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Biochemical and Biophysical Research Communications 322 (2004) 1131-1139

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Hippocalcin in the olfactory epithelium: a mediator of second messenger signaling[☆]

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Received 23 July 2004 Available online 10 August 2004

Abstract

Intracellular Ca²⁺ plays an important role in a variety of second messenger cascades. The function of Ca²⁺ is mediated, in part, by Ca²⁺-binding proteins such as calmodulin, calretinin, calbindin, neurocalcin, recoverin, and visinin-like proteins (VILIPs). These proteins are highly expressed in rat olfactory receptor neurons (ORNs) and are localized to distinct intracellular regions. In the present study, we have identified another Ca²⁺-binding protein, hippocalcin, in the rat olfactory epithelium (OE). Olfactory/brain hippocalcin shows high sequence homology with hippocalcins expressed in mice and humans. Hippocalcin was predominantly localized to the olfactory cilia, the site of the initial events of olfactory signal transduction, and was found to regulate the activity of ciliary adenylate cyclases (ACs) and particulate guanylyl cyclases (GCs) in a Ca²⁺-dependent manner. These data indicate that hippocalcin is expressed in rat ORNs, and is likely to regulate second messenger cascades in a Ca²⁺-dependent manner.

Ca²⁺ regulates diverse cellular functions in a variety of cell types, including neurons. In neurons, Ca²⁺-dependent regulation is involved in a number of critical neuronal processes such as excitation, neurotransmitter release, synaptic plasticity, and gene transcription [1–4]. In most cases, Ca²⁺-binding proteins regulate these Ca²⁺-related functions (see a review [5]). Olfaction is an excellent example of a signaling system that utilizes Ca²⁺ in a number of ways (see reviews [6,7]). In response to odorants, ORNs display a rapid and transient increase in intracellular Ca²⁺ levels. Briefly, transduction begins when odorants interact with specific receptors present on the cilia of

* Corresponding author. Fax: +1 410 614 8033. E-mail address: cmoon@jhmi.edu (C. Moon). ORNs. These G-protein-coupled receptors activate ACs to produce cAMP (Fig. 1). As intracellular cAMP levels increase, olfactory cyclic nucleotide-gated channels (oCNCs) open to allow an influx of Na⁺ and Ca²⁺, leading to the generation of an action potential, which transduces signals to the olfactory bulb. Later, Ca²⁺ is pumped out of ORNs by Na⁺/Ca²⁺ exchangers and Ca²⁺-ATPases present in the cilia and dendritic knobs, thus maintaining Ca²⁺ homeostasis and returning the cell to electrical neutrality [8–10] (Fig. 1). Other second messengers, such as cyclic GMP, are also produced by odorant stimulation [11–13].

Olfactory signal transduction pathways are modulated by changes in intracellular Ca²⁺ concentrations (see reviews [6,7]), and specific Ca²⁺-binding proteins mediate these changes. Thus, identification and characterization of the Ca²⁺-binding proteins within the olfactory system can provide important information about the complex signal transduction involved in olfaction. To date, olfactory tissues have been reported to contain

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^{**} Abbreviations: AC, adenylate cyclase; GC, guanylyl cyclase; GCAP, guanylyl cyclase-activating protein; NCS, neuronal Ca²⁺ sensor; OE, olfactory epithelium; ORN, olfactory receptor neuron; PKG, cGMP-dependent protein kinase; VILIP, visinin-like protein.

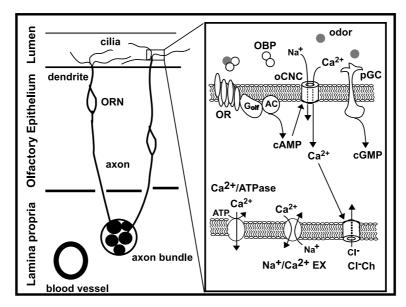


Fig. 1. Olfactory signal transduction. Signaling cascades that mediate the initial phase of odorant detection, desensitization, and return to Ca^{2+} homeostasis. See the text for details. *Abbreviations:* AC, adenylate cyclase; Cl^- Ch, chloride channel; G_{olf} , olfactory G protein; OBP, odorant-binding protein; oCNC, olfactory cyclic nucleotide-gated channel; OR, odorant receptor; ORN, odorant receptor neuron; and pGC, particulate guanylyl cyclase.

