

Plasma Membrane Ca^{2+} -ATPase Isoform 2a Is the PMCA of Hair Bundles

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Mechanoelectrical transduction channels of hair cells allow for the entry of appreciable amounts of Ca^{2+} , which regulates adaptation and triggers the mechanical activity of hair bundles. Most Ca^{2+} that enters transduction channels is extruded by the plasma membrane Ca^{2+} -ATPase (PMCA), a Ca^{2+} pump that is highly concentrated in hair bundles and may be essential for normal hair cell function. Because PMCA isozymes and splice forms are regulated differentially and have distinct biochemical properties, we determined the identity of hair bundle PMCA in frog and rat hair cells. By screening a bullfrog saccular cDNA library, we identified abundant PMCA1b and PMCA2a clones as well as rare PMCA2b and PMCA2c clones. Using immunocytochemistry and immunoprecipitation experiments, we showed in bullfrog sacculus that PMCA1b is the major isozyme of hair cell and supporting cell basolateral membranes and that

PMCA2a is the only PMCA present in hair bundles. This complete segregation of PMCA1 and PMCA2 isozymes holds for rat auditory and vestibular hair cells; PMCA2a is the only PMCA isoform in hair bundles of outer hair cells and vestibular hair cells and is the predominant PMCA of hair bundles of inner hair cells. Our data suggest that hair cells control plasma membrane Ca^{2+} -pumping activity by targeting specific PMCA isozymes to distinct subcellular locations. Because PMCA2a is the only Ca^{2+} pump present at appreciable levels in hair bundles, the biochemical properties of this pump must account fully for the physiological features of transmembrane Ca^{2+} pumping in bundles.

Key words: hair cells; stereocilia; cochlea; calcium; calcium pump; isozymes

Ca^{2+} is a key modulator of mechanoelectrical transduction by hair cells, the sensory cells of the inner ear. Transduction is elicited by deflection of the hair bundle, the sensory organelle of the hair cell (for review, see Hudspeth et al., 2000). Bundle deflection leads directly to the opening of cation-selective transduction channels, which permit significant entry of Ca^{2+} (Lumpkin et al., 1997; Ricci and Fettiplace, 1998). Entering Ca^{2+} regulates both fast (Crawford et al., 1991) and slow (Eatock et al., 1987) adaptation in hair cells and also triggers bundle movements opposing the stimulus (Benser et al., 1996; Ricci et al., 2000). In addition, Ca^{2+} may regulate the formation of the transduction apparatus (Zhao et al., 1996).

Each hair bundle is composed of dozens to hundreds of actin-filled stereocilia and, with the exception of the mammalian cochlea, a single axonemal cilium called the kinocilium. Because the bundle is exposed to an unusual extracellular fluid, endolymph, which is high in K^+ and low in Na^+ , Ca^{2+} cannot be removed from bundles with $\text{Na}^+/\text{Ca}^{2+}$ exchangers. Furthermore, stereocilia lack intracellular stores to sequester Ca^{2+} . Consequently, hair bundles rely on mobile Ca^{2+} buffers (Ricci et al., 1998) and

the plasma membrane Ca^{2+} -ATPase (PMCA) to regulate Ca^{2+} levels in stereocilia (Lumpkin and Hudspeth, 1998; Ricci et al., 1998; Yamoah et al., 1998); indeed, the majority of the Ca^{2+} that enters through transduction channels is removed by PMCA (Lumpkin and Hudspeth, 1998).

Four PMCA genes (*PMCA1–4*) have been identified; further isoform diversity is generated by alternative splicing in two regions, A and C (Keeton et al., 1993). Although the functional consequences of alternative splicing within region A are poorly understood (Hilfiker et al., 1994), splicing in region C alters the affinity of PMCA for calmodulin (Enyedi et al., 1994) and removes or adds phosphorylation sites (Enyedi et al., 1997).

All PMCA isoforms and many of their splice variants are expressed in the mammalian cochlea (Crouch and Schulte, 1996; Furuta et al., 1998). Furthermore, hair cells contain high levels of PMCA (Crouch and Schulte, 1995; Apicella et al., 1997; Street et al., 1998; Yamoah et al., 1998); levels are particularly high in the hair bundle (Crouch and Schulte, 1995; Yamoah et al., 1998). Although the subcellular distribution of each isozyme and functional significance for expression of multiple PMCA genes remain unknown, at least one of these isozymes, PMCA2, is required for auditory and vestibular function (Kozel et al., 1998; Street et al., 1998; Takahashi and Kitamura, 1999). This requirement for PMCA2 in the inner ear, coupled with the high concentration of PMCA in bundles and essential functions of Ca^{2+} in transduction, led us to surmise that Ca^{2+} regulation in hair bundles was essential for auditory and vestibular function. To better understand the bundle Ca^{2+} regulation and the consequences of

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Table 1. Isozyme-selective antibodies

Name	Specificity	Antigen
F1N	Bullfrog PMCA1	PMCA1 amino acids 1–96
F2N	Bullfrog PMCA2	PMCA2 amino acids 1–99
F2a	Bullfrog PMCA2a, rat PMCA2a	CSSPTSASAAAAGQG (C terminus)
Fb	Bullfrog PMCA1b/2b, rat PMCA1b/2b/3b	CSSSPGSPHLSLETSL (C terminus)
F2v	Bullfrog PMCA2v	CSLPVGGSTHPPSHPPAATDG (v variant)
NR1	Rat PMCA1	ANNSVAYSGVKNSIKEANC (N terminus)
NR2	Rat PMCA2	TNSDFYSKNQRNESSC (N terminus)
NR3	Rat PMCA3	ANSSIEFHPKQQQREVPC (N terminus)
NR4 (JA9)	Rat PMCA4	PMCA4 (mouse monoclonal)

PMCA activity in stereocilia, we identified PMCA isoforms and splice variants in auditory and vestibular epithelia and determined that the hair bundle isoform was PMCA2a.

MATERIALS AND METHODS

Isolation of bullfrog PMCA clones. PMCA1-specific and PMCA2-specific DNA fragments were synthesized from bullfrog liver and lung RNA by RT-PCR. Total RNA was subjected to reverse transcription (Superscript II RNase H-reverse transcriptase, Life Technologies, Grand Island, NY) and was followed by PCR, using degenerate primers (5' to 3': Df1, GAYGCNTGYGARACNATG; Dr1, GGCGCAAGCTTRTCRTANACRTTNCNCCCACAT; Dr2, RAANGTRTANACYTTRTA). The amplification protocol comprised 5 min at 94°C, 30 cycles of 20 sec at 94°C, 2 min at 50°C (–0.5°C for each consecutive cycle), 2 min at 72°C, followed by 14 cycles of 20 sec at 94°C, 2 min at 50°C, 2 min at 72°C, and a final 10 min at 72°C. PCR products were cloned (TA Cloning Kit, Invitrogen, Carlsbad, CA) and sequenced. To generate probes for library screening, we amplified PMCA1 or PMCA2 fragments by PCR and labeled them with [³²P]dCTP, using random hexamers (Amersham Pharmacia Biotech, Piscataway, NJ). The labeled PCR products were used to screen a bullfrog (*Rana catesbeiana*) sacculus cDNA library in Lambda ZAP Express (Stratagene, La Jolla, CA) constructed by Stefan Heller (Rockefeller University, New York, NY); ~600,000 plaques were screened with each probe. To obtain insert-containing pBK-CMV phagemids, we performed *in vivo* excisions on isolated clones by following the manufacturer's protocol (Stratagene).

Analysis of PMCA1 and PMCA2 splicing. Purified total RNA (RNeasy Kit, Qiagen, Valencia, CA) was primed with random hexamers and was reverse transcribed (ThermoScript RT-PCR System, Life Technologies). Nested primers that flank splice regions A and C in bullfrog PMCA1 (F1Cf1, GCAGAGAGGGAGTTACGCCG; F1Cf2, GGTCAGACTTATGGTTTAG; F1Cr1, CATTGGAAGTCCAAGGAGC; F1Cr2, GCACTTCTGCCTGCTGC) and PMCA2 (F2Af1, CTCATGTGATGGAAGGCTC; F2Af2, GAAGAATGCTTGTAACAGC; F2Ar1, TGCCTA TCTGCAGTCCGAG; F2Ar2, GGTCAGCTTGCCCTGTAGG; F2Cf1, CGCCTGACACAGAAGGAGG; F2Cf2, GGGAGTAAAGAAGAGGGC; F2Cr1, CTTCTCTGCCGGGCATCC; F2Cr2, GCAGTGCCATCTCCAGC) or rat PMCA2 region A (R2A+1, GGACGGATGGTGGTACTG; R2A+2, GCTGTGGGTGTCAACTCTC; R2A–1, AGATGGCTGTGGCGTTACC; R2A–2, ACCACCAGCACCGTCACAC) were used in PCR analysis for variability within these regions. The amplification protocol comprised 4 min at 94°C, 30 cycles of 30 sec at 94°C, 30 sec at 50–55°C, 45 sec at 72°C, and a final 10 min at 72°C.

Isozyme-selective antibodies. Antibody names, specificities, and antigens used for their generation are listed in Table 1. Synthetic PMCA peptides, designed with an added N- or C-terminal cysteine residue, were used for the production of antisera (GeneMed Synthesis, South San Francisco, CA). To purify antipeptide antibodies, we coupled peptides to SulfoLink resin (Pierce, Rockville, IL) via their terminal cysteine residues at a density of 1 mg of peptide/ml of resin. Antipeptide antisera, diluted with 10 vol of 25 mM Tris, pH 8, were passed over a 0.5–1 ml peptide column three to four times. After washing the column with 20 vol of 25 mM Tris, pH 8, and 10 vol of 500 mM NaCl/25 mM Tris, pH 8, we eluted the antibodies with 100 mM glycine, pH 2.5, followed by 100 mM CAPS (3-(cyclohexylamino)-1-propane sulfonic acid), pH 11. Antibodies eluted

with acid or base were neutralized, pooled, and dialyzed against PBS containing 0.02% sodium azide. The production of polyclonal antibodies specific for rat PMCA isoforms was described previously (Filoteo et al., 1997); they were subjected to affinity purification with methods similar to those described above. The monoclonal antibody NR4 (JA9) was raised against human erythrocyte PMCA and was directed toward an epitope very near the N terminus of isoform 4 (Adamo et al., 1992; Caride et al., 1996).

Glutathione S-transferase (GST) or *Escherichia coli* maltose-binding protein (MBP) was fused to N-terminal amino acids of PMCA1 (amino acids 1–96) and PMCA2 (amino acids 1–99) in the vectors pGEX-6P-1 (Amersham Pharmacia Biotech) and pMAL-p2 (New England Biolabs, Beverly, MA), respectively. Fusion proteins were expressed in *E. coli* and purified by glutathione or amylose affinity chromatography. Purified GST fusion proteins were injected into rabbits for the production of isoform-selective antibodies (Covance Research Products, Denver, PA). Antibodies were affinity purified on MBP-PMCA or GST-PMCA fusion protein columns, which were constructed by coupling fusion proteins to CNBr-Sepharose (Amersham Pharmacia Biotech) at a density of 1–5 mg of protein/ml of agarose. Antibodies against the N terminus of PMCA1 and PMCA2 were purified as described for antipeptide antibodies, with an additional negative selection step. To remove antibodies that recognize either PMCA1 and GST, first we passed GST-PMCA2 antiserum over a GST-PMCA1-Sepharose column. The flow through from this step subsequently was passed over an MBP-PMCA2-Sepharose column; bound antibodies were eluted as described above. Similarly, GST-PMCA1 antiserum first was passed over a GST-PMCA2-Sepharose precolumn and then purified on an MBP-PMCA1-Sepharose column.

