and 4) were present, and the other (cell 2) encompassed the entire apical portion of the hair cell. We also obtained tomographic data, but no reconstructions, from one striolar type I cell (cell 5) and a second type II cell (cell 6). In each reconstruction, we modeled the thick and thin filament bundles of the SO, the stereociliary rootlets (SRs), the cuticular plate, subcuticular mitochondria, and the membranes of the hair cell and surrounding calyx (the afferent ending innervating a type I hair cell) to obtain a 3D model of the SO in relation to these other structures.

Structure of the SO

Each SO was composed of 35–40 thick bundles of filaments and an equal number of thin bundles (Fig. 1A). These bundles had thicknesses of 34.6 ± 1.25 nm (mean ± SD), n = 59, and 11.3 ± 0.45 nm, n = 54, respectively. Individual filaments within the thick and thin bundles measured ~10 nm and 6 nm, respectively. Thick bundles were located immediately subjacent to the cell membrane (Fig. 1A and B), and exhibited a periodicity of ~128 ± 3.1 nm (n = 56); thin bundles were located midway between adjacent thick ones. The bundles of intertwined filaments (Fig. 1A, Inset) extended radially, like spokes, from the plasma membrane ~130 nm into the hair cell (Fig. 1B). Along the length of the organelle, bundles could be oriented vertically (Figs. 1A and 2A and B) or spirally (Figs. 1 E and F and 2F). Sometimes both orientations were found in restricted regions of a single organelle.

In type I hair cells, several bundles extended upwards from their circumferential location below the cuticular plate and inserted into the apical membrane surrounding the kinocilium.
Two neighboring SRs merged, seemingly without interruption, and insert into the calyx membrane (Fig. 1). Note also the cross-bundled and thin (arrowheads) filament bundles are composed of several thinner, spiral-bundled filaments (e.g., 1–4), measuring ~10 nm and ~6 nm, respectively. Note also the cross-filaments (small arrows), which EM Immunogold studies indicate are likely spectrin (Fig. 5 B and C; and insets). (Scale bar, 0.2 μm.) (B) Cross-section through the neck of a type I cell showing radial distribution of thick (arrows) and thin (arrowheads) SO filament bundles. They extend from the hair cell membrane to a depth of ~130 nm into the cell. (Scale bar, 0.2 μm.) (C and D) Apical insertion of a thick SO filament bundle (arrows) adjacent to the kinocilium. The site of insertion is a dimpled area (arrowhead) between the kinocilium and nearby SRs. (D) A SO filament bundle angles downward (arrows), approaching the cell membrane and continuing into the neck of the type I cell. (Scale bar, 0.5 μm; also applies to E and F.) Details of hair-cell neck region in cell 2, obtained with SLICER mode in IMOD. (E) Thick filament bundles of the SO merge together (arrowshead). (Scale bar, 1 μm.) (F) Higher magnification of the area enclosed in the box in E shows possible connections (arrows) between thick filament bundles of the SO and the calyx membrane bridging the intercellular cleft (dashed white lines show locations of hair-cell and calyx membranes). (Scale bar, 0.25 μm.)

Cal, calyx; CP, cuticular plate; M, mitochondria.

Connections of the SO to the Kinocilium and Some Stereocilia. The linkage between the membrane surrounding the kinocilium and the SO was seen in several cells. On occasion, SO bundles were observed to connect to SRs. This link was seen in a tomogram section for cell 1 (Fig. 2A) and in the resulting model (Fig. 2B). Two neighboring SRs merged, seemingly without interruption, into separate SO thick bundles. A similar observation was made in serial TEM photos (Fig. S1). In both instances, the SRs continuous with SO filaments were located on the circumference of the hair bundle. The SRs were slightly thicker than their continuations with the SO (Fig. 2A), suggesting that the two elements may differ in protein composition, or that, if both are made up of the same proteins, the latter is bundled differently.

