Identification of the Hair Cell Soma-1 Antigen, HCS-1, as Otoferlin

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ABSTRACT

Hair cells, the mechanosensitive receptor cells of the inner ear, are critical for our senses of hearing and balance. The small number of these receptor cells in the inner ear has impeded the identification and characterization of proteins important for hair cell function. The binding specificity of monoclonal antibodies provides a means for identifying hair cell-specific proteins and isolating them for further study. We have generated a monoclonal antibody, termed hair cell soma-1 (HCS-1), which specifically immunolabels hair cells in at least five vertebrate classes, including sharks and rays, bony fish, amphibians, birds, and mammals. We used HCS-1 to immunoprecipitate the cognate antigen and identified it as otoferlin, a member of the ferlin protein family. Mutations in otoferlin underlie DFNB9, a recessive, nonsyndromic form of prelingual deafness characterized as an auditory neuropathy. Using immunocytochemistry, we find that otoferlin is associated with the entire basolateral membrane of the hair cells and with vesicular structures distributed throughout most of the hair cell cytoplasm. Biochemical assays indicate that otoferlin is tightly associated with membranes, as it is not solubilized by alterations in calcium or salt concentrations. HCS-1 immunolabeling does not colocalize with ribeye, a constituent of synaptic ribbons, suggesting that otoferlin may, in addition to its proposed function in synaptic vesicle release, play additional roles in hair cells.

Keywords: balance, hearing, auditory system, vestibular system, ribeye

INTRODUCTION

Hair cells are the mechanoreceptors of the inner ear that detect sound and accelerations of the head. In fish and aquatic amphibians, hair cells in the lateral line organs detect fluid movements that are important for schooling behaviors, locomotion in turbulence, detection of prey, and the avoidance of predators (Dikgraaf 1933; Ghysen and Dambly-Chaudiere 2007).

Hair cells reside in sensory epithelia with their apically located hair bundles projecting into the lumen of the inner ear where they are poised to detect minute displacements. The hair bundle is a highly organized structure composed of actin-filled stereocilia arranged in rows of increasing height. Deflections of the order of a few nanometers are sufficient to increase the open probability of cation-selective transduction channels
located at stereociliary tips. Transduction channel opening leads to membrane depolarization and culminates in synaptic vesicle release at synapses located in the basolateral portion of the cell (Hudspeth 1989; LeMasurier and Gillespie 2005). Thus, apical and basolateral regions of the hair cell have distinct functions that are likely to depend upon different sets of proteins. Other subcellular regions within the hair cell include the cuticular plate, an actin-rich meshwork that anchors the hair bundle; the pericellular necklace, a vesicle-rich region encircling the cuticular plate; and the kinocilium, an axonemal cilium located at the tall edge of the hair bundle.

In all organisms, hair cells are few in number. Approximately 80,000 hair cells are present in the eight hair cell containing end organs of the chicken inner ear, and approximately 12,000 are present in the six end organs in the mouse inner ear. Such low numbers of hair cells, which require harvesting through time-consuming microdissections of animal ears, make it difficult to biochemically identify and characterize proteins important for hair cell function. The generation of immunological reagents against proteins in the inner ear has proved to be a useful alternative approach to the identification of molecules that are important for auditory and vestibular function (Richardson et al. 1990; Goodyear and Richardson 1992, 1999, 2003; Ahmed et al. 2006; McGee et al. 2006). Thus, apical and basolateral portions of the cell (Hudspeth 1989; LeMasurier and Gillespie 2005). Thus, apical and basolateral regions of the hair cell have distinct functions that are likely to depend upon different sets of proteins. Other subcellular regions within the hair cell include the cuticular plate, an actin-rich meshwork that anchors the hair bundle; the pericellular necklace, a vesicle-rich region encircling the cuticular plate; and the kinocilium, an axonemal cilium located at the tall edge of the hair bundle.

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**MATERIALS AND METHODS**

**Animals**

All animal procedures followed NIH guidelines and were approved by the Animal Care and Use Committee at the West Virginia University School of Medicine, University of Virginia School of Medicine, Washington University School of Medicine, or the United Kingdom Home Office and the local ethical committee of the University of Sussex.

**Antibody generation and purification**

Sensory epithelia were delaminated from the utricles of 1–21 day post-hatch (PH) White Leghorn chicks after treatment with thermolysin (Sigma), 500 μg/ml for 60 min at 37°C. The epithelia were lightly fixed in 1% paraformaldehyde for 15 min, washed by centrifugation, and frozen in liquid nitrogen. Six-week-old female BALB/C mice were injected intraperitoneally with 30 sensory epithelia emulsified in Ribi adjuvant (Ribi Immunocore Research, Hamilton, MT, USA) at day 0 and day 21. At day 60, an additional 50 sensory epithelia were injected. Four days after the final immunization, each mouse was killed, its spleen was removed, and the splenocytes were fused with SP/0 myeloma cells. Hybridomas were then selected and expanded. Hybridoma supernatants from over 400 wells were screened initially by ELISA on wells coated with 1 μg/well of antigen homogenate. Wells that were positive by ELISA were expanded further and secondarily screened by immunohistochemistry on cryostat sections of utricles from chickens. Cells producing supernatants that tested positive by immunohistochemistry were cloned twice by limiting dilution to obtain single clones. The monoclonal antibody used in this study is the HCS-1 antibody, also called mAb76.