Immunocytochemistry. Auditory and vestibular organs from postnatal day 21 (P21) or older rats were dissected in MEM (Life Technologies) with an added 25 mM HEPES, pH 7.5. Bullfrog sacculi were dissected in bullfrog saline solution [(in mM) 110 NaCl, 2 KCl, 2 MgCl₂, 3 D-glucose, 10 HEPES, pH 7.25] containing 4 mM CaCl₂. Hair cells were isolated as described previously (Yamoah et al., 1998). Tissues or dissociated hair cells were fixed in 3% formaldehyde in saline for 30 min, washed in PBS, and then permeabilized and blocked for 1 hr in blocking solution (PBS, 0.2% saponin, 3% donkey serum, and 1% bovine serum albumin). Formaldehyde was obtained as a 16% solution in sealed ampoules from Electron Microscopy Sciences (Fort Washington, PA). Tissues were incubated overnight at room temperature with 2.5 μg/ml primary antibody in the blocking solution, washed in PBS, and then incubated with a secondary antibody (5 μg/ml Cy5-conjugated donkey anti-rabbit IgG or donkey anti-mouse IgG; Jackson ImmunoResearch Laboratories, West Grove, PA) and 0.25 μM FITC-phalloidin (Sigma, St. Louis, MO) in the blocking solution for 1–3 hr. In some experiments utriculi were colabeled with an antibody against the calcium-binding protein calretinin (1:200; Chemicon, Temecula, CA). Cells and tissues were washed in PBS, mounted with Vectashield (Vector, Burlingame, CA), and viewed with Plan Apochromat 40' (numerical aperture, 1.00) and 60' (numerical aperture, 1.40) oil lenses on a Nikon TE 300 inverted microscope with a Bio-Rad MRC 1024 confocal imaging system (Hercules, CA). Acquired images were processed with Scion Image (version 1.62a) and Photoshop (version 5.03; Adobe Systems, San Jose, CA).

SDS-PAGE and immunoblotting. Proteins were separated by SDS-PAGE that used 10% acrylamide gels with a 150:1 acrylamide-to-bisacrylamide ratio and were transferred to polyvinylidene fluoride blot-

ting membranes in 7.5% methanol and 7.5 mM CAPS, pH 11, at 100 V for 2 hr with cooling. To enhance protein mobilization from the acrylamide gel, we added hemoglobin to the samples and to the pretransfer equilibration solution, as described previously (Gillespie and Gillespie, 1997). PMCA was detected with the 5F10 antibody (Affinity BioReagents, Golden, CO) as described previously (Yamoah et al., 1998), with a 5F10 dilution of 1:5000, alkaline phosphatase-conjugated goat anti-mouse IgG (Jackson ImmunoResearch Laboratories) at 1:20,000, 0.1 mM of the chemiluminescence reagent CSPD [disodium 2-chloro-5-(4-methoxyphosphoryl)-1,2-dioxetane-3,2'-(5'-chloro)-tricyclo[3.3.1.1^{3,7}] decan-4-yl)-1-phenyl phosphate; Tropix, Bedford, MA], and 10% Sapphire (Tropix), an enhancing reagent.

Immunoprecipitation. Hair bundles were purified from bullfrog sacculi by using the twist-off technique (Gillespie and Hudspeth, 1991). For each experiment we estimated the number of saccular equivalents of bundles that had been isolated so that one saccular equivalent is 100% recovery. Detergent-soluble proteins were extracted (Yamoah et al., 1998) from bullfrog residual maculae or purified hair bundles with 250 μ l of immunoprecipitation solution [1% Triton X-100 and (in mM) 1 DTT, 150 NaCl, 25 HEPES, pH 7.5, plus 10 μ M leupeptin, 10 μ M pepstatin, 200 μ M PMSF, and 0.1 mg/ml hemoglobin]. Extracts were incubated on ice for 4 hr with 6 μ g of affinity-purified anti-PMCA antibody. Immune complexes were precipitated by adding 15 μ l of donkey anti-rabbit IgG conjugated to Sepharose, incubating for 2 hr at 4°C with gentle agitation, and centrifuging to sediment the antibody–Sepharose complex. Donkey anti-rabbit IgG–Sepharose was prepared by coupling 2 mg of purified donkey anti-rabbit IgG (Jackson ImmunoResearch Laboratories) to 1 ml of CNBr–Sepharose. After three 250 μ l washes of immunoprecipitation solution, the immune complexes were recovered by incubating at 65°C for 20 min in an SDS-PAGE sample buffer that included 2% SDS and 100 mM DTT. For sequential immunoprecipitations additional anti-PMCA antibodies (6 μ g) were added to the supernatant from the first immunoprecipitation, and the immune complexes were isolated as described above. In some cases a third immunoprecipitation was performed. Proteins from the final supernatant were precipitated with 6 vol of acetone at –20°C and then solubilized in SDS-PAGE sample buffer. Proteins were electrophoresed and transferred to blotting membranes; PMCA was detected with 5F10 as described above.

Immunoelectron microscopy. Postembedding immunoelectron microscopy was performed as described previously (Petralia et al., 1997). Rat cochleae were positioned gently onto a 400-mm-thick slice of Bacto-agar (Difco Laboratories, Detroit, MI) and mounted on filter paper glued to an aluminum specimen holder. Then the specimens were quick frozen by contact with a liquid nitrogen-cooled sapphire block of a Life Cell CF0100 quick-freezing machine (Research and Manufacturing, Tucson, AR) and promptly transferred to liquid nitrogen. Frog sacculi were fixed in 4% paraformaldehyde plus 0.5% glutaraldehyde in 0.1 M phosphate buffer, cryoprotected in 30% glycerol, and frozen in liquid propane. All frozen samples were freeze substituted in 1.5% uranyl acetate in methanol at –90°C, infiltrated with Lowicryl HM-20 resin at –45°C, and polymerized with ultraviolet light. Ultrathin sections were obtained on a Ultracut microtome, collected in nickel grids, and used in immunocytochemistry. Sections were picked up on grids and were incubated in 0.1% sodium borohydride and 50 mM glycine in 5 mM Tris and 0.1% Triton X-100, pH 7.4 (TBST) for 10 min, blocked with 10% normal goat serum in TBST for 10 min, and then incubated for 2 hr in primary antibody in 1% normal goat serum/TBST. Grids were rinsed three times in TBST, blocked with 1% goat serum/TBST for 10 min, and then were incubated in gold-labeled secondary antibodies (10 nm gold particles conjugated to goat anti-rabbit IgG, diluted 1:20; Sigma) in 1% serum/0.5% polyethylene glycol ($M_r = 20,000$) in TBST for 1 hr. Grids were washed in TBST and water, stained with uranyl acetate, and observed in JEOL 1010 or Zeiss EM902 transmission electron microscopes.

Other methods. Amino acid alignments were created with the MegAlign software (DNASTar, Madison, WI). By convention (Carafoli and Stauffer, 1993; Strehler and Zacharias, 2001), PMCA isozymes are referred by their isoform numeral (1–4), splicing region C variant by small case letters (*a–f*), and splicing region A variants by small case letters (*v–z*) (e.g., PMCA2av). In some cases we dropped one or both of the splicing region letters (e.g., PMCA2 with the F2N antibody, which recognized all PMCA2 forms, or PMCA2a with the F2a antibody, which recognized the splicing region C form but did not distinguish splicing region A variants).

RESULTS

Isolation of bullfrog sacculus PMCA cDNA clones

To identify PMCA isoforms of bullfrog sacculus hair cells, we first cloned several PMCA cDNAs from a sacculus library. A vestibular organ ~1 mm in diameter, the sacculus contains ~2500 hair cells and twice as many supporting cells surrounded by a simple epithelium consisting of nonsensory cells (Jacobs and Hudspeth, 1990). Because no bullfrog PMCA sequences were available, we first aligned mammalian PMCA sequences and identified several regions of complete amino acid identity in all isozymes from all available species. Using degenerate oligonucleotides corresponding to these regions, we amplified PMCA fragments by using RT-PCR with RNA isolated from bullfrog tissues. Using Df1 and Dr1 primers, we amplified a 1.2 kb PCR product from bullfrog liver; all clones isolated from this PCR reaction were shown by sequencing to be closely related to mammalian PMCA1 (data not shown). Df1 and Dr2 primers produced a single 0.35 kb bullfrog lung PCR product, shown by cloning and sequencing to be PMCA2 (data not shown). Although not exhaustive, this PCR screen produced useful clones for the identification of bullfrog PMCA1 and PMCA2.

We used the cloned PCR products to screen a bullfrog sacculus cDNA library. Using the PMCA1 probe, we identified ~150 positive clones (of 600,000 screened), one of which appeared to be full length (Figs. 1, 2). At sequencing, this clone included a 1.3 kb unspliced intron with 39 stop codons, shared approximately equally by all three frames. To confirm that this insert was not present in the mature mRNA, we sequenced three other library clones, none of which had the insert. The corrected sequence of 4575 bases, tentatively identified as PMCA1, encoded a protein of 1214 amino acids (134,560 Da). Using the PMCA2 probe, we isolated ~120 clones (of 700,000 screened), one of which appeared to be full length (Figs. 1, 2). This clone also was spliced incompletely; to determine the correct sequence, we sequenced two other library clones and five PCR products that were amplified from bullfrog sacculus cDNA. This large clone also had a two nucleotide deletion in the coding sequence; we sequenced two additional library clones to correct the sequence. Because few cDNAs have been cloned from *R. catesbeiana* libraries, we are uncertain whether the high frequency of untranslatable clones reflects a species phenomenon or simply poor luck. The corrected sequence of 4416 bases, tentatively identified as PMCA2, encoded a protein of 1213 amino acids (132,801 Da).

We analyzed predicted amino acid sequences of bullfrog PMCA clones to determine their relationship to known PMCA isozymes. For each bullfrog sequence, high amino acid identity with other PMCAs extends throughout all cytoplasmic loops and the 10 putative transmembrane domains. The phosphoenzyme intermediate aspartate residue present in all P-type ATPases (Carafoli and Stauffer, 1993) is conserved in bullfrog PMCA1 (residue 469) and PMCA2 (residue 513). In addition, key residues in transmembrane domains that are critical for Ca^{2+} pumping in PMCA [P422, E423, P426, A854, N879, M882, D883, Q971, and E975 in human PMCA4 (Guerini et al., 1996, 2000)] were present in bullfrog PMCA1 (residues 426, 427, 430, 860, 885, 888, 889, 975, and 981) and PMCA2 (residues 470, 471, 474, 903, 928, 931, 932, 1020, and 1024). By examining the N-terminal ~15 amino acids, which vary widely between different isozymes, we determined that our sequences were related most closely to PMCA1 and PMCA2 (Fig. 2A,B). At the amino acid level, bullfrog PMCA1 was 88.7% identical to rat PMCA1, and bullfrog PMCA2

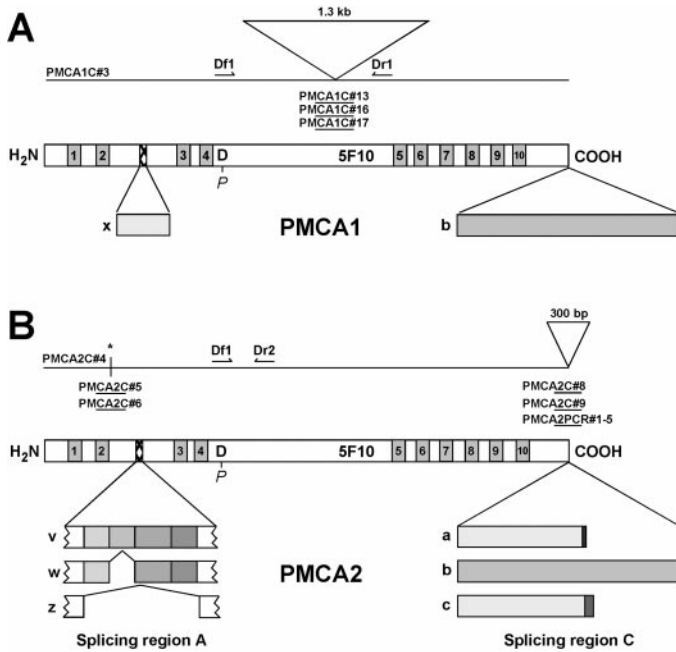


Figure 1. Cloning, structure, and alternative splicing of bullfrog PMCA isozymes. *Horizontal lines* indicate the extent of each cDNA that is sequenced, and the lines are aligned with a diagram of PMCA. Ten putative transmembrane domains (*shaded and numbered*), the phosphointermediate aspartate residue, and the 5F10 antibody binding site are indicated. *A*, Bullfrog sacculus PMCA1. *PMCA1C#3* indicates the clone that was used to derive most of the sequence for PMCA1; at the indicated 1.3 kb insert three other clones were sequenced too. *Df1* and *Dr1* indicate the positions of primers used in bullfrog liver RT-PCR to generate a PMCA1-specific probe. Splicing region A and C variants are indicated; we detected only *x* and *b*. *B*, Bullfrog sacculus PMCA2. *PMCA2C#4* indicates the clone that was used to derive most of the sequence for PMCA2. At the position indicated with an asterisk, the site of a 2 bp deletion, we sequenced two additional clones to confirm the correct sequence. In addition, we sequenced at least two library clones and five PCR products at splicing region C; only *PMCA2C#4* included a 300 bp unspliced insert. Variants *v*, *w*, and *z* were detected at splicing region A; variants *a*, *b*, and *c* were detected at splicing region C. *Df1* and *Dr2* indicate positions of primers used in bullfrog lung RT-PCR to generate a PMCA2-specific probe.

was 90.1% identical to rat PMCA2 (excepting alternative splicing regions A and C). These high sequence similarities establish that these isozyms are the bullfrog orthologs of PMCA1 and PMCA2.