**Stereociliar Rootlets.** We measured the diameters of stereocilia and SRs across all rows of the hair bundle in two hair cells. Stereocilia nearest the kinocilium (rows 1–3), were significantly thicker than those in rows 4–10 (stereocilia: rows 1–3, 140.6 ± 5.5, n = 34 vs. rows 4–10, 116.0 ± 3.4, n = 114, P < 0.05; rootlets: 46.1 ± 1.7, n = 34 vs. 38.0 ± 1.9, n = 114, P < 0.05). In all type I hair cell reconstructions, the largest SRs, those nearest the kinocilium, bent to form an angle of 110° within the cuticular plate (cell 2) (Fig. 2C and Movie S2). A type I cell from the striolar region (Fig. 2C, Inset), oriented in an appropriate plane, showed several rootlets bending within the cuticular plate. These long rootlets, on passing through the cuticular plate (Fig. 2C, Inset), were organized into two groups (Fig. 2E) that converged toward separate areas on the hair-cell membrane opposite the kinocilium (the SR insertion area) (Fig. 2D). The arrangement is best seen by observing the SR insertion area from different angles, as in the movie (Movie S2). SRs further from the kinocilium exhibited less and less bending and most appeared to remain within the cuticular plate (Fig. 2F and Movie S2).

**Subcuticular Mitochondria.** Within the confines of the SO, there is a set of exceptionally large mitochondria compared with those in the rest of the type I cell or in type II cells. Because mitochondrial function (Ca2+ homeostasis and source of ATP) is related to overall size, we used our tomograms to measure the volumes and surface areas of mitochondria from the same portion of the cell (the subcuticular region, ~6 μm below the apical cell membrane) in type I and neighboring type II cells (Table S1). Table S1 indicates that mitochondria inside the SO in type I hair cells are two-times larger in surface area and three- to four-times larger in volume than those in type II cells. In one reconstruction (cell 1), a few SRs emerging from the underside of the cuticular plate ended on subcuticular mitochondria and appeared to be “tethered” to them (Fig. 3A and B). In cell 2, other mitochondria were intimately associated with long stretches of SO bundles (Fig. 3 C and D). We observed similar close associations in each of the other reconstructions: for example, cell 4 (Fig. 4D).

**SO in Vestibular Type II Hair Cells.** EM results from the partial reconstruction of a type II cell (cell 4) are shown in Fig. 4. The SO is more extensive in the type II hair cell than in type I cells. It is longer, broader in extent, and the thick bundles are wider (compare Figs. 4A–C and S1) and can extend deeper into the hair cell (~400 nm, compared with ~130 nm for type I hair cells). Spherical structures surround the SO and are more numerous than in type I cells. Merging and morphing filaments were observed. Viewing the model from a vantage point below the cuticular plate (Fig. 4D), we can see that the SO is not connected to stereocilia in this cell, but it is closely associated with four mitochondria along its breadth, similar to type I cells (Fig. 3 C and D).

**Protein Composition of the SO.** We have identified one protein in the SO (Fig. 5). Using an antibody to α-2 spectrin, we labeled the SO in vestibular hair cells in both confocal (Fig. 5A) and EM immunogold (Fig. 5B and C) experiments. Quantification of the EM label for α-2 spectrin indicated a nonhomogeneous distribution of gold particles, with a tendency for them to be located immediately adjacent to the thick filament bundles (Fig. 5C) (χ2 test of homogeneity, χ2 = 23.1, df = 7, P < 0.002). Western blots (Fig. S24) confirmed the presence of α-2 spectrin in the inner ear. The identities of the bands at 150 kD and 285 kD were confirmed as α-2 spectrin with mass spectrometry (Fig. S2B).
SO Formation in Relation to Development of the Ear. Using α-2 spectrin as a marker for the SO, we performed confocal immunofluorescent (Fig. 5 D–F) and TEM (Fig. 5 G–J) studies on developing rats during the first two postnatal weeks, a period of rapid hair-cell differentiation and afferent development (21). The SO was not present at birth (postnatal day 0, P0) (Fig. 5D), but developed rapidly during the first week, making its first appearance at P2 (Fig. 5 E, G, and H), and appearing to be fully developed by P6 (Fig. 5 F, I, and J). Because transducer currents develop between embryonic day (E)16 and E17 in the mouse (22), and perhaps 1–2 d later in rats, the initial development of transduction does not require the SO, but it may be required for hair-cell maturation.