**Immunoprecipitation**

Dissected tissues that had been harvested and stored at −80°C were solubilized in cold phosphate-buffered saline (PBS; 137 mM NaCl, 2.7 mM KCl, 4.3 mM Na2HPO4, and 1.5 mM KH2PO4, pH 7.4) containing a cocktail of protease inhibitors (1 mM PMSF, 2 mM benzamidine, 1 μg/ml leupeptin, and 1 μg/ml pepstatin or 0.2 mM PMSF, 10 μg/ml apronin, 0.5 μg/ml leupeptin, and 0.7 μg/ml pepstatin) and 1% Triton X-100 (TX-100), centrifuged at 15,000 gmax for 10 min at 4°C, and aliquots of the supernatant were mixed with protein A Sepharose beads (Sigma Aldrich, St. Louis, MO, USA) charged either with mAb HCS-1 or an irrelevant mAb of the same isotype (anti-HCA mAb H27; Goodyear and Richardson, unpublished) or the anti-V5 antibody (Invitrogen, Carlsbad, CA, USA) overnight. Beads were washed extensively with PBS containing 0.1% TX-100, and bound proteins were eluted with 2× concentrated sample buffer (20% glycerol, 20 mM EDTA, 250 mM Tris-HCl, pH 8.0, 0.02% bromophenol blue, 6% sodium dodecyl sulfate,
10% β-mercaptoethanol) and analyzed by SDS-PAGE. To determine the solubility of otoferlin, tissues were sequentially extracted in 150 mM NaCl, 10 mM sodium phosphate, pH 7.2 (PBS); 5 mM ethylene glycol bis(2-aminoethyl ether)-N,N′,N′,N′-tetraacetic acid in 150 mM NaCl, 10 mM HEPES, pH 7.2 (EGTA); 1 M NaCl, 10 mM HEPES, pH 7.2 (High Salt); 10 mM HEPES, pH 7.2 (Low Salt); and 0.1% TX-100 in 150 mM NaCl, 10 mM HEPES, pH 7.2 (TX-100). Extracts were centrifuged at 70,000 g max for 15 min at 4°C, and the resultant supernatants were immunoprecipitated in the presence of 0.1% TX-100 and analyzed as described above.

Mass spectrometry

Each of the two protein bands precipitated from chick inner ears by mAb HCS-1 was excised from a Coomassie-stained acrylamide gel, and the gel slices were washed in water, reduced, carboxymethylated, and digested with trypsin. The resultant tryptic peptides were desalted, concentrated, and applied to a capillary liquid chromatography system coupled to a quadrupole time-of-flight mass spectrometer (LC-MS/MS; Qtof2, Waters). The MS/MS data obtained were used to search the NCBI databases employing the MASCOT search engine (http://www.matrixscience.com). Probability-based MASCOT scores were used to evaluate identification; matches with a P<0.05 for random occurrence were considered significant.

Immunocytochemistry

Cryosections of chicken, dogfish, gerbil, and guinea pig

Chicken, gerbil, and guinea pig tissues were fixed for 1 h at room temperature in 3.7% formaldehyde in 0.1 M sodium phosphate buffer, pH 7.4. Inner ears from the smooth dogfish, Mustelus canis, were fixed in 3.7% formaldehyde in elasmobranch Ringer’s solution (280 mM NaCl, 540 mM urea, 10 mM KCl, 4.5 mM NaHCO3, 10 mM CaCl2, 50 μM Na2HPO4, 0.1% dextrose; Marine Biological Laboratory Chemical Room, Woods Hole, MA, USA) overnight. Fixed tissues were washed in PBS, cryoprotected with 30% sucrose in PBS overnight, embedded in 1% low gelling point agarose (type VII agar, Sigma Aldrich), frozen, and cryosectioned at ~30°C. Frozen sections were mounted on gelatin-coated slides. Tissues from 3-week-old gerbils and a 3-month-old guinea pig were decalcified in 0.5 M EDTA, pH 8.0, for 24–48 h prior to cryoprotection; tissues from 2-day-old chicks and mature dogfish were processed without decalcification. Sections were preblocked with 10% horse serum in Tris-buffered saline (TBS; 10 mM Tris-HCl, pH 7.4, 150 mM NaCl) and incubated overnight at room temperature with either the HCS-1 antibody alone or with mAb HCS-1 and anti-CdBP2 (1:100, BD Biosciences, San Jose, CA, USA), which detects ribeye, anti-Lamp1 (1:500; Abcam Inc, Cambridge, MA, USA), and anti-protein tyrosine phosphatase receptor Q (anti-Ptprq; 1:100, mAb D10 supernatant) in 10% horse serum in TBS. Following washing in TBS, bound antibodies were detected with species-specific secondary antibodies (1:200 FITC goat anti-mouse IgG2a, Serotech, Oxford, UK and 1:100 rhodamine swine anti-rabbit, Dako). In samples that were only immunolabeled with the HCS-1 antibody, filamentous actin was stained with Texas Red phalloidin (1:300, Invitrogen). Images were obtained with an Axioskop epifluorescent microscope or a LSM 510 confocal microscope (Zeiss, Thornwood, NY, USA) and colorized with Photoshop 7.0 software (Adobe Systems, Inc., San Jose, CA, USA).