Because of the importance of alternative splicing for PMCA regulation, we examined bullfrog sacculus PMCA1 and PMCA2 library clones and RT-PCR products from sacculus RNA for alternative splicing in regions A and C. For PMCA1 we detected only *x* and *b* variants at sites A and C (Figs. 1, 2, Table 2).

The splicing of PMCA2 was more complex. The full-length PMCA2 clone used a novel splicing region A variant, which we named *v* (Fig. 2C). Interestingly, the cDNA sequence of this variant begins with GT, the consensus splice donor dinucleotide, and ends with AG, the consensus splice acceptor dinucleotide (Green, 1991), suggesting that this sequence can be removed by alternative splicing. PCR analysis of >50 library clones indicated that all were of splice form *v*; in addition, RT-PCR analysis on bullfrog sacculus RNA showed that, in addition to the abundant *v*, PMCA2 splicing region A variants *w* and *z* were also present (Table 2). The *v* splice form was not specific to the inner ear, however, but also was found in bullfrog PMCA2 in brain, kidney,

lung, and retina (data not shown). The full-length PMCA2 clone incorporated the *a* splice variant in splicing region C (Fig. 2D); in other library clones *a* was the predominant form (98 of 103 clones), although *b* (4 of 103) and *c* (1 of 103) also were detected (Table 2). The *a* variant was the only splicing region C sequence identified by RT-PCR from sacculus RNA (Table 2). The two major PMCA transcripts in bullfrog sacculus hence encoded PMCA1bx and PMCA2av.

Localization of PMCA isoforms in bullfrog sacculus

To localize sacculus PMCA isoforms, we generated polyclonal antibodies selective for specific PMCA isozymes and splice variants (see Table 1) and used them for immunocytochemistry. Because fusion proteins used for generation of the F1N and F2N antibodies contained substantial sequence identity within the PMCA coding regions, we generated selective antibodies by performing sequential negative and positive affinity selection. Affinity-purified F1N recognized PMCA1, but not PMCA2, fusion proteins on immunoblots, whereas F2N recognized PMCA2, but not PMCA1, fusion proteins (data not shown). All antibodies also exhibited minimal cross-reactivity with other bullfrog sacculus proteins on blots (see Fig. 5A) and therefore were useful tools for localizing PMCA isozymes in bullfrog sacculus tissue.

Antibodies listed in Table 2 were used to detect PMCA isozymes within the bullfrog sacculus (Fig. 3). The PMCA1 N terminus antibody (F1N) strongly labeled basolateral membranes of hair cells (Fig. 3A); although PMCA1 usually was excluded completely from apical membranes, in rare instances we observed faint hair bundle labeling. Plasma membranes of supporting cells also were labeled with PMCA1-selective antibodies. An identical labeling pattern was observed with the use of an antibody (Fb) that was raised against a peptide sequence within the *b* splice forms of bullfrog PMCA1 and PMCA2 (Fig. 3B,C); because rat PMCA3b contains an identical sequence, this antibody also may detect a putative bullfrog PMCA3b. Labeling with either antibody was usually more intense in hair cells with larger cell body diameters. No labeling was observed when an irrelevant affinity-purified antibody was used; furthermore, the peptide used to raise the Fb antiserum blocked labeling with Fb. Both observations indicate the specificity of the PMCA1 immunolabeling.

Hair bundles in bullfrog sacculi were labeled intensely by antibodies selective for the PMCA2 N terminus (F2N; Fig. 3D,E), for the PMCA2a splice form (F2a; Fig. 3F), or for the PMCA2v splice form (F2v; Fig. 3G). Labeling often appeared to be concentrated near stereociliary tips, although sacculus-to-sacculus variability was noted. In addition, PMCA2-selective antibodies labeled the apical surface of the hair cell and a punctate ring surrounding the cuticular plate (Fig. 3D). For each of these features the labeling was specific; no labeling was observed with an irrelevant primary antibody, and the peptide used to raise the F2v antibody blocked hair cell labeling with F2v. No PMCA2 labeling was detected in the basolateral membrane of hair cells, in supporting cells, or in extramacular nonsensory epithelial cells of the bullfrog sacculus.

We used immunoelectron microscopy to localize PMCA1 and PMCA2 at the ultrastructural level (Fig. 4). Using postembedding labeling on freeze-substituted sacculus tissue, we found strong PMCA1 immunoreactivity along membranes of hair cells (Fig. 4A) and supporting cells (data not shown). PMCA2 immunoreactivity was observed only along stereocilia membranes and at the apical surface of the hair cell (Fig. 4B–E). No PMCA2 labeling

A

<i>fPMCA1bx</i>	MAANSVAYSCVKNSTENHSGTIGVQGLRDMELRSTDAOKIQEYGVDFYIC	57
<i>hPMCA1bx</i>	MDMANNVAYSGVKNSTLKAANNDGDFGIFLAFRLRAIMELRSTDAOKIQEYGVDFYIC	60
<i>rPMCA1bx</i>	MDMANNVAYSGVKNSTLKAANNDGDFGIFLAFRLRAIMELRSTDAOKIQEYGVDFYIC	60
<i>fPMCA1bx</i>	SRKTSPHDLSGSLADLERRQVFGKRIIPPKPKTKFLQVWEALQDVTLITLTAATL	117
<i>hPMCA1bx</i>	TKLKTSPIEGLSGNPADLERRQVFGKRIIPPKPKTKFLQVWEALQDVTLITLTAATL	120
<i>rPMCA1bx</i>	TKLKTSPIEGLSGNPADLERRQVFGKRIIPPKPKTKFLQVWEALQDVTLITLTAATL	120
<i>fPMCA1bx</i>	SLGLSPYRPPDEKFLDGGESGSAFSGPSSGSAWTEGAAITLLSVVYVLTAFNDWSKERQ	177
<i>hPMCA1bx</i>	SLGLSPYQPPEDNALCCQVSVGEELEGEGEETGWIETGAAITLLSVVYVLTAFNDWSKERQ	180
<i>rPMCA1bx</i>	SLGLSPYQPPEDNALCCQVSVGEELEGEGEETGWIETGAAITLLSVVYVLTAFNDWSKERQ	180
<i>fPMCA1bx</i>	FRGLQRIEQQKFTVIRGGQVQIPVADITVGGDIAQRYGGLPADGLFIQGNLKDIE	237
<i>hPMCA1bx</i>	FRGLQRIEQQKFTVIRGGQVQIPVADITVGGDIAQRYGGLPADGLFIQGNLKDIE	240
<i>rPMCA1bx</i>	FRGLQRIEQQKFTVIRGGQVQIPVADITVGGDIAQRYGGLPADGLFIQGNLKDIE	240
<i>fPMCA1bx</i>	SSLTGESDVKRNLKXPLLSTGTHVMEGSSAVVATLWNSOTGIIPTLLAG--EPEEK	297
<i>hPMCA1bx</i>	SSLTGESDVKRNLKXPLLSTGTHVMEGSSAVVATLWNSOTGIIPTLLAGGEEEEK	300
<i>rPMCA1bx</i>	SSLTGESDVKRNLKXPLLSTGTHVMEGSSAVVATLWNSOTGIIPTLLAGGEEEEK	300
<i>fPMCA1bx</i>	EKRRKRRKQDDVFNRRKAKAQDGAAMEMQLKSEGGDGGDGRKRNALPKRERS	354
<i>hPMCA1bx</i>	KDLKRRKRNKQDGAAMEMQLKSEGGDGGDGRKRNALPKRERS	360
<i>rPMCA1bx</i>	KDLKRRKRNKQDGAAMEMQLKSEGGDGGDGRKRNALPKRERS	360
<i>fPMCA1bx</i>	VGQSKTKLAVQIKGKGLMSAIVTIVLVYVVIDFTVVRKRWLAETPIYIVQVFKF	414
<i>hPMCA1bx</i>	VGQSKTKLAVQIKGKGLMSAIVTIVLVYVVIDFTVVRKRWLAETPIYIVQVFKF	420
<i>rPMCA1bx</i>	VGQSKTKLAVQIKGKGLMSAIVTIVLVYVVIDFTVVRKRWLAETPIYIVQVFKF	420
<i>fPMCA1bx</i>	HGVTVLVAVPEGLPLAVTISLAYSVKMMKMDNLRHLDACETMGNATAICSDKTGTL	474
<i>hPMCA1bx</i>	HGVTVLVAVPEGLPLAVTISLAYSVKMMKMDNLRHLDACETMGNATAICSDKTGTL	480
<i>rPMCA1bx</i>	HGVTVLVAVPEGLPLAVTISLAYSVKMMKMDNLRHLDACETMGNATAICSDKTGTL	480
<i>fPMCA1bx</i>	TNMRMTVQAVINIKHYRSTEDASLSESLKLNLTGSLVNCAYTSKILLPEKGGGLPRH	534
<i>hPMCA1bx</i>	TNMRMTVQAVINIKHYRSTEDASLSESLKLNLTGSLVNCAYTSKILLPEKGGGLPRH	540
<i>rPMCA1bx</i>	TNMRMTVQAVINIKHYRSTEDASLSESLKLNLTGSLVNCAYTSKILLPEKGGGLPRH	540
<i>fPMCA1bx</i>	VGNKTECALIGPLDRLKRDYQVRNEIPEDHFKVYTFNSRRMSVTLKNVDSYRQFS	594
<i>hPMCA1bx</i>	VGNKTECALIGPLDRLKRDYQVRNEIPEDHFKVYTFNSRRMSVTLKNVDSYRQFS	600
<i>rPMCA1bx</i>	VGNKTECALIGPLDRLKRDYQVRNEIPEDHFKVYTFNSRRMSVTLKNVDSYRQFS	600
<i>fPMCA1bx</i>	KGASEILKCKPKFLSANGKARVFRPRORDLVKTVIPKASEGLRTICLAFDFPAGEB	654
<i>hPMCA1bx</i>	KGASEILKCKPKFLSANGKARVFRPRORDLVKTVIPKASEGLRTICLAFDFPAGEB	660
<i>rPMCA1bx</i>	KGASEILKCKPKFLSANGKARVFRPRORDLVKTVIPKASEGLRTICLAFDFPAGEB	660
<i>fPMCA1bx</i>	EPDWNENDVNGTICIAVAVTEQVFRVPEVPAIKKQORAGITVVMVGDNTNARAAT	714
<i>hPMCA1bx</i>	EPDWNENDVNGTICIAVAVTEQVFRVPEVPAIKKQORAGITVVMVGDNTNARAAT	720
<i>rPMCA1bx</i>	EPDWNENDVNGTICIAVAVTEQVFRVPEVPAIKKQORAGITVVMVGDNTNARAAT	720
<i>fPMCA1bx</i>	KGILLHPGDFLCEGDFNRRIRNEKGELEQERDKIWPKLRVLRASSTDKHTLVKGI	774
<i>hPMCA1bx</i>	KGILLHPGDFLCEGDFNRRIRNEKGELEQERDKIWPKLRVLRASSTDKHTLVKGI	780
<i>rPMCA1bx</i>	KGILLHPGDFLCEGDFNRRIRNEKGELEQERDKIWPKLRVLRASSTDKHTLVKGI	780
<i>fPMCA1bx</i>	IDSTVQRQVAVTGDGNDGPAKRAQVDFAMGAGTQVAREASDILLTDNFTSIVR	834
<i>hPMCA1bx</i>	IDSTVQRQVAVTGDGNDGPAKRAQVDFAMGAGTQVAREASDILLTDNFTSIVR	840
<i>rPMCA1bx</i>	IDSTVQRQVAVTGDGNDGPAKRAQVDFAMGAGTQVAREASDILLTDNFTSIVR	840
<i>fPMCA1bx</i>	AVMGRNVYDTSKFLQPLTQVAVVAVTGAFTGACITQDSFLKAVQMLWNLIMDFLASE	894
<i>hPMCA1bx</i>	AVMGRNVYDTSKFLQPLTQVAVVAVTGAFTGACITQDSFLKAVQMLWNLIMDFLASE	900
<i>rPMCA1bx</i>	AVMGRNVYDTSKFLQPLTQVAVVAVTGAFTGACITQDSFLKAVQMLWNLIMDFLASE	900
<i>fPMCA1bx</i>	ALATEPPTESLILKPKYGRNKLRSMMNLLGHAFVQVAVVTLFAGEKFFDIDSGR	954
<i>hPMCA1bx</i>	ALATEPPTESLILKPKYGRNKLRSMMNLLGHAFVQVAVVTLFAGEKFFDIDSGR	960
<i>rPMCA1bx</i>	ALATEPPTESLILKPKYGRNKLRSMMNLLGHAFVQVAVVTLFAGEKFFDIDSGR	960
<i>fPMCA1bx</i>	NAPLHAPPSEHYTVTNTFVWMLFNEINARKIHCERNVFEIENNAICTVIGLTFVQ	1014
<i>hPMCA1bx</i>	NAPLHAPPSEHYTVTNTFVWMLFNEINARKIHCERNVFEIENNAICTVIGLTFVQ	1020
<i>rPMCA1bx</i>	NAPLHAPPSEHYTVTNTFVWMLFNEINARKIHCERNVFEIENNAICTVIGLTFVQ	1020
<i>fPMCA1bx</i>	LIIVQGGKFPSCSELSIQWLSIFLQKSTLNGQLSITPISRLKFLKAGHGQCKE	1074
<i>hPMCA1bx</i>	LIIVQGGKFPSCSELSIQWLSIFLQKSTLNGQLSITPISRLKFLKAGHGQCKE	1080
<i>rPMCA1bx</i>	LIIVQGGKFPSCSELSIQWLSIFLQKSTLNGQLSITPISRLKFLKAGHGQCKE	1080
<i>fPMCA1bx</i>	IEPEELAEVETDHAERELRRGQILWFRGLNRITQIRVNAFRSSYLEGKXPSSRS	1134
<i>hPMCA1bx</i>	IEPEELAEVETDHAERELRRGQILWFRGLNRITQIRVNAFRSSYLEGKXPSSRS	1140
<i>rPMCA1bx</i>	IEPEELAEVETDHAERELRRGQILWFRGLNRITQIRVNAFRSSYLEGKXPSSRS	1140
<i>fPMCA1bx</i>	IHNFMTHPEFRLEDSEPIFLDIDTDAEDDAPTRKSSPPSPHNNNAVDSGHIHTEM	1194
<i>hPMCA1bx</i>	IHNFMTHPEFRLEDSEPIFLDIDTDAEDDAPTRKSSPPSPHNNNAVDSGHIHTEM	1200
<i>rPMCA1bx</i>	IHNFMTHPEFRLEDSEPIFLDIDTDAEDDAPTRKSSPPSPHNNNAVDSGHIHTEM	1200
<i>fPMCA1bx</i>	NRSATSSSGPLSLETSL	1214
<i>hPMCA1bx</i>	NRSATSSSGPLSLETSL	1220
<i>rPMCA1bx</i>	NRSATSSSGPLSLETSL	1220