Discussion

Previous studies provided descriptions of the SO (7–11, 15, 16, 18) but were unable to generate a 3D view of the structure. In this study we did so using EMT. Our reconstructions show that the SO of type I hair cells has the shape of an inverted, open-ended cone (a frustum), conforming to the shape of the hair-cell neck. This cone consists of 35–40 thick filamentous bundles and an equal number of thin bundles, with an apical extension near the kinocilium, connected to the neighboring apical hair-cell membrane. In the lower, constricted neck region, thick filament bundles occasionally merge with each other. The upper ends of at least a few thick circumferential filament bundles of the SO connect to SRs and several mitochondria appear tethered to SRs and to SO bundles. The lower portions of the SO appear to insert into the plasma membrane. These associations with other elements of the cell suggest that the SO may be involved in regulation of hair-cell transduction, as well as contribute to the characteristic shape of the type I hair cell. In the type II cell, the SO has a more planar appearance and is not connected to stereocilia or the apical membrane, suggesting a predominantly structural role.

Our data provide a fresh view of the relationship between the stereocillar bundle and the cuticular plate. In most accounts, individual stereocilia have been thought to insert into the cuticular plate and terminate there. The cuticular plate is then envisioned as providing a platform that contributes to the stereocilia returning to their normal upright positions after bending (6). That SRs actually traverse the cuticular plate has been noted only occasionally in the literature (17, 23). Our data are unique in showing that the SRs reach the plasma membrane opposite the kinocilium. Stabilization of the cuticular plate itself and how it might be anchored has only been adequately explained in cochlear outer hair cells: the cuticular plate extends to the apical-lateral wall of the cell (23–25), and stereocilia insert in the lateral cell membrane while still within the cuticular plate (23). This process is not the case in inner hair cells nor in vestibular hair cells (26). What provides a foundation for the cuticular plate and thus stabilizes its position? Jaeger et al. (27) showed that the cuticular plate in bullfrog hair cells was connected to the axial cytoskeleton through a well-defined bundle of microtubules. The present data show that the hair bundle and cuticular plate are even better fixed at three points on the cell membrane: the rootlets bend and extend through the cuticular plate, inserting into the plasma membrane at two points opposite the kinocilium, and the apical SO inserts into the apical cell membrane surrounding the kinocilium. These three attachments potentially form a tripod-like structure, coupling stereocilia and the kinocilium with subcuticular cytoskeletal elements, possibly affecting mechanotransduction.

The bending of SRs inside the cuticular plate is an intriguing finding. The only previous description of such bending is a single photomicrograph of a bent SR (26). There may be a molecular...
correlate for this bend. Actin is the major component of SRs (16, 23, 28). The particular angle of the bend in the SRs (110°) is similar to that produced by the actin-related proteins, Arp2/3 (actin-related protein 2/3 complex) (29–31) and coronin (32). In our material, only the longest and thickest rootlets, derived from stereocilia nearest the kinocilium and passing through the cuticular plate, exhibited this 110° angle; shorter rootlets (those that did not emerge from the cuticular plate) exhibited less acute angles.

Another prominent feature is the aggregation of large mitochondria in the subcuticular region of type I cells. Mitochondrial functional capacity is related to both overall size and internal structure (33). Such subcuticular aggregation of mitochondria with large volumes and surface areas, and “tethering” (34, 35) indicate the need for tight control of calcium homeostasis or that the SO, SRs, and other structures in the apical part of the hair cell may have high energetic requirements. Recent experiments in cochlear outer hair cells demonstrated that apical mitochondria can act to block Ca2+ diffusion into the hair-cell soma (36). In addition, mitochondrial cristata structure has recently been related to the high metabolic rate of synapses at the calyx of Held (33). Further studies can provide structural details relevant to the functional status of these apical mitochondria (e.g., ratios of cristae to outer membrane surface area, and analysis of cristata junction diameter and density).