Zebrafish cryosections

Larval zebrafish were euthanized at 72 h postfertilization and fixed overnight in 4% paraformaldehyde (prepared in 0.1 M phosphate buffer, pH 7.4). Following thorough rinsing in PBS, fish were cryoprotected by overnight incubation in 30% sucrose in PBS, embedded in OCT compound (Miles Diagnostics, Elkhart, IN, USA), and cryosectioned at 10 μm thickness. Specimens were blocked in 2% normal horse serum, 2% normal goat serum, 1% bovine serum albumin, and 0.2% TX-100 in PBS and were then incubated overnight in HCS-1 (1:500) and anti-PAX2 (1:200; Zymed, South San Francisco, CA, USA), with 2% horse serum and 0.2% TX-100 in PBS. Sections were then rinsed in PBS and incubated for 2 h in Alexa-488 anti-mouse IgG (1:500; Invitrogen) and Cy-3 anti-rabbit IgG (1:500; Amersham, Piscataway, NJ, USA). Following washing, specimens were mounted on microscope slides in 9:1 glycerol/PBS and imaged on a Nikon Eclipse 2000 inverted microscope with epifluorescent illumination.

Bullfrog saccular wholemounts and isolated cells

Bullfrog sacculi were fixed in 3% paraformaldehyde in cold PBS for 25 min, washed in PBS, and permeabilized in 0.1% TX-100 in PBS for 1 h. Following a PBS wash, the tissue was blocked in 5% goat serum in PBS for 1 h and incubated overnight at 4°C in 4.5 μg/ml HCS-1 antibody in blocking solution. Bound antibodies were detected with Alexa-488 goat anti-mouse secondary antibody (Invitrogen), and filamentous actin was labeled with 35 nM Alexa-568 phalloidin (Invitrogen). Samples were mounted in Vectashield (Vector Laboratories, Burlingame, CA, USA) and viewed with a LSM 510 META confocal microscope (Zeiss). Isolated bullfrog saccular hair cells (Lumpkin and Hudspeth 1995) were allowed to settle onto coverslip dishes (Mattek, Ashland, MA,
USA) coated with 1 mg/ml concanavalin A. The cells were fixed for 25 min in 3% paraformaldehyde in cold PBS, washed for 10 min in PBS, and permeabilized for 20 min in 0.1% saponin in PBS. Dishes were washed for 5 min in PBS and blocked for 1 h in 10 mg/ml bovine serum albumin (fraction V; EMD Biosciences, San Diego, CA, USA) and 5% goat serum in PBS. Primary antibody solutions of 4.5 μg/ml HCS-1 antibody with or without 2.5 μg/ml F2b anti-plasma membrane Ca^{2+} ATPase b isofrom (anti-PMCAb) antibody (Dumont et al. 2001; gift from P. Gillespie, Oregon Health & Science University) in blocking solution were incubated with the cells overnight at room temperature. Following two 10-min washes with PBS/0.1% Tween-20, dishes were incubated at room temperature for 2 h with blocking solution containing 33 nM Alexa-568 phallolidin and either 13 μg/ml Alexa 488 goat anti-mouse antibody alone to detect bound mAb HCS-1 or 13 μg/ml goat anti-mouse Alexa 488 and 13 μg/ml Alexa-568 goat anti-rabbit (Invitrogen) to visualize both otoferlin and PMCAb. Dishes were then washed 2×10 min in PBS/0.1% Tween-20 and 1×10 min in PBS and mounted in Vectashield. Images were obtained using an LSM 510 META confocal microscope with a 100×, 1.4 NA objective.

Preparation of expression vectors

Full-length otoferlin was amplified by polymerase chain reaction (PCR) with Pfu polymerase using cDNA from P2 chicken utricle as the template. The 5′ half of the cDNA was amplified with primers GgOt-oR5 (GATCCGCCTAGCCACCATTGCTTCAGCTG CAGCT, NheI site underlined) and GgOt-oR6 (CAAAAGGATCCAAAGGCCCAC, including a T=C point mutation at base 11 to introduce a BamHI site, underlined). PCR product was gel-purified, cut with NheI and BamHI, and ligated into the vector backbone of pEGFP-actin (Clontech) that had been cut with NheI and BamHI, dephosphorylated, and gel-purified to remove the EGFP-actin insert. After transformation into Escherichia coli XL1 blue cells (Stratagene), recombinant plasmids were identified by PCR and by restriction enzyme digestion of plasmid DNA. One plasmid was selected for use in the second cloning step, linearized with BamHI, and dephosphorylated. The 3′ half of the cDNA was amplified as above, using primers GgOt-oF7 (GGATCCCTTTTCCGATCCCTT, including point mutation A>G at base 11 to create a BamHI site, underlined) and GgOt-oR4 (CCCCGATCCCTTTGAGACGCAGGAC, BamHI site underlined). The PCR product was gel-purified, cut with BamHI, and ligated into the prepared plasmid. After transformation into E. coli XL1 Blue, full-length otoferlin clones were identified by PCR and restriction enzyme digests of plasmid DNA. Clone pOtofFL5 was sequenced fully to confirm that the insert encoded full-length otoferlin. pOtofFL5 was transfected into cells using Lipofectamine 2000. After overnight incubation, cell monolayers were fixed in 3.7% formaldehyde in 0.1 M sodium phosphate buffer, pH 7.5, blocked in TBS containing 10% horse serum and 0.1% TX100 for 1 h, and then incubated overnight with HCS1 mAb diluted 1 in 500 in TBS plus 10% horse serum. After washing three times in TBS, Alexa 555 conjugated goat anti-mouse IgG2a (1 in 500 in TBS/HS) was added for 1 h, and monolayers were washed three times in TBS, mounted with Vectashield, and photographed on a Zeiss Axioplan fluorescence microscope or a Zeiss LSM510 confocal microscope.