B

<i>fPMCA2av</i>	MGMVNSDFYSYKQBNFVNLHSGTIGVQGLRDMELRSTDAOKIQEYGVDFYIC	60
<i>hPMCA2az</i>	MDMTNSDFYSYKQBNFVNLHSGTIGVQGLRDMELRSTDAOKIQEYGVDFYIC	60
<i>rPMCA2aw</i>	MDMTNSDFYSYKQBNFVNLHSGTIGVQGLRDMELRSTDAOKIQEYGVDFYIC	60
<i>fPMCA2av</i>	KPSFVGLPCTADPDKRQVYVSNFIPPKPKTKFLQVWEALQDVTLITLTAATL	120
<i>hPMCA2az</i>	KPSFVGLPCTADPDKRQVYVSNFIPPKPKTKFLQVWEALQDVTLITLTAATL	120
<i>rPMCA2aw</i>	KPSFVGLPCTADPDKRQVYVSNFIPPKPKTKFLQVWEALQDVTLITLTAATL	120
<i>fPMCA2av</i>	LSFYRPPDEKFLDGGESGSAFSGPSSGSAWTEGAAITLLSVVYVLTAFNDWSKERQ	180
<i>hPMCA2az</i>	LSFYRPPDEKFLDGGESGSAFSGPSSGSAWTEGAAITLLSVVYVLTAFNDWSKERQ	180
<i>rPMCA2aw</i>	LSFYRPPDEKFLDGGESGSAFSGPSSGSAWTEGAAITLLSVVYVLTAFNDWSKERQ	180
<i>fPMCA2av</i>	FRGLQRIEQQKFTVIRGGQVQIPVADITVGGDIAQRYGGLPADGLFIQGNLKDIE	240
<i>hPMCA2az</i>	FRGLQRIEQQKFTVIRGGQVQIPVADITVGGDIAQRYGGLPADGLFIQGNLKDIE	240
<i>rPMCA2aw</i>	FRGLQRIEQQKFTVIRGGQVQIPVADITVGGDIAQRYGGLPADGLFIQGNLKDIE	240
<i>fPMCA2av</i>	SSLTGESDVKRNLKXPLLSTGTHVMEGSSAVVATLWNSOTGIIPTLLAG--EPEEK	300
<i>hPMCA2az</i>	SSLTGESDVKRNLKXPLLSTGTHVMEGSSAVVATLWNSOTGIIPTLLAGGEEEEK	300
<i>rPMCA2aw</i>	SSLTGESDVKRNLKXPLLSTGTHVMEGSSAVVATLWNSOTGIIPTLLAGGEEEEK	300
<i>fPMCA2av</i>	EKRRKRRKQDDVFNRRKAKAQDGAAMEMQLKSEGGDGGDGRKRNALPKRERS	360
<i>hPMCA2az</i>	KDLKRRKRNKQDGAAMEMQLKSEGGDGGDGRKRNALPKRERS	360
<i>rPMCA2aw</i>	KDLKRRKRNKQDGAAMEMQLKSEGGDGGDGRKRNALPKRERS	360
<i>fPMCA2av</i>	VGQSKTKLAVQIKGKGLMSAIVTIVLVYVVIDFTVVRKRWLAETPIYIVQVFKF	420
<i>hPMCA2az</i>	VGQSKTKLAVQIKGKGLMSAIVTIVLVYVVIDFTVVRKRWLAETPIYIVQVFKF	420
<i>rPMCA2aw</i>	VGQSKTKLAVQIKGKGLMSAIVTIVLVYVVIDFTVVRKRWLAETPIYIVQVFKF	420
<i>fPMCA2av</i>	HGVTVLVAVPEGLPLAVTISLAYSVKMMKMDNLRHLDACETMGNATAICSDKTGTL	480
<i>hPMCA2az</i>	HGVTVLVAVPEGLPLAVTISLAYSVKMMKMDNLRHLDACETMGNATAICSDKTGTL	480
<i>rPMCA2aw</i>	HGVTVLVAVPEGLPLAVTISLAYSVKMMKMDNLRHLDACETMGNATAICSDKTGTL	480
<i>fPMCA2av</i>	TNMRMTVQAVINIKHYRSTEDASLSESLKLNLTGSLVNCAYTSKILLPEKGGGLPRH	540
<i>hPMCA2az</i>	TNMRMTVQAVINIKHYRSTEDASLSESLKLNLTGSLVNCAYTSKILLPEKGGGLPRH	540
<i>rPMCA2aw</i>	TNMRMTVQAVINIKHYRSTEDASLSESLKLNLTGSLVNCAYTSKILLPEKGGGLPRH	540
<i>fPMCA2av</i>	VGNKTECALIGPLDRLKRDYQVRNEIPEDHFKVYTFNSRRMSVTLKNVDSYRQFS	600
<i>hPMCA2az</i>	VGNKTECALIGPLDRLKRDYQVRNEIPEDHFKVYTFNSRRMSVTLKNVDSYRQFS	600
<i>rPMCA2aw</i>	VGNKTECALIGPLDRLKRDYQVRNEIPEDHFKVYTFNSRRMSVTLKNVDSYRQFS	600
<i>fPMCA2av</i>	KGASEILKCKPKFLSANGKARVFRPRORDLVKTVIPKASEGLRTICLAFDFPAGEB	660
<i>hPMCA2az</i>	KGASEILKCKPKFLSANGKARVFRPRORDLVKTVIPKASEGLRTICLAFDFPAGEB	660
<i>rPMCA2aw</i>	KGASEILKCKPKFLSANGKARVFRPRORDLVKTVIPKASEGLRTICLAFDFPAGEB	660
<i>fPMCA2av</i>	EPDWNENDVNGTICIAVAVTEQVFRVPEVPAIKKQORAGITVVMVGDNTNARAAT	720
<i>hPMCA2az</i>	EPDWNENDVNGTICIAVAVTEQVFRVPEVPAIKKQORAGITVVMVGDNTNARAAT	720
<i>rPMCA2aw</i>	EPDWNENDVNGTICIAVAVTEQVFRVPEVPAIKKQORAGITVVMVGDNTNARAAT	720
<i>fPMCA2av</i>	KGILLHPGDFLCEGDFNRRIRNEKGELEQERDKIWPKLRVLRASSTDKHTLVKGI	780
<i>hPMCA2az</i>	KGILLHPGDFLCEGDFNRRIRNEKGELEQERDKIWPKLRVLRASSTDKHTLVKGI	780
<i>rPMCA2aw</i>	KGILLHPGDFLCEGDFNRRIRNEKGELEQERDKIWPKLRVLRASSTDKHTLVKGI	780
<i>fPMCA2av</i>	IDSTVQRQVAVTGDGNDGPAKRAQVDFAMGAGTQVAREASDILLTDNFTSIVR	840
<i>hPMCA2az</i>	IDSTVQRQVAVTGDGNDGPAKRAQVDFAMGAGTQVAREASDILLTDNFTSIVR	840
<i>rPMCA2aw</i>	IDSTVQRQVAVTGDGNDGPAKRAQVDFAMGAGTQVAREASDILLTDNFTSIVR	840
<i>fPMCA2av</i>	AVMGRNVYDTSKFLQPLTQVAVVAVTGAFTGACITQDSFLKAVQMLWNLIMDFLASE	900
<i>hPMCA2az</i>	AVMGRNVYDTSKFLQPLTQVAVVAVTGAFTGACITQDSFLKAVQMLWNLIMDFLASE	900
<i>rPMCA2aw</i>	AVMGRNVYDTSKFLQPLTQVAVVAVTGAFTGACITQDSFLKAVQMLWNLIMDFLASE	900
<i>fPMCA2av</i>	ALATEPPTESLILKPKYGRNKLRSMMNLLGHAFVQVAVVTLFAGEKFFDIDSGR	960
<i>hPMCA2az</i>	ALATEPPTESLILKPKYGRNKLRSMMNLLGHAFVQVAVVTLFAGEKFFDIDSGR	960
<i>rPMCA2aw</i>	ALATEPPTESLILKPKYGRNKLRSMMNLLGHAFVQVAVVTLFAGEKFFDIDSGR	960
<i>fPMCA2av</i>	NAPLHAPPSEHYTVTNTFVWMLFNEINARKIHCERNVFEIENNAICTVIGLTFVQ	1020
<i>hPMCA2az</i>	NAPLHAPPSEHYTVTNTFVWMLFNEINARKIHCERNVFEIENNAICTVIGLTFVQ	1020
<i>rPMCA2aw</i>	NAPLHAPPSEHYTVTNTFVWMLFNEINARKIHCERNVFEIENNAICTVIGLTFVQ	1020
<i>fPMCA2av</i>	LIIVQGGKFPSCSELSIQWLSIFLQKSTLNGQLSITPISRLKFLKAGHGQCKE	1080
<i>hPMCA2az</i>	LIIVQGGKFPSCSELSIQWLSIFLQKSTLNGQLSITPISRLKFLKAGHGQCKE	1080
<i>rPMCA2aw</i>	LIIVQGGKFPSCSELSIQWLSIFLQKSTLNGQLSITPISRLKFLKAGHGQCKE	1080
<i>fPMCA2av</i>	IEPEELAEVETDHAERELRRGQILWFRGLNRITQIRVNAFRSSYLEGKXPSSRS	1140
<i>hPMCA2az</i>	IEPEELAEVETDHAERELRRGQILWFRGLNRITQIRVNAFRSSYLEGKXPSSRS	1140
<i>rPMCA2aw</i>	IEPEELAEVETDHAERELRRGQILWFRGLNRITQIRVNAFRSSYLEGKXPSSRS	1140
<i>fPMCA2av</i>	IHNFMTHPEFRLEDSEPIFLDIDTDAEDDAPTRKSSPPSPHNNNAVDSGHIHTEM	1200
<i>hPMCA2az</i>	IHNFMTHPEFRLEDSEPIFLDIDTDAEDDAPTRKSSPPSPHNNNAVDSGHIHTEM	1200
<i>rPMCA2aw</i>	IHNFMTHPEFRLEDSEPIFLDIDTDAEDDAPTRKSSPPSPHNNNAVDSGHIHTEM	1200
<i>fPMCA2av</i>	NRSATSSSGPLSLETSL	1214
<i>hPMCA2az</i>	NRSATSSSGPLSLETSL	1214
<i>rPMCA2aw</i>	NRSATSSSGPLSLETSL	1214