Previous immunohistochemical studies have provided data on proteins in this part of the cell. The cuticular plate of vestibular hair cells is immunoreactive for actin, tropomyosin, and myosin (16, 37). Demémes and Scarfone (38) demonstrated fodrin (α-2 spectrin) immunoreactivity in the SO and cuticular plate and suggested that fodrin participates in a Ca2+-dependent cross-linking of actin filaments. Although we have not yet been able to confirm actin, we have confirmed by EM immunogold, Western blot, and mass spectrometry that α-2 spectrin is an integral, major component of the SO.

Physiological studies support a contractile hypothesis for something in the neck of the type I, but not type II, hair cell. K+ depolarization induces not only reversible shape changes in type I hair cells (39–41), but also a rise in cytosolic Ca2+ concentration (42) that could trigger such changes. Rising Ca2+ levels have been suggested as a potential substrate for mechanotransduction (3), with the large SO-associated mitochondria potentially playing a role (36, 43) by promoting reuptake of Ca2+ after mechano-transduction. A prior study (44) has shown that in the presence of local elevated calcium levels, fodrin (α-2 spectrin) can change from a diffuse distribution within the cytoplasm to a submembranous position, or a patch. If this were also the case in hair cells, it is conceivable that the SO could change its dimensions and rigidity depending on variations in intracellular Ca2+ concentration (with the latter regulated, in part, by the large subcuticular mitochondria), and thus influence hair bundle or cuticular plate movements, or tilting of the cell neck. Any of these actions could alter the sensitivity of the transduction apparatus, and thus, mechanotransduction (45). Rüsch and Thurm (46) showed that hair bundles at different sites on the sensory epithelium exhibit differences in amplitude and in the time course of deflection. Our data suggest that some of these differences could be because of variations in cuticular plate size, hair-bundle size, SO shape, cross-linking, or perhaps even mitochondrial size.

Our findings suggest a dual function for the SO in type I hair cells. One structural role is as a cytoskeletal specialization...
 responsible for the constricted neck characteristic of these hair cells (10, 15, 18). The architecture of the SO in type I and type II cells is one of many differences, besides $I_{K,L}$ (the outward potassium current present in type I hair cells), of a functional difference between type I and type II cells (47). Another potential structural role is to provide a tripartite foundation for the cuticular plate. Neither structural requirement is obviously present in type II hair cells, nor was a tripartite arrangement present in the one type II hair cell we reconstructed. Its columnar shape is similar to many columnar epithelial cells lacking an SO. Concerning a dynamic role, the overall architecture of the apical region in type I cells—a cluster of large mitochondria constrained between the SO and the cuticular plate, the plasma membrane and the adjacent calyx, the dense-cored vesicles—suggests that the SO is part of a larger functional complex regulating mechanotransduction or other physiological functions. Further analysis of the morphology, composition, and location, in combination with physiological studies, should shed light on its function.

Materials and Methods

Animals. Normal adult chinchillas (Chinchilla lanigera) were used for tomographic studies. Adult Long-Evans rats (Rattus norvegicus), weighing 230–330 g, were used for immunocytochemistry studies. For the development studies, 14 Long-Evans rat pups were studied. Two pups were killed on each day (beginning at birth, P0 to P6), one for immunofluorescent studies and the other for TEM. The Institutional Animal Care and Use Committee at the University of Illinois approved procedures involving animals.

EMT. We used serial sections of chinchilla utricular macula for conventional TEM and for IVEM. Our TEM methods were described previously (45). We analyzed a total of 41 tomograms using research facilities at the National Center for Microscopy and Imaging Research. For each reconstruction, a series of images at regular tilt increments was collected as described previously (30).

Three-Dimensional Reconstruction. Pixel sizes in our reconstructions were 2.8 nm and 1.96 nm. IMOD software (46) was used for alignment. Warped reconstructions were processed with TxBR software (47). Measurements were made with 3dmod (“Object info” feature). Stereocilia and rootlet diameters were measured at the same distance above the cell membrane (5.6 nm). Volumes and surface areas of mitochondria below the cuticular plate and within 6 μm of the apical surface of the cell were also measured. Movies were made with Amira software (version 5.2.1, Mercury/TGS).

Antibodies. Primary antibodies were: mouse anti–α-2 spectrin (Chemicon), goat anticalretinin (Chemicon). Secondary antibodies for confocal experiments were: goat antirabbit, antigoat, anti-mouse, or antigeen. All antibodies were used at 1:200 dilutions.