Three overlapping fragments of otoferlin were amplified by PCR with Pfu polymerase (Stratagene) using cDNA from P2 chicken utricle as the template. Products obtained with primer pairs GgOt-oF1 (CAGATCTCAGTGTGGCTCTGAGCTGCAGCT, XhoI) and GgOt-oR1 (CCGGTGATCCCTAGGG CAGGTAGCCTTTGCTTC, BamHI) and GgOt-oF2 (CAGATCTCAGTGTGGCTCTGAGCTGCAGCT, XhoI) and GgOt-oR2 (CCGGTGAT CCCCCCTGATATTTCAGCTGTG, BamHI), and GgOt-oF3 (CAGATCTCAGTGTGGCTCTGAGCTGCAGCT, XhoI) and GgOt-oR4 (CCGGTGATCCCTAGCCAGGCTTTGCTTC, BamHI) were gel-purified, digested with XhoI and BamHI, and ligated into XhoI and BamHI cut pEGFP-actin (Clontech). PCR and restriction digests of plasmid DNA were used to identify clones encoding EGFP-otoferlin fusion proteins. All inserts were confirmed to be free of errors by DNA sequencing. All three constructs expressed EGFP-tagged protein in mammalian cells that was not recognized by the HCS-1 mAb, possibly as a result of misfolding caused by the tag (data not shown).

RT-PCR

Total RNA from 1 day post-hatch chick tissues was isolated using Trizol (Invitrogen, Paisley, UK) then treated with RNase-free DNase I to remove traces of genomic DNA (Applied Biosystems, Warrington, UK). Randomly primed first-strand cDNA was synthesized from 1 μg of total RNA using AMV reverse transcriptase (Promega, Southampton, UK) and the reaction diluted to 100 μl final volume. Otoferlin and GAPDH PCR products were amplified from 2 μl aliquots of the reverse transcriptase (RT) reactions using Bioline Taq polymerase (Bioline, UK) and primers GgOt-oF3 and GgOt-oR2 (see above) and GAPDHF1 (GCTGTGATTTGTTG GAGTC) and GAPDHR1 (TCAGCAGCAGCCTTCAC TAC). Aliquots (5 μl) of the PCR reactions were run on...
1.5% agarose gels, stained with ethidium bromide, and photographed under UV illumination.

RESULTS

Isolation and characterization of hair cell soma-1 antibody

As a means to identify and characterize proteins important for inner ear function, we immunized mice with emulsified chicken inner ear sensory epithelia that had been briefly exposed to a dilute aldehyde fixative. We generated a panel of 400 wells containing hybridoma cells. Supernatants from these wells were initially screened by ELISA on a chicken inner ear homogenate. Cells from the wells that gave a positive signal by ELISA were subsequently expanded and screened by immunohistochemistry on frozen sections of chicken utricles. Using this approach, 40 antibodies were isolated that recognized proteins in specific cell types of the inner ear. Of these, four antibodies were further characterized: two IgG2b antibodies bound specifically to supporting cells (data not shown), one IgM antibody bound to the cuticular plate of the hair cell (data not shown), and one IgG2a antibody bound to an antigen specifically found in the hair cell soma. This latter antibody was named the HCS-1 antibody.

The specific localization of the HCS-1 antigen to the soma of utricular hair cells (Fig. 1A) led us to further examine its expression in other hair cell containing end organs in a variety of species. In the chicken, the HCS-1 antigen was present in all hair cells of the basilar papilla (Fig. 1B and data not shown) and other hair cell containing end organs (data not shown). In the inner ear of the guinea pig (Fig. 1C, D) and gerbil (not shown), the HCS-1 antigen was present in the inner hair cells of the cochlea, but not in the outer hair cells, and was detectable at low levels in the vestibular hair cells of the maculae. In addition, the HCS-1 antigen was present in the utricular, saccular, and lateral line hair cells in zebrafish (Fig. 1E); in the utricular and saccular hair cells of the dogfish (Fig. 1F and data not shown); and in the saccular hair cells of the bullfrog (Fig. 1G). This mAb does not detect denatured protein on immunoblots (data not shown) suggesting that it recognizes a conformation-dependent epitope.