C

<i>fPMCA2av</i>	SPKTSPHDLSGSLADLERRQVFGKRIIPPKPKTKFLQVWEALQDVTLITLTAATL	59
<i>fPMCA2w</i>	SRKTSPHDLSGSLADLERRQVFGKRIIPPKPKTKFLQVWEALQDVTLITLTAATL	48
<i>fPMCA2x</i>	SRKTSPHDLSGSLADLERRQVFGKRIIPPKPKTKFLQVWEALQDVTLITLTAATL	47
<i>fPMCA2y</i>	SRKTSPHDLSGSLADLERRQVFGKRIIPPKPKTKFLQVWEALQDVTLITLTAATL	40
<i>fPMCA2z</i>	SRKTSPHDLSGSLADLERRQVFGKRIIPPKPKTKFLQVWEALQDVTLITLTAATL	9
<i>fPMCA2v</i>	GTIYKAKQQDGAAG	74
<i>fPMCA2w</i>	SSSQKAKQQDGAAG	63
<i>fPMCA2x</i>	SSSQKAKQQDGAAG	32
<i>fPMCA2y</i>	SSSQKAKQQDGAAG	49
<i>fPMCA2z</i>	SSSQKAKQQDGAAG	18

D

<i>fPMCA2a</i>	STGIEVWNTFKSGASFGALRRQSSITSQNDVTVNSPVSISLNAISSPFSASAAAAG	60
<i>fPMCA2b</i>	STGIEVWNTFKSGASFGALRRQSSITSQNDVTVNSPVSISLNAISSPFSASAAAAG	7
<i>fPMCA2c</i>	STGIEVWNTFKSGASFGALRRQSSITSQNDVTVNSPVSISLNAISSPFSASAAAAG	60
<i>fPMCA2a</i>	GG	62
<i>fPMCA2b</i>	KAFRSYLEGKXPDSRTSIIHNFMTHPEFKIESQPHIIDDITLEDHPALKQSSQPP	67
<i>fPMCA2c</i>	HPRREGIP	68
<i>fPMCA2b</i>	SPNKNNAVDSGHIHTTETSASSTSSPFSISLNAISSPFSASAAAAG	106

Figure 2. Sequence analysis of bullfrog sacculus PMCA1bx and PMCA1av isozymes. *Highlighted* residues are those with identity to the aligned residues in at least one other PMCA. Splicing region A is *underlined in light gray*; splicing region C is *underlined in dark gray*. *A*, Bullfrog sacculus PMCA1bx (*fPMCA1bx*) aligned with human PMCA1bx (*hPMCA1bx*) and rat PMCA1bx (*rPMCA1bx*). Protein translation likely initiates at nucleotides 340–342; this is the first AUG codon of a long open reading frame and matches the second AUG of the mammalian PMCA1 sequence. *B*, Bullfrog sacculus PMCA2av (*fPMCA2av*) aligned with human PMCA2az (*hPMCA2az*) and rat PMCA2aw (*rPMCA2aw*). Nucleotides 328–330 correspond to the first AUG codon of a long open reading frame and match the known translation start site of mammalian PMCA2. *C*, Bullfrog sacculus splicing region A variant PMCA2av (*fPMCA2av*) aligned with rat PMCA2 splice region A variants *w*, *x*, *y*, and *z* (*rPMCA2w*, *rPMCA2x*, *rPMCA2y*, *rPMCA2z*, respectively). *D*, Bullfrog sacculus PMCA2 splice region C variants identified in the sacculus cDNA library.

Table 2. PMCA splice variants in bullfrog sacculus

Isozyme	Primers	Product size (bp)	Sacculus library	Sacculus cDNA
PMCA1x	F1A+2/F1A–2	326	++++	++++
PMCA1b	F1C+2/F1C–2	439	++++	++++
PMCA2v	F2A+2/F2A–2	410	++++	++++
PMCA2w	F2A+2/F2A–2	374	–	+
PMCA2z	F2A+2/F2A–2	239	–	+
PMCA2a	F2C+2/F2C–2	613	++++	++++
PMCA2b	F2C+2/F2C–2	386	+	–
PMCA2c	F2C+2/F2C–2	559	+	–

Isozymes: PMCA isoforms cloned from the sacculus cDNA library or amplified by RT-PCR from bullfrog sacculus RNA. Primers: The second pair of primers from a set of nested primers used for PCR. Product size: Expected PCR product sizes, given in base pairs, for each primer pair. Symbols signify abundant (++++; >80%), rare (+; <10%), and undetected (–) splice variants identified among sacculus library clones or PCR products from sacculus cDNA.

was seen along hair cell basolateral membranes or in supporting cells.

These immunolabeling data indicate that PMCA1b is a major isoform of hair cell basolateral membranes, and PMCA2a is a major isoform of hair bundles. Nevertheless, other PMCA isoforms, not detected with our antibodies, also could be present in sacculi. Although the NR3 antibody, raised against rat PMCA3, did not label bullfrog sacculi (data not shown), its cross-reactivity with a putative bullfrog PMCA3 isoform is unknown. Because immunolabeling cannot exclude the presence of PMCA3 or other isoforms, we turned to protein biochemistry to address the identity of hair bundle and somatic PMCA isoforms.

Protein immunoblot analysis of bullfrog sacculus PMCA isoforms

We analyzed PMCA1 and PMCA2 by protein immunoblotting, using each of the antibodies we previously had used for immunocytochemistry. With each antibody, as well as with the pan-PMCA antibody 5F10, we detected two to three bands of 130–170 kDa in whole bullfrog sacculus (Fig. 5A). Only the antibody against the ν form of PMCA2 (F2v) did not react efficiently with sacculus PMCA isoforms on immunoblots. The higher molecular mass forms of PMCA were not unexpected; PMCA isoforms (particularly PMCA2) often migrate in SDS-PAGE with a second band ~20 kDa larger than the expected protein product (Hilfiker et al., 1994).

Immunoprecipitation of bullfrog sacculus PMCA isoforms

We used an immunoprecipitation approach to confirm the identity of the principal PMCA isoforms of bullfrog hair cell somas and hair bundles. We precipitated PMCA isoforms from sacculus protein extracts with selective antibodies and then analyzed all precipitated and unprecipitated PMCA isoforms by immunoblotting with the pan-PMCA antibody 5F10. Because 5F10 cross-reacts with all four mammalian PMCA isoforms (Caride et al., 1996) as well as with PMCA isoforms in a wide range of species (Borke et al., 1989; de Talamoni et al., 1993; Benaim et al., 1995), we expected to analyze all sacculus PMCA isoforms simultaneously in these experiments.

To perform these immunoprecipitation experiments, we solubilized PMCA from tissue extracts with Triton X-100 and then sequentially precipitated each extract with isoform-selective PMCA antibodies. Proteins remaining in the supernatant after

immunoprecipitation were precipitated with acetone. For a comprehensive examination of PMCA isoforms, including those for which we had no specific antibody, we required complete acetone precipitation of remaining PMCA isoforms; by examining total PMCA before and after acetone precipitation, we demonstrated that this precipitation was efficient (Fig. 5B). PMCA was detected in immunoprecipitates and the final supernatant by SDS-PAGE and immunoblotting with 5F10.

We first determined which PMCA isoforms were present in bullfrog residual macula (Gillespie and Hudspeth, 1991), the sacculus tissue remaining after hair bundle isolation (Fig. 5C,D). This tissue contains bundleless hair cells, supporting cells, and a few nonsensory extramacular cells. When F1N was used as the first precipitating antibody (Fig. 5C), most of the PMCA in the residual macula extract was precipitated. When Fb was used before F1N, Fb precipitated most PMCA (Fig. 5D). PMCA1 precipitated by F1N, but not Fb (Fig. 5D), may be another PMCA1 splicing region C variant, although our molecular cloning and RT-PCR experiments only detected the *b* form. Alternatively, proteolysis might have eliminated the Fb antibody epitope. Very little PMCA2 was present in the residual macula tissue (Fig. 5C,D). PMCA precipitated by Fb, but not F1N, in Figure 5C may be PMCA3b; in addition, there was a small amount of PMCA that was not precipitated by any of our antibodies, which may have been other PMCA3 splice variants or PMCA4. The results from these immunoprecipitation experiments thus demonstrate that the principal residual macula isoform is PMCA1b, consistent with immunocytochemical results (see Fig. 3).

We next examined the identity of PMCA isoforms in purified hair bundles (Gillespie and Hudspeth, 1991). F2a immunoprecipitated all PMCA from a hair bundle protein extract; neither F2N nor F1N precipitated any additional PMCA, and no PMCA remained in the final supernatant (Fig. 5E). Besides showing that the only bundle PMCA is PMCA2a, the data also demonstrated the effectiveness of the hair bundle purification. Although only a few sacculus equivalents of PMCA1 give a strong 5F10 signal (e.g., Fig. 5C,D), we saw no significant 5F10 reactivity in 47 sacculus equivalents of hair bundles when PMCA1 antibodies were used for immunoprecipitation (Fig. 5E).

Our experimental conditions insured that all 5F10-reactive PMCA isoforms from the sacculus, presumably all sacculus PMCA molecules, could be analyzed in a single experiment. The data therefore established conclusively that PMCA2a is the hair bundle PMCA isoform in the bullfrog sacculus and that PMCA1b accounts for a majority of the PMCA in the rest of the sacculus.