a variety of Ca²⁺-binding proteins, including calmodulin [14], calretinin [15], p26olf [16], calbindin-D28k [15], neurocalcin [17,18], recoverin [18], VILIP [19], and guanylyl cyclase-activating protein 1 (GCAP1) [13]. We have identified another Ca²⁺-binding protein, hippocalcin, in rat OE.

Hippocalcin is a Ca²⁺-binding protein of the neuronal Ca²⁺ sensor protein (NCS) family. It was first thought to be expressed exclusively in the hippocampus [20], but later was found in other brain regions, including the neocortex, caudate-putamen, taenia tecti, claustrum, olfactory tubercle, anterior olfactory nucleus, and the granule cell and glomerular layers of the olfactory bulb [21]. It has a primary structure containing three putative Ca²⁺-binding sites (EF-hands) and a N-terminal myristoylation site. Several studies have revealed that myristoylation is necessary for its Ca²⁺-dependent membrane association [20,22,23]. The N-terminal myristoyl moiety on hippocalcin interacts with lipid bilayers and facilitates interaction with other membrane proteins [20]. Hippocalcin is also expressed in *Drosoph*ila melanogaster and is found primarily in the brain [24]. The fact that hippocalcin is expressed throughout the central nervous system of adult flies, particularly in the neuropil, suggests that hippocalcin is perhaps involved in synaptogenesis. Similar implications are suggested by studies of post-natal developmental expression of hippocalcin in rat retina. These studies demonstrate that compared to other Ca²⁺ sensor proteins, hippocalcin is expressed later in development at a time when synaptogenesis is taking place [25]. Here, we demonstrate another important role for hippocalcin in ORNs, modulation of second messenger signaling.

Materials and methods

Antibody. Rabbit anti-hippocalcin antibodies were a gift from Dr. M. Kobayashi (Toho University School of Medicine, Tokyo, Japan). Calculation of the phylogenetic tree. The phylogenetic tree was constructed from aligned sequences using the ClustalW program by the neighbor-joining method [26] using the Neighbor program in the

PHYLIP package (Phylogeny Inference Package version 3.6a, 2002, distributed by Dr. J. Felsenstein, Department of Genetics, University of Washington, Seattle, WA). Gaps were removed from the calculations. The evolutionary distances of the sequences were estimated based on Dayhoff's PAM matrix [27] using the program of Protdist of the PHYLIP package (version 3.6a). Bootstrap re-samplings were performed in the Seqboot program of the PHYLIP package (version 3.6a). Accession numbers of amino acid sequences are indicated in the figure legend.

Cilia preparation. Cilia were prepared as previously published [13,28]. Briefly, male Sprague–Dawley rat nasal turbinates were dissected, pooled, and washed in EDTA (2mM) in Ringer's solution (112mM NaCl, 3.4mM KCl, 2.4mM NaHCO₃, and 2mM Hepes, pH 7.4) at 4°C. The tissue was centrifuged at 5000g for 5min, and the pellet was re-suspended in deciliation buffer. The bathing medium was supplemented with CaCl₂ to a final concentration of 10mM and agitated gently end-over-end for 20min at 4°C. The deciliated epithelium was removed by centrifugation for 5min at 1500g. The supernatant containing isolated cilia was centrifuged for 10min at 12,000g and the resulting pellet containing the isolated cilia was washed twice in 10mM Tris–HCl, 3mM MgCl₂, and 1mM EDTA, pH 8.0. The final cilia pellet was re-suspended in 20 µl of 10 mM Tris–HCl, 3 mM MgCl₂, and 1 mM EDTA, pH 8.0, and stored at -80°C before use.