Fig. 5. The α-2 spectrin is one protein component of the SO. (A) Confocal microscopy shows cuticular plate (arrows) and SO (arrowheads) immunoreactivity for α-2 spectrin (red). Calretinin antibody (green) distinguishes type II (II) from type I (I) hair cells. (Inset) Higher magnification. Velocity reconstruction of α-2 spectrin label (red) in a type II hair cell shows the SO, which appears to hang down in two large flaps (arrowheads) from the cuticular plate (CP). (Scale bars, 2 μm.) (B) EM immunogold with an antibody to α-2 spectrin (gold particles) localizes the protein to the cuticular plate (arrowheads) in a type I cell (I) and to the SO (black box) in a type II cell (II). (Inset) Higher magnification of the area enclosed within the box in B. Gold particles (arrows) had a tendency to be located over the cross-links between the thick and thin filament bundles. (Scale bars, 0.5 μm.) (C) Quantification of the α-2 spectrin EM immunogold results. In 14 SO profiles, such as this example (SO) from a type II hair cell (II), gold particles were identified within an area delineated by a white line circumscribing the SO profile. Starting with each thick filament bundle, intervals between thick bundles were divided into eight equally spaced samples running parallel to the thick bundles, and the interval into which each particle fell was determined. (Inset) For 363 particles, sums in the eight sample bins are illustrated, superimposed upon a schematic of the SO. A $\chi^2$ test of homogeneity ($\chi^2 = 23.1, df = 7, P < 0.002$) indicated a preference for intervals immediately adjacent to the thick filament bundles. (D–F) Confocal microscopy of α-2 spectrin antibody in developing rat crista in the first postnatal week. (D) At birth (P0), there is no label in the cuticular plate (CP, short arrow), and only weak immunoreactivity in the upper half of the hair cell. (E) At P2, α-2 spectrin antibody labels the cuticular plate less intensely than at P6 (F), when it also labels parts of the lateral membrane in the region of the SO. (Scale bars, 5 μm.) (G–J) Normal TEM of developing rat hair cells. At P2 (G and H), the SO is just beginning to form in a hair cell of indeterminate type. The calyx ending, which defines type I hair cells, typically does not begin to form until P4 (21). (Scale bars in G and H, 0.5 μm.) At P6 (I and J), both the calyx (Cal) surrounding the type I cell and the SO (arrowheads) are well developed. Several stereocilia rootlets can also be observed within the cuticular plate in I. (Scale bars, 0.5 μm in I and 2 μm in J.)


More detailed descriptions of EM tomography, immunogold quantification, mass spectrometry, and Western blot analysis are given in SI Materials and Methods.

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Supporting Information

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SI Materials and Methods

Animals. Normal adult chinchillas (Chinchilla lanigera) were used for tomographic studies. Adult Long-Evans rats (Rattus norvegicus), weighing 230–330 g, were used for immunocytochemistry studies. For the development studies, 14 Long-Evans rat pups were studied. Two pups were killed on each day [from birth (postnatal day 0, P0) to P6]; one for immunofluorescent studies and the other for transmission electron microscopy (TEM). Procedures involving animals were approved by the Institutional Animal Care and Use Committee at the University of Illinois at Chicago.

Fixation for Tomographic Studies. Five adult chinchillas were deeply anesthetized with Nembutal (80 mg/kg) and perfused transcardially with 200 mL physiological saline containing heparin (2,000 IU) followed by 8 mL/g body weight of a triadethylene fixative, as described previously (1). After perfusion, vestibular epithelia were dissected in 0.1 M PB at 4 °C.

Electron Microscope Tomography. We used serial sections of chinchilla utricular macula for conventional TEM and for intermediate-voltage electron microscopy (IVEM). Our TEM methods have been described previously (1).

Semithin sections (0.5 µm) for IVEM were sectioned with a diamond Histoknife (Diatome) and serial sections were collected onto Formvar-coated or Formvar/Luxel-coated, carbon-coated, 1 × 2-mm single slot grids (Ted Pella). All sections were stained with 2% (wt/vol) uranyl acetate (10 min) and Sato lead stain (10 min), and then carbon-coated again. Fiducial cues consisting of 20-nm colloidal gold particles were deposited on one side of each section.