PMCA isoforms in rat organ of Corti and utricle

To determine the generality of the bullfrog sacculus localization results, we examined PMCA isoform distribution in auditory and vestibular organs of rat, where we could use antibodies selective for each of the four known PMCA isoforms (see Table 1; Filoteo et al., 1997). To confirm that PMCA isoforms 1–4 are the only PMCA isoforms present in vertebrates, we used the BLAST search algorithm (Altschul et al., 1990) to search for genes related to PMCA within the human genome (<http://www.ncbi.nlm.nih.gov/genome/seq/HsBlast.html>). Each human PMCA cDNA identified only its own gene and the other three PMCA genes and, in one case, a sarcoplasmic–endoplasmic reticulum Ca^{2+} -ATPase (data not shown). These data suggest that, if any other PMCA isoforms are present in the genome, they are related much more distantly to PMCA isoforms 1–4 than each is to the others. By performing immunocytochem-

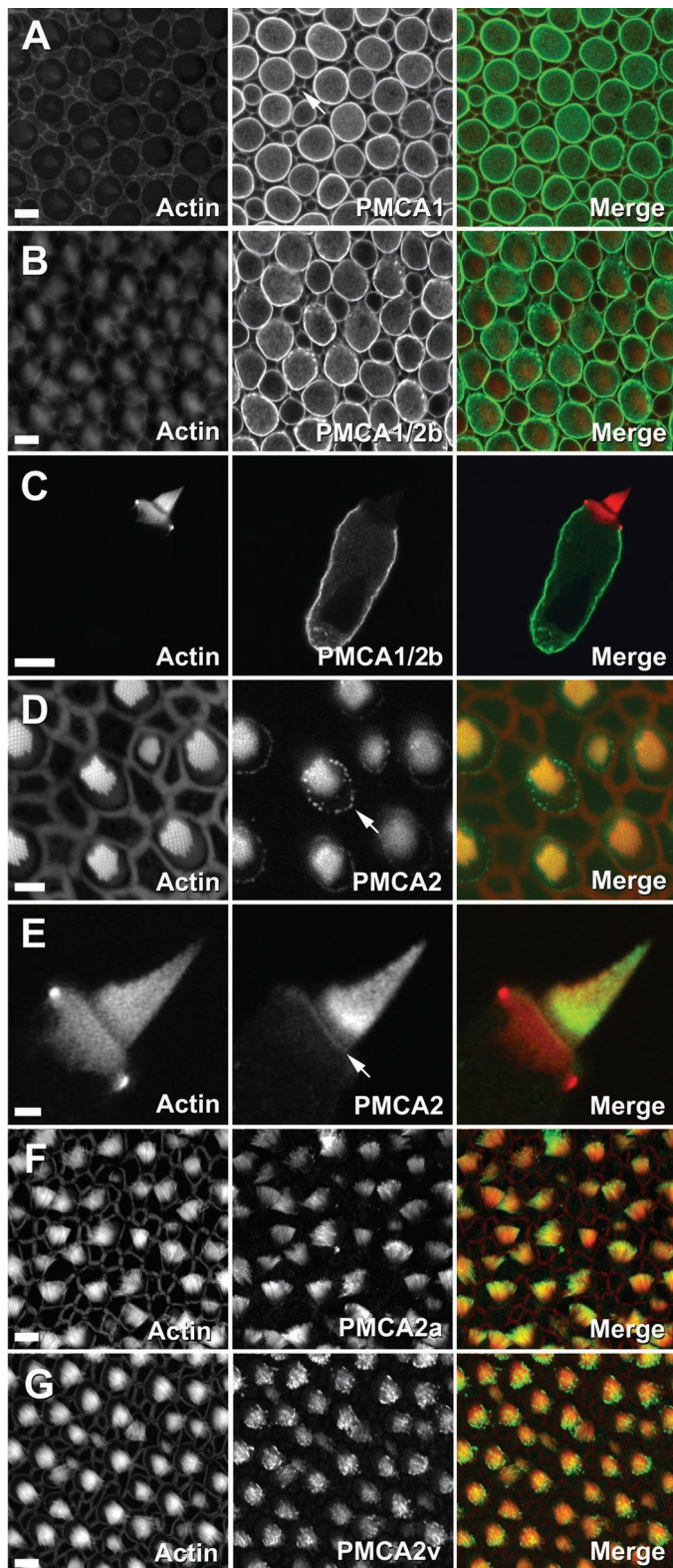


Figure 3. Localization of bullfrog sacculus PMCA isozymes by immunofluorescence. *Left columns*, Actin (FITC-phalloidin); *middle columns*, PMCA; *right columns*, combined actin (red) and PMCA (green). *A*, F1N labeling for PMCA1. Shown is a cross section through a bullfrog sacculus; plasma membranes of hair cells and supporting cells (*arrow*) are labeled. *B*, Fb labeling for PMCA1b and PMCA2b. Shown is a cross section through bullfrog sacculus hair cells and supporting cells. *C*, Fb labeling for PMCA1b and PMCA2b. The hair bundle is unlabeled. *D*, F2N labeling for PMCA2. Shown are apical surfaces of bullfrog sacculus hair

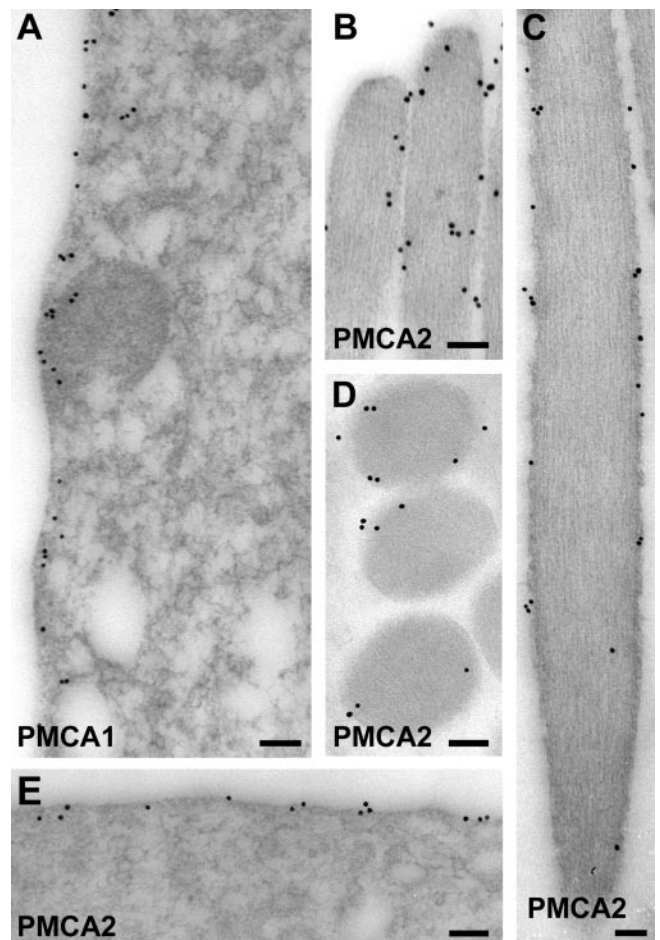


Figure 4. Localization of bullfrog sacculus PMCA isozymes by immunoelectron microscopy. *A*, F1N labeling for PMCA1. Shown is a hair cell basolateral membrane. *B–E*, F2N labeling for PMCA2. *B*, Stereociliary tips. *C*, Stereocilium shaft. *D*, Cross section through stereocilia. *E*, Hair cell apical surface. Scale bars, 100 nm.

istry in rat tissues, we therefore were able to use antibodies against all PMCA gene products.

We first examined the organ of Corti, the sensory epithelium of the mammalian auditory system. In this tissue PMCA2 immunoreactivity was restricted in hair cells to bundles and apical surfaces (Fig. 6*B,F*). Labeling was intense in bundles of outer hair cells; bundles of inner hair cells had far less immunoreactivity, which occasionally appeared to be concentrated at stereociliary tips. An identical labeling pattern was observed with F2a (Fig. 6*C*), indicating that PMCA2a is present in hair bundles.

PMCA1 was located on basolateral membranes of inner hair cells (Fig. 6*A*) but was absent from their hair bundles. PMCA1 was never observed in basolateral membranes of outer hair cells, although on occasion we observed very low PMCA1 labeling on apical surfaces (data not shown). PMCA1 also was detected in the

←

cells and supporting cells; there is PMCA2 labeling in stereocilia and in the pericuticular necklace (*arrow*). *E*, F2N labeling for PMCA2. Shown is labeling in an isolated hair cell; there is strong labeling near the base of the tallest stereocilia and labeling along the apical surface (*arrow*). *F*, F2a labeling for PMCA2a. Shown is a whole-mount view of hair bundles. *G*, F2v for PMCA2v. Shown is a whole-mount view of hair bundles. Scale bars: *A–C*, *F*, *G*, 10 μ m; *D*, 5 μ m; *E*, 2 μ m.

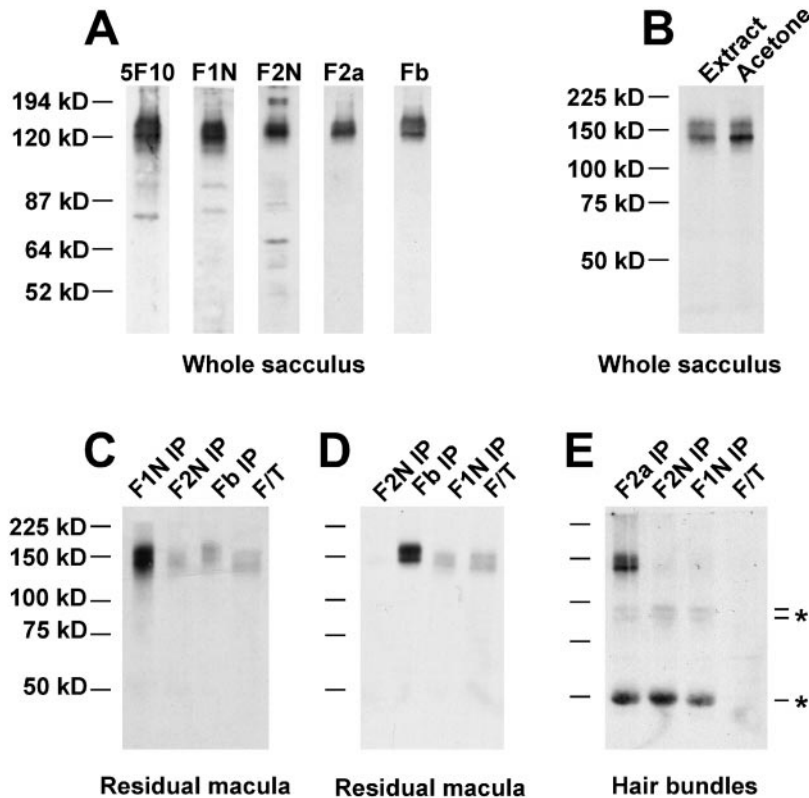


Figure 5. Protein immunoblot and immunoprecipitation analysis of bullfrog sacculus PMCA isozymes. *A*, Immunoblotting of whole bullfrog sacculus with 5F10 and isozyme-selective antibodies. One sacculus equivalent was used for each lane. *B*, 5F10 immunoblot indicating efficiency of acetone precipitation. Extracts from bullfrog sacculi either were added directly to sample buffer (*Extract*) or were precipitated with acetone (*Acetone*). Band intensities were similar, indicating that acetone precipitation was efficient. *C*, 5F10 detection of PMCA sequentially immunoprecipitated from a residual macula extract (five sacculus equivalents) with PMCA-selective antibodies. *F/T*, Flow through, those proteins that did not bind to any of the precipitating antibodies. *D*, Same immunoprecipitation and detection conditions as in *C*, except that the antibody order is changed. *E*, 5F10 detection of PMCA sequentially immunoprecipitated from a hair bundle extract (47 sacculus equivalents) with PMCA-selective antibodies. Indicated bands (*) were derived from the precipitating antibodies (data not shown).

Deiter's cell cups, where these cells cradle the base of outer hair cells. PMCA3 was observed only in inner hair cells; this isozyme was located prominently in a ring near the inner hair cell cuticular plate and at lower levels on the apical surface of inner hair cells (Fig. 6*D*). PMCA3 labeling was detected occasionally in the basolateral plasma membrane of inner hair cells and, at very low levels, in bundles of inner hair cells. Fb, which recognizes PMCA1b, 2b, and 3b, labeled the plasma membrane of inner hair cell somas and Deiter's cell cups in a similar pattern to that obtained with NR1 (data not shown). Surprisingly, Fb also labeled plasma membranes of inner and outer pillar cells as well as phalanges of Deiter's cells. None of the other antibodies against rat PMCA, which recognize all known isozymes, labeled these structures. PMCA4 was not detected in the organ of Corti.