Cyclase assay. Cyclase assays were performed as previously described [29] with some modifications. The reaction was initiated by adding olfactory cilia (5µg) and purified hippocalcin (40µM) to the stimulation solution containing (final concentration): 1.3 mM [γ -³²P]ATP (19,000–22,000 dpm/nmol; New England Nuclear), 50 mM Hepes (pH 7.8), 60 mM KCl, 20 mM NaCl, 10 mM MgCl₂, 0.4 mM EGTA, an appropriate concentration of CaCl₂ (0.1–10 µM free Ca²⁺), and 1 mM 3-isobutyl-1-methyl-xanthine (IBMX). Free Ca²⁺ was calculated using the MaxChelator program (www.stanford.edu/~cpatton/webmaxc2.htm). The reaction was terminated after 2 min stimulation.

cAMP formation was measured with a scintillation counter [30–45]. For guanylyl cyclase (GC) assays, $1.3\,\text{mM}$ [α - 32 P]GTP (19,000–22,000 dpm/nmol) was used and cGMP [30] formation was measured after 10 min stimulation.

Expression and purification of hippocalcin. The entire coding region of hippocalcin was sub-cloned into pET17b (Novagen, Madison, WI, USA) using *Hin*dIII and *Bam*HI sites, and expressed in BL21 (DE3) co-transfected with a pBB131 construct containing the sequence for *N*-myristoyl transferase. Bacterially expressed hippocalcin was purified according to the instructions for insoluble proteins (T7 Tag Purification Kit; Novagen).

Immunohistochemical analysis. Immunohistochemistry was performed following the Vectastain ELITE protocol (Vector Laboratories). Tissue sections were permeabilized for 1 h in PBS containing 0.1% Triton X-100 and blocked for 1 h in PBS containing 1% BSA and 4% normal serum. Section slides were then incubated overnight at 4°C in PBS containing the rabbit anti-hippocalcin antibody at a dilution of 1:500. The next day, slides were rinsed with PBS and incubated with Vectastain (Vector Laboratories) biotinylated secondary antibody (1:1000) for 30 min, and then incubated in 0.5% H₂O₂ for 10 min at room temperature. Slides were then incubated in the avidin–biotin reagent for 30 min, rinsed in PBS, and developed using 0.25 mg/ml diaminobenzidine (Sigma) in 50 mM Tris, pH 7.4. For pre-absorption of anti-hippocalcin antibody, the rabbit anti-hippocalcin antibody was incubated with purified hippocalcin (1.5 μg) on an orbit rotator overnight at 4°C and used as a primary antibody.

Gel electrophoresis and immunoblot analysis. Whole rat adult nasal epithelium and tissues were homogenized in RIPA buffer (50 mM Tris, pH 7.4, 150 mM NaCl, 1% (v/v) Triton X-100, 1% (v/v) deoxycholic acid, and 0.1% SDS) and incubated on ice for 30 min. The extracts were cleared by centrifugation for 15 min in an Eppendorf microfuge at 13,000 rpm at 4°C. Supernatants (40 μg of protein per gel lane) were subjected to SDS/PAGE on a 4–15% gradient gel. Protein content was measured using Bradford assay with BSA as a standard. The separated proteins were transferred to PVDF membrane and the membranes were probed with rabbit anti-hippocalcin antibody. Horseradish peroxidase-conjugated goat anti-rabbit IgG (Roche Molecular Biochemicals) was used as a secondary antibody at 1:5000 dilutions. Bands were visualized by enhanced chemiluminescence (Amersham–Pharmacia).

Isolation of the hippocalcin cDNA clone. Total RNA was purified from OE tissues by the acid guanidinium thiocyanate phenol/chloroform extraction method [46]. For cDNA synthesis, polyadenylated RNA was further isolated using mRNA purification kit (Invitrogen). An oligo(dT)-primed rat OE cDNA library was constructed in Uni-ZAP XR (Stratagene) from poly(+) RNA isolated from rat OE tissues by using a ZAP-cDNA synthesis kit (Stratagene).