Hair cell soma-1 antigen is otoferlin

To identify the HCS-1 antigen, we used the antibody to immunoprecipitate proteins from a Triton X-100 soluble extract of the chicken inner ear and subjected the immunoprecipitated protein to mass spectrometry. When resolved by SDS-PAGE, the immunoprecipitated...
protein consistently migrated as a high molecular weight doublet, two bands with apparent molecular masses of 230 and 210 kDa (Fig. 2A). Both bands yielded a number of tryptic peptides with masses corresponding to peptides derived from the sequences for mouse and human otoferlin, a member of the ferlin family of proteins which is known to be critical for normal auditory function (Yasunaga et al. 1999, 2000; Migliosi et al. 2002; Mirghomizadeh et al. 2002; Tekin et al. 2005; Ebermann et al. 2007; Longo-Guess et al. 2007). Seven peptides unique to the otoferlin sequence were obtained from the 230-kDa band, one of which was present within a larger peptide fragment (Fig. 2B). The 210-kDa band yielded six of the same peptides along with an additional six peptides (underlined with asterisks).

Location of tryptic peptides in the sequence of mouse cochlear otoferlin. Tryptic digestion followed by mass spectrometry identified seven unique peptides to the otoferlin sequence for the 230-kDa immunoprecipitated protein (underlined). One of these peptides (italics and double underlined) overlapped a larger peptide. The 210-kDa immunoprecipitated protein identified six of the same peptides obtained for the higher molecular weight band and an additional six peptides (underlined with asterisks).

To confirm that the HCS-1 antibody recognizes otoferlin, we transiently transfected cultured cell lines with plasmid DNAs encoding the complete open reading frame of chick otoferlin, espin-EGFP, or a combination of both plasmids and then stained the cultures with the HCS-1 antibody. HCS-1 immuno-reactive cells were observed in cultures that had been transfected with otoferlin alone (Fig. 3A) but not in cells that had been sham-transfected (Fig. 3B) or transfected with espin-EGFP alone (not shown). In polarized epithelial cell lines that had been co-transfected with both plasmids, otoferlin was detected within the cell soma (Fig. 3C) but not within the highly elongated microvilli that form on the apical surface of the cell in response to espin over-expression (Fig. 3D). Otoferlin is therefore excluded from the microvilli of polarized epithelial cells lines in vitro, as it is from the stereocilia of hair cells in vivo. We also transiently over-expressed three C-terminally EGFP-tagged otoferlin fragments that together encompassed the entire open reading frame of otoferlin in a number of different cell lines, including HEK 293, MDCK, and Caco2 cells, and in supporting cells in primary cultures derived from the chicken utricle. Immunoreactivity for HCS-1 was not detected in any cell type expressing any of these otoferlin constructs (not shown) suggesting that the

FIG. 2. Identification of HCS-1 antigen as otoferlin. A Immunoprecipitation from chick sensory epithelia using the HCS-1 antibody. Coomassie-stained SDS-PAGE gel revealing a high molecular weight doublet immunoprecipitated from a Triton X-100 extract of chicken inner ear (arrow, left lane). These proteins were not immunoprecipitated with the H27 antibody, an isotype-matched irrelevant antibody (right lane). Immunoprecipitated bands were excised and subjected to mass spectrometry. Bands corresponding to the immunoglobulin heavy- and light chains are indicated by HC and LC, respectively. Molecular mass markers in kilodaltons are indicated to the left. B Location of tryptic peptides in the sequence of mouse cochlear otoferlin. Tryptic digestion followed by mass spectrometry identified seven unique peptides to the otoferlin sequence for the 230-kDa immunoprecipitated protein (underlined). One of these peptides (italics and double underlined) overlapped a larger peptide. The 210-kDa immunoprecipitated protein identified six of the same peptides obtained for the higher molecular weight band and an additional six peptides (underlined with asterisks).
fragments do not fold correctly and providing further evidence that HCS-1 recognizes an epitope that depends on the tertiary structure of the protein.

Otoferlin expression and localization

Recent reports have demonstrated a role for otoferlin protein in auditory hair cell synaptic vesicle release (Roux et al. 2006; Heidrych et al. 2009). Using the HCS-1 antibody, we first sought to determine the tissue distribution of the otoferlin protein in the chick. Otoferlin immunoreactivity was not detected in cryosections of the chicken forebrain, cerebellum, gizzard, heart, intestine, kidney, liver, lung, muscle, retina, skin, or tongue (Fig. 4A–L) but was in utricle (Fig. 4M). Using RT-PCR with primers that span two introns and should only amplify a 381-bp fragment from cDNA, expression was detected in at least four of these tissues (forebrain, cerebellum, kidney, and retina; Fig. 4N). HCS-1 may either recognize an ear-specific epitope or an epitope that is masked in all tissues other than the ear. Alternatively, it may only detect cells in which otoferlin is expressed at very high levels.