We also examined PMCA in the rat utricle, a vestibular organ. PMCA2 immunoreactivity was intense in hair bundles and was absent from the rest of the tissue (Fig. 7*B,C*); furthermore, the vestibular bundle isoform was PMCA2a (Fig. 7*D*). F2N also strongly labeled hair bundles of the semicircular canal (Fig. 7*E*). Although many vestibular hair bundles were labeled intensely with antibodies against PMCA2, a subset of bundles had lower levels of this isozyme. These bundles usually belonged to hair cells that were immunoreactive for the calcium-binding protein calretinin; consistent with their generally smaller bundles, these cells may be immature type II hair cells (Dechesne et al., 1991, 1994). Basolateral membranes of utricular hair cells were labeled with antibodies against PMCA1 and PMCA3. PMCA1 labeling was observed at substantial levels only in a fraction of hair cells (Fig. 7*A*). By contrast, PMCA3 labeling was apparent in nearly all hair cells, although it was usually considerably lower in calretinin-positive cells (Fig. 7*D*). Like the supporting cells of the bullfrog sacculus, supporting cells of the rat utricle target PMCA1 to their plasma membranes, albeit at a significantly lower level than in hair cells. We did not observe PMCA4 expression in rat vestibular organs.

We observed no immunolabeling of rat vestibular organs or organ of Corti by using an antibody against the ν splice variant of PMCA2 (data not shown). To confirm the absence of this splice form in rat hair cells, we performed RT-PCR on cochlea and utricle RNA to examine the splicing in region A. Using primers that flank splicing region A, we found no evidence for the expression of a rat ν form, although we did amplify, clone, and sequence w , x , and z forms (see Table 2).

Because we used antibodies against all four known PMCA isozymes (see Table 1), these data demonstrated that PMCA2a is the major bundle isoform in rat hair cells, as it is in bullfrog hair cells. By contrast, hair cells and supporting cells of the rat cochlea and vestibular system use a variety of PMCA isozymes in their basolateral membranes.

DISCUSSION

Hair cells target PMCA isozymes to discrete subcellular locations. In auditory and vestibular epithelia PMCA2a was the predominant calcium pump of hair bundles; by contrast, PMCA1b was observed only on plasma membranes of hair cell and supporting cell somas. Although we did not detect PMCA4 in auditory or vestibular epithelia, we did observe PMCA3 in mammalian hair cells. This differential localization of PMCA isoforms presumably reflects the need for independent Ca^{2+} regulation in different cells and within distinct subcellular compartments, such as the hair bundle (Fig. 8).

PMCA isoforms in the bullfrog sacculus

In bullfrog sacculus we found prominent expression of PMCA1bx and PMCA2av. Furthermore, sequential immunoprecipitation experiments suggested that PMCA2a is the only plasma membrane calcium pump in bullfrog hair bundles, and PMCA1b is the principal one of the basolateral membrane. This conclusion rests on the assumption that the monoclonal antibody 5F10 recognizes

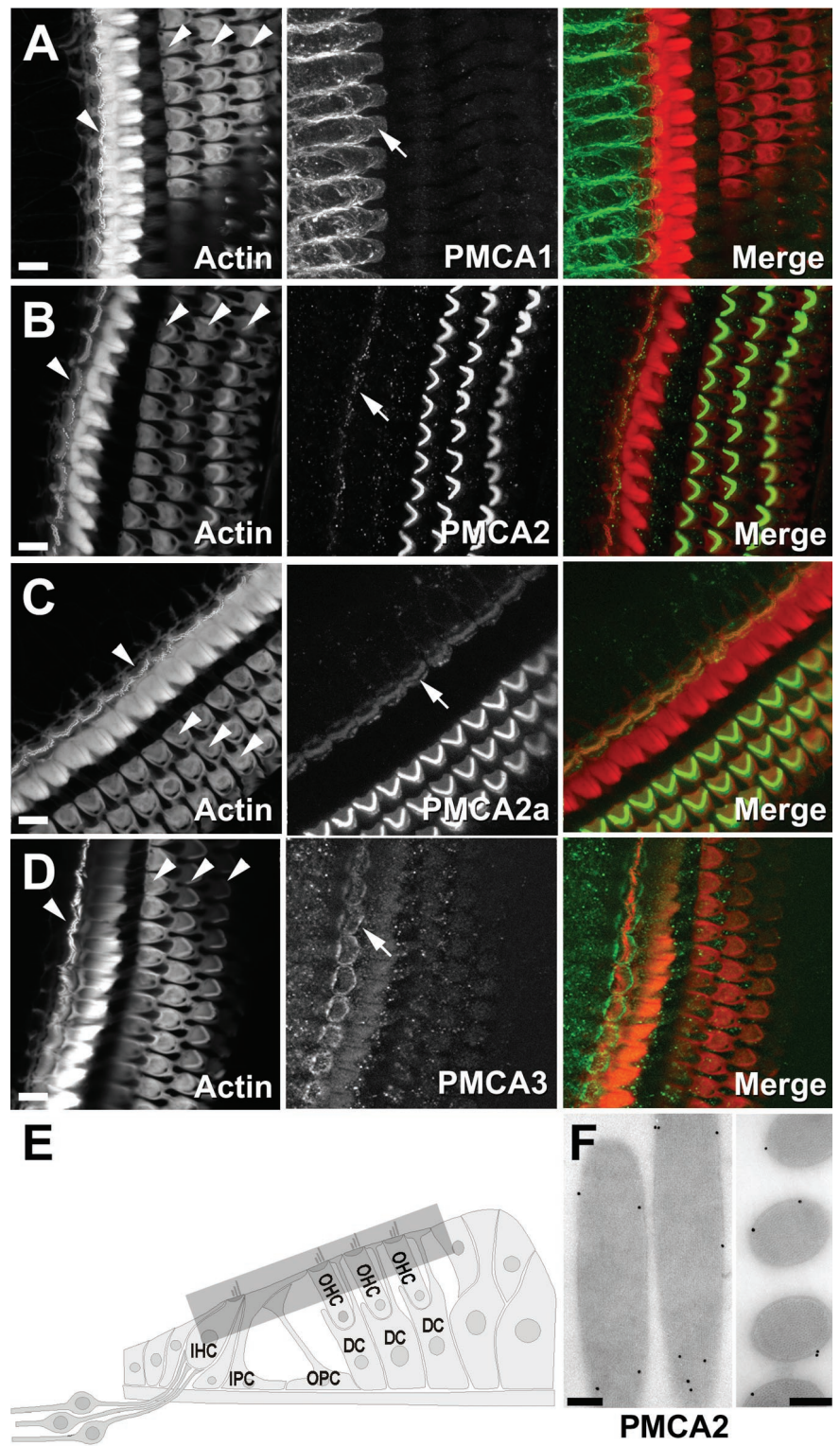


Figure 6. Localization of rat organ of Corti PMCA isozymes by immunofluorescence and immunoelectron microscopy. Shown are cross sections through the rat organ of Corti. *Left columns*, Actin (FITC-phalloidin); *middle columns*, PMCA immunolabeling; *right columns*, combined actin (red) and PMCA (green). In the *left columns* the *arrowheads* point to the single row of inner hair cells and three rows of outer hair cells. *A*, NR1 labeling for PMCA1. Basolateral membranes of inner hair cells were labeled (*arrow*). *B*, NR2 labeling for PMCA2. Hair bundles of outer hair cells were labeled intensely; modest immunoreactivity was present on bundles of inner hair cells (*arrow*). *C*, F2a labeling for PMCA2a. Stereocilia of inner (*arrow*) and outer hair cells were labeled. *D*, NR3 labeling for PMCA3. Immunoreactivity was restricted to the pericuticular necklace (*arrow*) and bundles of inner hair cells. Bundle labeling was very low, when detectable. *E*, Diagram of the mammalian organ of Corti indicating the approximate region of projected confocal images. *IHC*, Inner hair cell; *OHC*, outer hair cell; *IPC*, inner pillar cell; *OPC*, outer pillar cell; *DC*, Deiter's cell. *F*, NR2 labeling for PMCA2; shown is postembedding immunoelectron microscopy on freeze-substituted rat cochlear tissue. *Left*, Stereociliary shafts and tip. *Right*, Cross section. PMCA2 is associated with plasma membranes. Scale bars: *A–D*, 10 μ m; *F*, 200 nm.

and binds all PMCA isoforms on protein immunoblots. Although true for all four mammalian PMCA isoforms (Caride et al., 1996), only when 5F10 can be tested against all bullfrog PMCAs can we determine conclusively that no other PMCA isoforms are located in hair bundles or hair cell plasma membranes. Nevertheless, the conclusion that PMCA2a is the bundle isoform was supported strongly by localization in the rat auditory and vestibular systems; with the use of antibodies against all four PMCA isozymes, PMCA2a was the only prominent hair bundle PMCA isozyme.

PMCA isoforms in the mammalian vestibular and auditory systems

In the organ of Corti the pan-PMCA monoclonal antibody 5F10 was used to localize PMCA to the surface of stereocilia of both inner and outer hair cells as well as along the basolateral membrane of inner hair cells (Crouch and Schulte, 1995; Apicella et al., 1997; Street et al., 1998). 5F10 immunoreactivity was absent from hair bundles of mutant mice that lacked PMCA2 expression,