We analyzed a total of 41 tomograms using research facilities at the National Center for Microscopy and Imaging Research. For each reconstruction, a series of images at regular tilt increments was collected as described previously (2). Tilt series were recorded with a Gatan 4 K K × 4 K camera at 5,000 × magnification. We used a magnification of 5,000x to comfortably include the entire circumference of the apical part of type I hair cell, the proximal part of the hair bundle, and most of the type I hair cell constriction (neck) in the field of interest. The z-resolution of the tomographic volumes was 8 nm, compared with the 70-nm thickness of ultrathin sections, which allowed us to visualize objects at a much higher effective thickness of ultrathin sections, which allowed us to visualize objects at a much higher effective z-resolution depth resolution (8 nm vs. 70 nm) compared with conventional EM. Some elements were re-examined at high magnification (20,000–25,000×).

Three-Dimensional Reconstruction. The pixel sizes in our reconstructions were 2.8 nm and 1.96 nm. The IMOD software package (3) was used for rough alignment, and in some cases for fine alignment and reconstruction. Warped reconstructions were processed (fine alignment and reconstruction) using the TxBR software package (4). Serial tomograms of hair cells were joined using Etemo (IMOD) and four final volumes were obtained: three type I hair cells, (cell 1 from 6 serial tomograms, cell 2 from 11 serial tomograms, and cell 3 from 5 serial tomograms), and one type II hair cell (cell 4, from 9 serial tomograms). For each final volume (tomogram), segmentation was performed by manual tracing in the planes of highest resolution and by automated isosurface rendering with the program 3dmod (IMOD). Reconstructions were visualized using 3dmod. Cell 5, a striolar type I hair cell, and cell 6, an extrastriolar type II hair cell, were visualized with 3dmod as eight and three serial tomograms, respectively, but not segmented. The program allows stepping through slices of the reconstruction in any orientation (SLICER option) and tracking or modeling features of interest in any of the three dimensions. For quantification, measurements were made using 3dmod (“Object info” feature). Sterocilia and rootlet perimeters were each measured at the same distance above the cell membrane (5.6 nm), and the volumes and surface areas of mitochondria below the cuticular plate within 6 µm of the apical surface of hair cell were also measured. Movies of surface-rendered volumes and slices through the reconstructions were made using Amira software (version 5.2.1, Mercury/TGS).

EM Immunogold Quantification. To test if there were a nonuniform distribution of particles, we examined 14 striated organelles (SO) profiles from 12 hair cells (five type I and seven type II) and divided them into intervals. Starting with each thick filament, an interval was divided into eight equally spaced samples running parallel to the thick filaments, and we determined the interval into which each particle fell. A χ² test of homogeneity using “a single classification with equal expectations” (χ² = 23.1, df = 7, P < 0.002) indicated a preference for the intervals immediately adjacent to the thick filament.

Antibodies. Primary antibodies used were: mouse anti α-2 spectrin (Chemicon), goat anti-calretinin (Chemicon). Secondary antibodies for confocal experiments (Chemicon) were: Alexa 488-conjugated donkey anti-goat; Alexa-594-conjugated donkey anti-mouse. Secondary antibody for EM immunogold experiments (Aurion kit, EM Sciences) was: gold-conjugated rabbit anti-mouse. We used calretinin antibody as a marker of hair-cell type (5).

Immunohistochemistry. Three animals were used for immunohistochemical studies and three for Western blots. Fixation, confocal microscopy, and EM immunogold procedures were identical to those described in a recent study (5), which can be consulted for details. Western blots were done to verify that the hair-cell staining found in rat endorgans was α-2 spectrin. For Western blots, animals were anesthetized (Nembutal, 80 mg/kg, i.p.) and decapitated. Vestibular endorgans, cochlea, vestibular ganglion, brain (positive control), and vibrissae (negative control) were harvested from three adult rats within 10 min of killing. Methods were identical to those published previously (5), except that the ECL detection kit was obtained from GE Healthcare Life Sciences. Antibody incubation conditions were mouse anti-α-2 spectrin antibody (diluted 1:2,000) for 4 h at room temperature, washed 3 × 5 min in PBS-Tween, and HRP-conjugated goat anti-mouse IgG (diluted 1:30,000) for 1 h at room temperature. The identities of bands from Western blots were confirmed with mass spectrometry in the University of Illinois at Chicago Proteomics Laboratory. Scaffold software (v.3.1.4.1, Proteome Software) was used to validate mass spectrometry-based peptide and protein identifications.