During inner ear development, otoferlin immunoreactivity was detected as early as E7 in the distal end of the chick basilar papilla, in a region where Ptpq (Goodyear et al. 1997) expression is detected by the HCA antibody (Fig. 5). Otoferlin staining clearly revealed that immature hair cells possess a thin cytoplasmic process that extends down to the basal surface of the epithelium (Fig. 5, E7–E8). As the hair cells mature, these processes retract (Fig. 5, E10). Double labeling studies with HCS-1 mAb and antibodies to the hair cell marker myosin VI at E7 and E8 (Fig. 6) indicate that HCS-1 immunoreactivity is confined to cells and cellular processes in which myosin VI can be detected. HCS-1 is therefore unlikely to be expressed by either supporting cells or hair cell progenitors.

In mature chick hair cells, double labeling for otoferlin and Lamp1, a marker for lysosomes and late endosomes (Griffiths et al. 1988), did not reveal a significant degree of co-localization (Fig. 7A). Co-labeling for otoferlin and ribeye, a known synaptic ribbon protein (Schmitz et al. 2000), failed to reveal any apparent co-localization of these two proteins in chick hair cells (Fig. 7B). Also, there was no evidence for otoferlin expression at the ribeye-positive synaptic ribbons that were immunolabeled within retinal photoreceptors and bipolar cells (Fig. 7C). In cryosections of chick basilar papilla, double immunolabeling for otoferlin and PMCAb (Fig. 7D), an isoform of the plasma membrane calcium ATPase (Dumont et al. 2001), indicates that otoferlin is associated with the basolateral membrane of the hair cell.

To better appreciate the expression of otoferlin at the subcellular level, we examined the bullfrog sacculus, the hair cells of which are relatively large and can readily be isolated from the sensory epithelium. We first confirmed antibody specificity in this tissue by immunoprecipitation. As seen for the chick inner ear, the HCS-1 antibody immunoprecipitated a doublet in the 225-kDa region from bullfrog saccules (Fig. 8A). Additional co-immunoprecipitating proteins were not evident on silver-stained SDS-PAGE gels, although some proteins may have been masked by the presence of the immunoglobulin heavy and light chains (Fig. 8A). In whole-mount preparations of the sacculus, otoferlin immunoreactivity is restricted to hair cells where it is abundant in the soma and, as in the chick, appears to be localized to the basolateral membrane (Fig. 1G and see Gale et al. 2000). As in the whole-mount preparations, otoferlin was present along the basolateral membrane of the isolated hair cells and in membranous structures within the cytoplasm of the isolated cells. This immunoreactivity substantially overlapped with that of the PMCAb (Fig. 8B), an abundant hair cell integral membrane protein (Dumont et al. 2001). In the cytoplasm of bullfrog hair cells, otoferlin was most often seen

FIG. 3. HCS-1 antibody staining of transfected cell lines. A HEK-293 cells transfected with plasmid construct encoding the full-length form of chick otoferlin stained with the HCS-1 antibody. B Sham-transfected HEK-293 cells stained with the HCS-1 antibody. C, D Caco-2 cell co-transfected with plasmid construct encoding a full-length form of chick otoferlin and a plasmid encoding EGFP-espin doubled-labeled with HCS-1 antibody (red in C, D) and antibody to EGFP (green in D). Arrows indicate the highly elongated microvilli that form on the EGFP-espin transfected cells. Scale bars=10 μm.
enriched in a region immediately between the cuticular plate and the nucleus (Fig. 8B). The patterns of otoferlin immunoreactivity in the isolated bullfrog hair cells and in the intact saccular epithelium did not show discrete “hot spots” in the basolateral portion of the cell as might be expected if the protein was specifically localized to the ~20 synaptic dense bodies present in that location (Isa and Hudspeth 1996; Lenzi et al. 1999). In a subset of isolated bullfrog saccular hair cells, an enrichment of otoferlin in the perinuclear region of the cell was observed. The localization in this region appeared to be related to the apparent health of the cells and was present in cells that contained large perinuclear vacuoles, which also were PMCAb immunopositive, and may be an indicator of the damage caused by the isolation procedure (data not shown). In approximately 30% of the isolated bullfrog saccular hair cells, otoferlin immunoreactivity was also present in the apical membrane (data not shown). We did not, however, observe apical membrane labeling in the whole-mount preparations of the bullfrog sacculus.

Solubility of hair cell otoferlin

To assess whether otoferlin is, as predicted, a transmembrane protein, we sequentially subjected chicken inner ear tissue with buffered saline containing a calcium chelator, high salt, low salt, and 0.1% Triton X-100 and then subjected the soluble protein fraction following each treatment to immunoprecipitation. Of these treatments, only extraction with 0.1% Triton X-100 resulted in the release of otoferlin into the soluble protein fraction (Fig. 9) suggesting that both isoforms are transmembrane proteins.

DISCUSSION

The identification and characterization of hair cell-specific proteins has been hindered by the relatively small numbers of hair cells that can be harvested from the inner ears of most vertebrates. We have used the monoclonal antibody, HCS-1, to immunoprecipitate its cognate antigen from chicken inner ears. Using mass spectroscopy, we have identified the antigen as otoferlin. We describe the expression of otoferlin across a number of species, its tissue distribution, subcellular localization, and developmental profile.