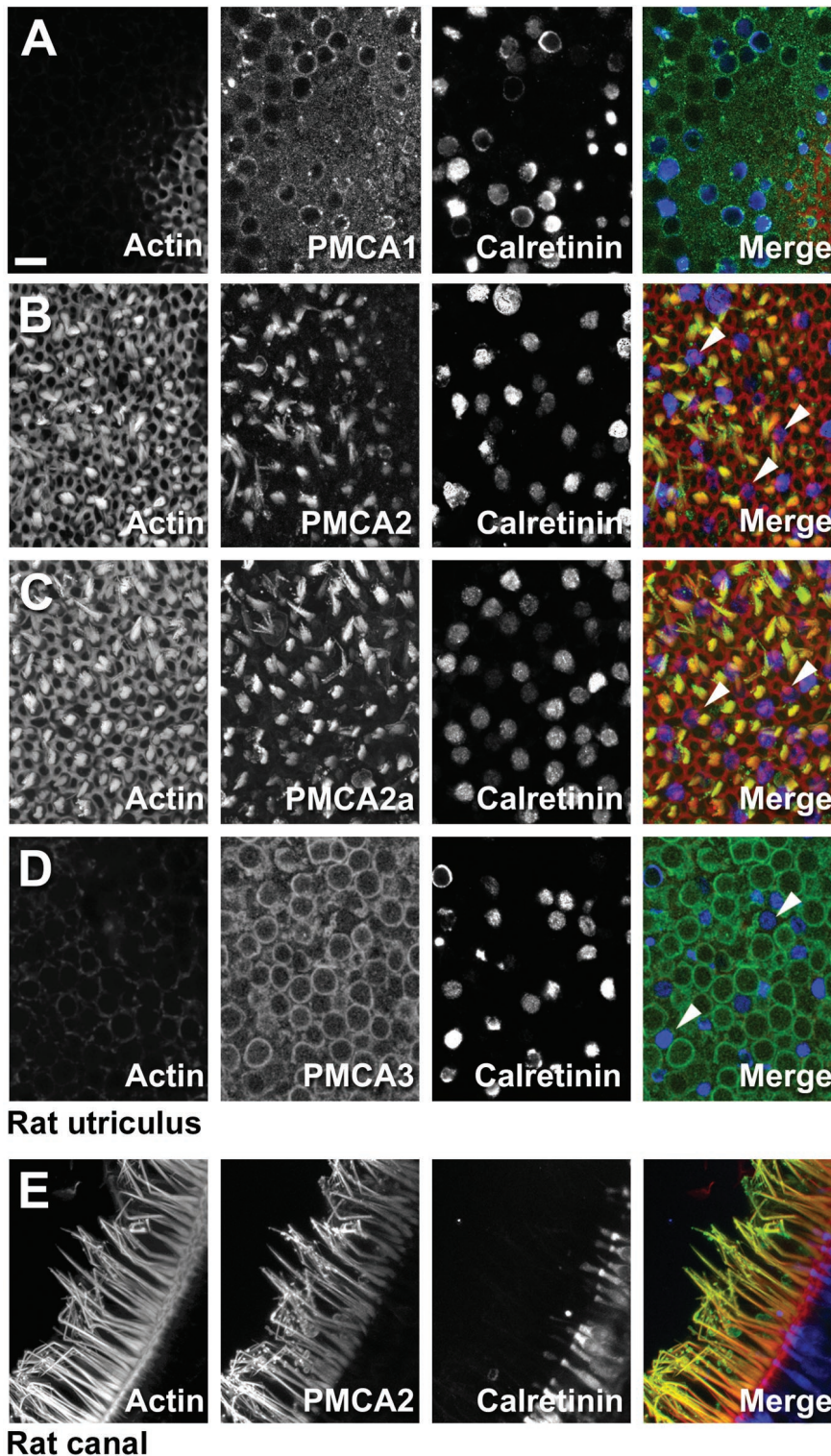


Figure 7. Localization of rat vestibular PMCA isozymes by immunofluorescence. Cross sections through the rat utricle (*A–D*) and ampulla (*E*). *Left columns*, Actin (FITC-phalloidin); *middle left columns*, PMCA immunolabeling; *middle right columns*, calretinin immunolabeling; *right columns*, combined actin (red), PMCA (green), and calretinin (blue). *A*, NR1 labeling for PMCA1. Shown is a single confocal section. *B*, NR2 labeling for PMCA2. Shown is a projection of confocal sections covering $\sim 10 \mu\text{m}$ depth. Some hair bundles with low PMCA2 labeling belong to hair cells with high levels of calretinin (*arrowheads*). *C*, F2a labeling for PMCA2a. Shown is a projection of confocal sections covering $\sim 10 \mu\text{m}$ depth. Hair bundles with low PMCA2a and high calretinin labeling are indicated by *arrowheads*. *D*, NR3 labeling for PMCA3. Shown is a single confocal section. Hair cells with low PMCA3 expression have high calretinin levels (*arrowheads*). *E*, NR2 labeling for PMCA2. Scale bar, $10 \mu\text{m}$ (applies to all panels).

suggesting that this isozyme was prominent there (Street et al., 1998; Takahashi and Kitamura, 1999). RT-PCR experiments identified PMCA1–4 gene products, as well as many of their splice variants, in cochlea (Crouch and Schulte, 1996; Furuta et al., 1998); furthermore, *in situ* hybridization experiments localized PMCA2 transcripts to outer hair cells and PMCA1 to inner hair cells (Furuta et al., 1998). Because PMCA labeling was prominent in outer hair cells and PMCA2 transcripts were localized there and because PMCA2b was the only PMCA2 splice

variant identified in cochlea by RT-PCR, PMCA2b was hypothesized to be the stereocilia isoform (Crouch and Schulte, 1995; Furuta et al., 1998).

We show here, however, that PMCA2a is the major hair bundle PMCA not only in bullfrog sacculus but also in bundles of rat organ of Corti, utricle, and semicircular canal. Furthermore, we did not observe hair bundle labeling using an antibody that recognizes the *b* sequence shared by PMCA1b, PMCA2b, and PMCA3b. Because PMCA2 protein levels in inner hair cells are

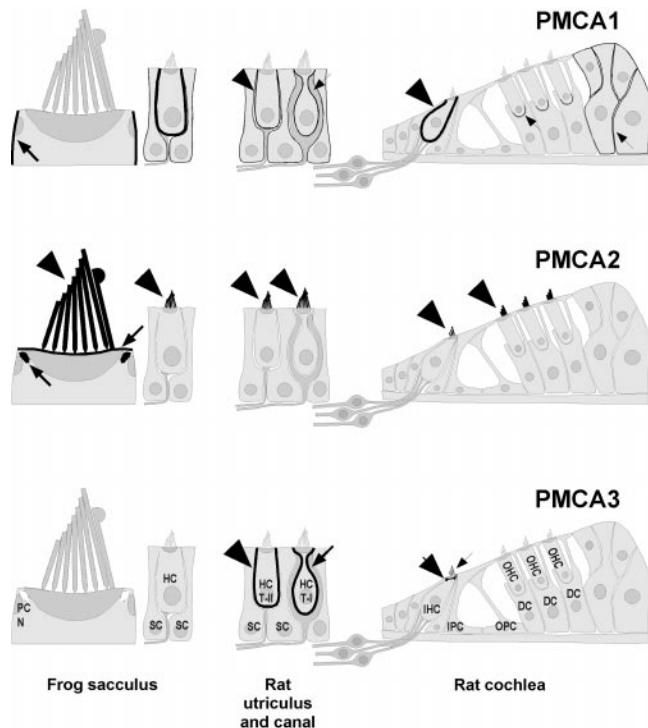


Figure 8. Schematic localization of PMCA isozymes in auditory and vestibular epithelia. PMCA isozyme location is indicated by *black shading* (intensity indicates relative amount) and by *arrows or arrowheads*. *PMCA1* was located on basolateral membranes of bullfrog and rat vestibular hair cells as well as inner hair cells. Occasional labeling was seen on outer hair cell apical surfaces. *PMCA2* was present at high levels on bundles and apical surfaces of vestibular hair cells and outer hair cells; levels were lower on bundles of inner hair cells. In bullfrog sacculus the labeling was also present in the pericuticular necklace. *PMCA3* labeling was strong in the basolateral membranes of rat utricle hair cells; this isozyme was also present near the apical surface of inner hair cells and, in some instances, in their bundles. *HC*, Hair cell; *PC N*, pericuticular necklace; *SC*, supporting cell; *HC T-I*, type I vestibular hair cell; *HC T-II*, type II vestibular hair cell.

very low, it is not surprising that *in situ* hybridization experiments did not detect *PMCA2* transcripts in inner hair cells (Furuta et al., 1998). Although we occasionally detected low levels of *PMCA1* in bundles of outer hair cells and *PMCA3* in bundles of inner hair cells, *PMCA2a* was detected consistently at moderate (inner hair cells) to very high (outer hair cells) levels, suggesting that this isozyme is the principal Ca^{2+} pump of cochlear hair bundles.

The presence of *PMCA1b* in the basolateral membrane of inner hair cells resembled its expression in the bullfrog and rat vestibular system. By contrast, no substantial amounts of *PMCA* were identified in basolateral membranes of outer hair cells (see Fig. 6*A*; Crouch and Schulte, 1995). This latter observation perhaps is not surprising, given the near-crystalline packing of outer hair cell intramembrane particles that are thought to correspond to the motor responsible for electromotility (Forge, 1991). *PMCA3* was also absent from outer hair cells, although it was present in inner hair cells. We consistently observed *PMCA3* in a ring below the apical surface that resembled the pericuticular necklace (Hasson et al., 1997), a region that is richly endowed with intracellular vesicles; in addition, *PMCA3* was observed occasionally in bundles of inner hair cells. A possible interpretation of this localization is that *PMCA3* is stored in the necklace

and is mobilized into hair bundles of inner hair cells under appropriate conditions. Finally, we found no evidence for *PMCA4* in the organ of Corti. Previous *in situ* hybridization experiments suggested that *PMCA4* is expressed in the cochlea at very low levels, with the exception of a short period of expression in inner hair cells that peaked at P12 (Furuta et al., 1998). Because our observations were confined to older rats, our inability to detect this isozyme was unsurprising.

Localization and targeting of PMCA isozymes

Within hair cells and supporting cells *PMCA* isozymes are targeted to the apical domain exclusively or to the basolateral domain exclusively. Although splice variants may allow localization within a domain, these sequences may not be those that dictate basolateral or apical targeting. Conclusive determination of targeting mechanisms for individual *PMCA* isozymes must await closer experimental examination, for example by the expression of *PMCA1bx-PMCA2av* chimeras in hair cells.

Within a domain *PMCA* isozymes did not always appear to be distributed uniformly within the membrane. Localization may be specified by splicing region C alternative exons. Human *PMCA2b* and *PMCA4b* bind via their C termini to several membrane-associated scaffolding proteins that contain PDZ domains (Kim et al., 1998). Bullfrog *PMCA1b* contains the same C-terminal sequence (-ETSL) as human *PMCA2b*, suggesting that it too could bind PDZ proteins, which could localize this isozyme within the hair cell.

PMCA2 sometimes appeared to be concentrated near stereocilia tips (see, for example, Fig. 3*G*) and in a broad band near the base of the hair bundle (see, for example, Fig. 3*E*; Yamoah et al., 1998). Bundle *PMCA* uses the *a* splicing variant, however, which apparently does not bind any PDZ-containing proteins (Kim et al., 1998). Localization of *PMCA2av* within stereocilia might rely instead on the novel *v* variant within splicing region A. The 12 extra amino acids encoded by *v* resemble sequences that bind to src homology 3 (SH3) domains; although matching the minimal sequence, PxxP, required for SH3 binding, *v* does not match the more stringent general consensus $\psi\text{Pxx}\psi\text{P}$, where ψ is a hydrophobic amino acid (Sudol, 1998). Although this putative SH3-binding domain conceivably could target *PMCA2av* to or localize it in bullfrog sacculus hair bundles, its absence from rat vestibular and auditory *PMCA2* suggests that other localization mechanisms may be used in rat and perhaps in bullfrog hair cells.

PMCA2a as the stereocilia calcium pump

We have shown previously that *PMCA* is present in bullfrog hair bundles at the very high membrane density of $\sim 2000/\mu\text{m}^2$ (Yamoah et al., 1998). Furthermore, this *PMCA* can be sufficiently active during transduction to generate a measurable outward current (Yamoah et al., 1998) and extrude most of the Ca^{2+} that enters a stereocilium during transduction (Lumpkin and Hudspeth, 1998). The data presented here show that the calcium pump responsible for this activity is *PMCA2a*.

Why is *PMCA2a* restricted to the hair bundle? The hair cell may target *PMCA2a* to the hair bundle simply so that a high calcium pump concentration can be established there. If all *PMCA2a* travels to the hair bundle, the hair cell can regulate the membrane density of this protein in the stereocilia by controlling *PMCA2a* transcription and translation.

In addition, *PMCA2* may have catalytic or regulatory properties that may be relevant for hair bundle physiology. First, *PMCA2* isoforms have significantly lower values of K_m for Ca^{2+}

than do other isoforms, indicating that they can lower Ca^{2+} to substantially lower levels than can other isozymes (Elwess et al., 1997; Strehler and Zacharias, 2001). Resting hair bundle Ca^{2+} is very low (Denk et al., 1995; Lumpkin and Hudspeth, 1998); bundles may use PMCA to lower Ca^{2+} to insulate them from Ca^{2+} fluxes in the soma (Lenzi and Roberts, 1994) or to permit specific bundle functions, such as tip link regeneration (Zhao et al., 1996). Second, calmodulin activates PMCA2 isozymes with a substantially greater affinity than with other PMCA isozymes; this affinity difference may be irrelevant, however, because of the high hair bundle calmodulin concentration ($\sim 70 \mu\text{M}$; Walker et al., 1993). Finally, the variant used at splicing region C may be relevant for hair bundle function. For example, protein kinase C phosphorylates the *a* form of PMCA2, preventing stimulation of the pump by calmodulin (Enyedi et al., 1997). Only when PMCA2a is expressed and purified can we determine all of the properties that are significant for hair bundle function, including the maximal pumping rate of the pump, H^+ stoichiometry (Hao et al., 1994), and dependence on ATP, Ca^{2+} , and calmodulin.

PMCA may have a broader role within hair bundles other than simply removing Ca^{2+} that enters during transduction or from the soma. Hair cells in lower vertebrates are capable of Ca^{2+} -triggered bundle movements (Benser et al., 1996; Ricci et al., 2000), which may be required for sensitive detection of periodic stimuli (Hudspeth et al., 2000; Manley, 2000). Indeed, a PMCA2a-established spatiotemporal Ca^{2+} gradient at the Ca^{2+} -sensitive site of the channel could provide the energy source for active bundle movements (Hudspeth, 1997). In the auditory system PMCA2a is present in an exceptionally high density in outer hair cells, which are responsible for local mechanical amplification (Dallos, 1992). Amplification, which is required for sensitive stimulus detection and sharp frequency discrimination, may be powered by somatic electromotility (Ashmore and Kolston, 1994) or hair bundle mechanical activity (Hudspeth et al., 2000). By establishing the necessary energy source, PMCA2a would play a central role in any bundle-based mechanism.

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