Mass Spectrometry. Analysis was performed by the University of Illinois at Chicago Research Resources Center’s Mass Spectrometry, Metabolomics, and Proteomics Facility. The in-gel trypic digestion was performed according to the protocol described by Kinter and Sherman (6). Briefly, the gel bands were cut into 1-mm³ pieces, rinsed, and dehydrated. The protein was reduced with DTT and alkylated with iodoacetamide in the dark, before overnight digestion with trypsin at 37 °C in 50 mM ammonium bicarbonate. Peptides were concentrated and analyzed with a Thermo Orbitrap Velos mass spectrometer using a chip-based HPLC system (Agi-
lent Chip Cube) adapted to run on the Orbitrap Velos (7) using collision-induced dissociation fragmentation.

**Database Searching.** Tandem mass spectra were extracted by Readm.exe (version 4.0.2, Institute for Systems Biology), converted to the Mascot generic format using MzXML2Search and then submitted to a Mascot search engine (version 2.2.04). Charge state deconvolution and deisotoping were not performed. Mascot was set up to search the SwissProt 57.15 database (selected for *Rattus* taxonomy), assuming the digestion enzyme was trypsin. Mascot was searched with a fragment ion mass tolerance of 0.60 Da and a parent ion tolerance of 10.0 ppm. Oxidation of methionine, acetylation of lysine, and the N terminus and iodoacetamide derivative of cysteine were specified in Mascot as variable modifications.

**Criteria for Protein Identification.** Scaffold software (v. 3.1.4.1, Proteome Software) was used to validate MS/MS-based peptide and protein identifications. Peptide identifications were accepted if they could be established at greater than 95.0% probability as specified by the Peptide Prophet algorithm (8) and if they contained at least two identified peptides. Protein probabilities were assigned by the Protein Prophet algorithm (9). Proteins that contained similar peptides and could not be differentiated based on MS/MS analysis alone were grouped to satisfy the principles of parsimony.


**Fig. S1.** Serial section TEM of SOs. Ultrathin sections from a serial section study of type I vestibular hair cells in chinchilla crista. (A and B) Actin stereociliar rootlets (SR) in a striolar type I hair cell are directly continuous with stereocilia (arrowheads) and enter the SO (SO, short arrows). In the lower part of the neck, long arrows show insertions of the thick SO filaments into the cell membrane. Some microtubules (Mt, long, thin arrows) in A and C connect SRs to the SO below the central portion of the cuticular plate. Vesicular structures (asterisks in B and C) line up in parallel with microtubules (Mt) and the SO. (D) Section through the median plane of a type I hair cell, showing gray bands (short arrows) that lie between the thick filaments of the SO forming a narrow band along the cell membrane. Our EM immunogold results indicate this gray band is α-2 spectrin. (Scale bar in D, 1 μm; applies to all panels.) CP, cuticular plate; M, mitochondria.

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A. Western blot results.

**Fig. S2.** Antibody validation. (A) Immunoblot of rat brain (positive control for α-2 spectrin), vibrissae (negative control), vestibular end organs (VO), vestibular ganglia (VG), and rat cochleae labeled with an anti-α-2 spectrin antibody. The blot shows two spectrin bands. The upper band corresponds to the full-length α-2 spectrin protein (∼285 kDa), the lower band to a well-characterized proteolytic fragment at ∼150 kDa (1). (B) Mass spectrometry results. (Upper) Results of our mass spectrometry runs on the upper (285 kDa) band in our immunoprecipitation gels. With 46% coverage and 118 unique peptides, there is 95–100% confidence that the band is α-2 spectrin. (Lower) Results of mass spec runs on the lower, 150-kDa band in our gels. Again, with 43% coverage and 108 unique peptides, there is a similar level of confidence that this band is also α-2 spectrin.