On the basis of its primary sequence, otoferlin is predicted to be a large single-pass transmembrane protein containing three to six protein kinase C conserved region 2 (C2) domains (Yasunaga et al. 1999). C2 domains are membrane-targeting regions that mediate Ca$^{2+}$-dependent membrane interactions (Cho and Stahelin 2006; Lemmon 2008). Highly expressed in the inner ear, otoferlin mRNA is also present in other tissues including the brain, liver, lung and heart, kidney, and testis (Yasunaga et al. 1999, 2000; Schug et al. 2006). Transcripts encoding two classes of otoferlin isoforms, resulting from alternative splicing, have been detected by RT-PCR in the human brain and cochlea. The predicted long isoforms (~230 kDa) contain all six C2 domains while the short isoforms (~140 kDa) contain only the last three C2 domains (Schmitz et al. 2000; Yasunaga et al. 2000). RT-PCR data suggest that only the larger otoferlin isoforms are present in the mouse brain and cochlea (Yasunaga et al. 2000). The HCS-1 antibody immunoprecipitates bands from chicken and frog inner ears that are consistent with the presence of transcripts encoding the larger otoferlin isoforms. Shorter isoforms could also be present in these ears, but may not be precipitated by this monoclonal antibody. Interestingly, HCS-1 immunoprecipitation of the chicken and frog inner ear lysates reveals two large otoferlin protein isoforms that migrate at ~210 and 230 kDa. Cochlear splice variants have not been described that would account for these two bands (Yasunaga et al. 2000), suggesting they may differ in apparent mass as a consequence of posttranslational modification. These immunoprecipitated proteins are not, however, recognized by antibodies against phosphorylated serine, threonine, or tyrosine residues (data not shown), suggesting that the observed heterogeneity is not due to phosphorylation. Both protein isoforms appear to be tightly associated with cell membranes and are not extracted by calcium chelation or either high- or low-ionic strength buffers suggesting that both isoforms contain the hydrophobic C-terminal domain and are indeed transmembrane proteins.

The HCS-1 antibody detects otoferlin located in hair cells of chick, zebrafish, dogfish, and frog inner ears, as well as hair cells in the zebrafish lateral line. These findings indicate that the epitope recognized by the HCS-1 antibody is conserved across many species and at least four vertebrate classes. In chick, the antibody is highly specific for hair cells and the antigen is not detected in organs other than the ear, indicating there may be hair cell-specific isoforms of otoferlin, that the expression levels of otoferlin in other cell types is much lower than in hair cells, or that the epitope is masked in those tissues in which expression can be detected by RT-PCR (i.e., brain, retina, and kidney). Consistent with previous reports of otoferlin mRNA and protein expression in adult mouse and rat cochlea (Judice et al. 2002; Roux et al. 2006; Schug et al. 2006; Brandt et al. 2007), otoferlin immunoreactivity is only detected in the inner hair cells of the adult guinea pig and gerbil cochlea. In the adult guinea pig saccule, HCS-1 immunoreactivity is weak. We do, however, detect robust otoferlin immu-
noreactivity in vestibular hair cells of chick, zebrafish, frog, and dogfish, and others have shown that otoferlin protein is present in the vestibular hair cells of the rat and mouse (Yasunaga et al. 1999; Roux et al. 2006; Schug et al. 2006). We attribute the reduced immunoreactivity in the guinea pig sacculus to the properties of the HCS-1 antibody, which appears to have limited ability to recognize otoferlin in some

![Diagram of HCS-1 antibody staining and otoferlin RT-PCR of multiple chick tissues.](image)

**Fig. 4.** HCS-1 antibody staining and otoferlin RT-PCR of multiple chick tissues. A–N Double labeling with the HCS-1 antibody (otoferlin) and phalloidin (F-actin) of cryosections of forebrain (A), cerebellum (B), gizzard (C), heart (D), intestine (E), kidney (F), liver (G), lung (H), muscle (I), retina (J), skin (K), tongue (L), and utricle (M) from the 1-day PH chicken. Scale bar in M is 40 μm and applies to all panels. N RT-PCR products for otoferlin (Otof) and glyceraldehyde 3-phosphate dehydrogenase (GAPDH) were amplified by RT-PCR from total RNA prepared from 1 day PH chick forebrain (F), cerebellum (C), gizzard (G), heart (H), intestine (I), kidney (K), lung (L), skeletal muscle (M), retina (R), skin (S), tongue (T), and utricle (U). 1, 2 Reverse transcription controls with no RNA added; 3 PCR reaction control. Reverse transcription reactions were performed with (+) or without (−) AMV reverse transcriptase. Otoferlin mRNA expression was detected in forebrain, cerebellum, kidney, retina, and utricle.
The HCS-1 antibody detects hair cells in the basilar papilla of the developing chick as early as E7. Hair cells have been detected in this organ with the HCA mAb and with a Tuj1 antibody at E6.5 (Bartolami et al. 1991; Molea et al. 1999). Otoferlin is therefore expressed within 12 h of some of the earliest markers for hair cell differentiation, and its expression patterns at these early stages clearly reveal that differentiating hair cells retain a cytoplasmic tail that extends right down to the basal surface of the epithelium (Fig. 6), as has been described previously for regenerating hair cells in the chick and the frog (Girod et al. 1989; Steyger et al. 1997; Stone and Rubel 2000).