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B. Mass spectrometry results.

**Upper band - 285 kDa**

SFF2_RAT (100k), 284.640 kDa

Spectrin alpha chain, brain CE/Rattus novegicus CIK-α-spectrin PE-1 SV-v2

118 unique peptides, 151 unique spectra, 271 total spectra, 1142/2472 amino acids (46% coverage)

**Lower band - 150 kDa**

SFF2_RAT (100k), 284.640 kDa

Spectrin alpha chain, brain CE/Rattus novegicus CIK-α-spectrin PE-1 SV-v2

108 unique peptides, 139 unique spectra, 275 total spectra, 1069/2472 amino acids (44% coverage)

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*Fig. S2.* Antibody validation. (A) Immunoblot of rat brain (positive control for α-2 spectrin), vibrissae (negative control), vestibular end organs (VO), vestibular ganglia (VG), and rat cochleae labeled with an anti-α-2 spectrin antibody. The blot shows two spectrin bands. The upper band corresponds to the full-length α-2 spectrin protein (285 kDa), the lower band to a well-characterized proteolytic fragment at ∼150 kDa (1). (B) Mass spectrometry results. (Upper) Results of our mass spectrometry runs on the upper (285 kDa) band in our immunoprecipitation gels. With 46% coverage and 118 unique peptides, there is 95–100% confidence that the band is α-2 spectrin. (Lower) Results of mass spec runs on the lower, 150-kDa band in our gels. Again, with 43% coverage and 108 unique peptides, there is a similar level of confidence that this band is also α-2 spectrin.

Table S1. Subcuticular mitochondrial dimensions

<table>
<thead>
<tr>
<th>Samples</th>
<th>Type I hair cells</th>
<th>Neighboring type II hair cells</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>n</td>
<td>SA (10^6 nm^2)</td>
</tr>
<tr>
<td>Cell 1 juxtastriolar</td>
<td>14</td>
<td>1.74 ± 0.79</td>
</tr>
<tr>
<td>Cell 2 extrastriolar</td>
<td>29</td>
<td>1.58 ± 0.73</td>
</tr>
<tr>
<td>Cell 3 extrastriolar</td>
<td>13</td>
<td>1.61 ± 0.81</td>
</tr>
<tr>
<td>Average</td>
<td>56</td>
<td>1.64 ± 0.09</td>
</tr>
</tbody>
</table>

Samples, region and cell number from which samples were taken; n, number of mitochondria measured for each type of cell; Values, mean ± SD. Only complete mitochondria from the subcuticular region (~6 μm below the apical cell membrane) in type I and neighboring type II hair cells were reconstructed. As mitochondrial function is related to overall size (in particular to surface area), mean volumes (Vol) and surface areas (SA) were measured for each organelle, and surface to volume ratios (SA/Vol) were calculated. Student t test was used to compute significance levels for type I vs. type II mitochondrial surface areas (P < 0.0001) and volumes (P < 0.0001).

Movie S1. Movie showing digital sections through the tomogram (six serial semithin sections at 0.5 μm per section, a total of 3 μm joined volume) and presentation of modeled apical structures from cell 1 (a juxtastriolar type I hair cell): hair-cell membrane, cuticular plate, subcuticular mitochondria, kinocilium, SRs, SO—thick and thin filaments—and connections between SRs and SO. Extracellular structures: calyx terminal, containing dense core vesicles. All labels have been colored to match the color-coding of each modeled structure.

Movie S2. Movie showing digital sections through the tomogram (11 serial semithin sections at 0.5 μm per section, a total of 5.5 μm of joined volume) and presentation of apical structures modeled from cell 2 (an extrastriolar type I hair cell): hair-cell membrane, cuticular plate, subcuticular mitochondria, kinocilium, SRs, centriole and kinociliar rootlets, SO—thick and thin filaments—and connections between SRs and SO. Extracellular structures: calyx terminal, containing dense core vesicles. Labels have been colored to match the color-coding of each modeled structure.