In mature hair cells, otoferlin is predominately cytoplasmic and in bullfrog saccular hair cells it is enriched in the supranuclear cytoplasm, the region between the nucleus and the cuticular plate. It also is present in the basolateral plasma membrane where it co-localizes with the membrane protein PMCA2b. After isolation of bullfrog saccular hair cells, otoferlin immunoreactivity was also detected at the apical membrane in a subset of the hair cells. We did not, however, observe apical membrane labeling in the hair cells present within the whole-mounted tissue suggesting that disruption of the cellular junctions, which delineate the apical and basolateral membrane domains, can alter the subcellular localization of this protein and that this protein may undergo specific targeting and relatively dynamic regulation.

Otoferlin is the member of the ferlin family of proteins and shares the highest protein sequence identity with dysferlin, a protein proposed to play an important role in plasma membrane repair (Bansal and Campbell 2004). Such repair occurs through the fusion of dysferlin-containing lysosomal vesicles to the plasma membrane. The similarity between dysferlin and otoferlin suggests that otoferlin may play a role in membrane-fusion events. While otoferlin is present near the basolateral membrane of hair cells, the lack of significant co-localization with Lamp1, a lysosomal and late endosomal marker (Griffiths et al. 1988), implies that it may not be involved in general membrane repair.
FIG. 7. Immunolocalization of otoferlin, Lamp1, ribeye, and PMCAb in chicken tissues. Confocal microscopy images of cryosections from chicken basilar papilla (A, B, and D) and retina (C) double-labeled in green with antibodies to otoferlin (HCS-1 antibody) and, in red, antibodies to either A Lamp1; a marker for lysosomes and late endosomes; B, C ribeye, the synaptic ribbon marker labeled with the anti-CtBP2 antibody, or D PMCAb, a component of the hair cell’s basolateral membrane. Scale bars in A–D=10 μm; scale bars in insets=2 μm.

FIG. 8. Immunoprecipitation of otoferlin from bullfrog saccule and subcellular localization in isolated bullfrog saccular hair cells. A Silver stain of a SDS-PAGE gel containing proteins immunoprecipitated from bullfrog sacculus with the anti-otoferlin antibody (HCS-1) or an irrelevant isotype-matched control antibody (anti-V5 antibody). Immunoglobulin heavy chain is observed at approximately 50 kDa. Molecular mass markers in kilodaltons are indicated to the right. B Confocal localization of actin (red), otoferlin (HCS-1 antibody, green), and PMCAb (blue) in an isolated saccular hair cell by confocal microscopy. Individual channels are shown to the left, merged image is shown to the right. Scale bar=5 μm.
Mutations in the human otoferlin gene result in nonsyndromic prelingual deafness, DFNB9, a recessive auditory neuropathy (Yasunaga et al. 1999; Rodriguez-Ballesteros et al. 2003; Varga et al. 2003, 2006; Tekin et al. 2005; Choi et al. 2009) and mice that carry otoferlin mutations also exhibit auditory deficits (Roux et al. 2006; Longo-Guess et al. 2007). Inner hair cells from otoferlin knockout animals exhibit impaired calcium-evoked exocytosis (Roux et al. 2006), as do immature outer hair cells (Beurg et al. 2008) leading to the hypothesis that otoferlin is critical for synaptic vesicle release in cochlear hair cells and that it may serve a role similar to the function of synaptotagmin in nerve terminals (Roux et al. 2006). Otoferlin has been shown to associate with syntaxin 1A, SNAP-25 (Roux et al. 2006; Ramakrishnan et al. 2009) and L-type Ca\(^{2+}\) channels (Ramakrishnan et al. 2009); however, it also interacts with myosin VI (Heidrych et al. 2009). Moreover, inner hair cells from hypothyroid rats exhibit exocytotic profiles similar to those in immature inner hair cells despite lacking otoferlin mRNA and protein (Brandt et al. 2007), and although otoferlin is expressed at high levels in vestibular hair cells (Yasunaga et al. 1999; Roux et al. 2006; Schug et al. 2006), vestibular function is unaffected in otoferlin mutants (Longo-Guess et al. 2007; Schwander et al. 2007). Otoferlin is not, therefore, essential for synaptic vesicle release in all types of hair cell. Otoferlin does not co-localize with ribeye, a synaptic ribbon protein, in the chick basilar papilla nor is it enriched in discrete regions that might correlate to the ~20 presynaptic dense bodies found in bullfrog saccular hair cells (Issa and Hudspeth 1996; Lenzi et al. 1999). In addition, otoferlin is highly expressed in both the superior and inferior regions of the chick basilar papilla, both during development and in the mature organ (data not shown), despite hair cells in the inferior region having little, if any, afferent innervation. Collectively, these data suggest that otoferlin may play additional roles in hair cells that are yet to be defined.

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