Degenerate oligonucleotide primers were designed based on conserved sequences of GCAP Ca²⁺-binding protein family members from various species. Hippocalcin was isolated by reverse transcriptase-polymerase chain reaction from the rat OE cDNA library using sense primer (5'-CATATGGGCAAGCAGAATAGCAAGC-3') and antisense primers (5'-AAGCTTCAGAACTGGGAAGCGCT-3'). Multiple DNA fragments were acquired through PCR and cloned into pSTBlue-1 (Novagen, Madison, WI, USA) using the Perfect Blunt Cloning Kit. Sequence analysis was used to identify full-length clones that were then sub-cloned into a pET17B expression vector.

Phosphorylation and dephosphorylation of cilia fractions. Phosphorylation reactions were performed according to the method of Moon et al. [13], with modifications. Protein kinases and phosphatases were obtained from Promega (Madison, WI). Isolated cilia fractions (200 μg) were prepared in a reaction buffer (200 μl) containing 50 mM Tris–HCl, pH 7.5, 10 mM MgCl₂, 0.4 mM EGTA, 10 μg/ml calmodulin, 1 mM LiCl₂, 1 mM 8-Br-cGMP, and 1 mM ATP.

For phosphorylation of cilia fractions, cAMP-dependent protein kinase catalytic subunit (120 U), cGMP-dependent protein kinase (300 U), and Ca²⁺/calmodulin PK (CaMK, 1 U) were added and

incubated for 30 min at room temperature. The cilia fractions were centrifuged for 2 min at 12,000g. The resulting pellet of cilia fraction was re-suspended in cyclase assay solution and assayed for AC activity

For dephosphorylation of cilia fractions, protein phosphatases 2A and 2B were used. Cilia fractions were incubated with protein phosphatase 2A (2.5 U) and protein phosphatase 2B (0.5 U) for 30 min at room temperature. Then, the cilia fractions were centrifuged, re-suspended in cyclase assay solution, and assayed for AC activity.

Control cilia fractions were kept for 30 min at room temperature and assayed for AC activity.

Statistical analysis. Apparent cyclase activities under various free Ca²⁺ concentrations and dose–responses of hippocalcin protein were analyzed statistically, using one-way ANOVA. Post testing was performed comparing all values to control (Dunnett method). Statistical significance is shown in each figure.

Results and discussion

Hippocalcin was identified in a rat OE cDNA library generated using poly(A)+ RNA isolated from rat OE. The complete rat hippocalcin sequence (GenBank AY442172) contained an open reading frame of 579 bp (193 amino acids). The rat hippocalcin showed high amino acid sequence homology to mouse and human hippocalcins from brains (100% and 99%, respectively). Sequence analysis of hippocalcin compared to VILIP-1, VILIP-2, and VILIP-3 showed that hippocalcin had the highest sequence homology to VILIP-3 (95%; Fig. 2A), but showed less homology to other VILIPs (66%; Fig. 2A).

Phylogenetic analysis of Ca²⁺-binding proteins suggests that the ancestors of NCSs were first duplicated into the ancestors of the GCAP subfamily and the other NCSs (Fig. 2B). High clustering probabilities of NCSs (100%) and GCAP1s (99.7%) substantiate this possibility. The ancestors of the other neuronal Ca²⁺ sensors diverged to form recoverin, VILIP-1, VILIP-2, VILIP-3, hippocalcin, and neurocalcin subfamilies. Hippocalcin is phylogenetically similar to members of the VILIP family, which are found in various regions of brains and known to be involved in cellular signal transduction [47]. Olfactory hippocalcin is 100% identical to the brain hippocalcin. Another OE Ca²⁺-binding protein, p26olf, is more closely related to calbindin/calretinin than to NCSs.

To investigate the role of hippocalcin in olfactory signal transduction, we determined the intracellular localization of hippocalcin in adult rat OE by immunohistochemical analysis. Hippocalcin immunoreactivity was predominantly localized to the apical layer of the OE, where the olfactory knob and cilia of mature ORNs reside (Fig. 3A), suggesting that hippocalcin may be involved in initial olfactory signal transduction or adaptation. Signal was not detectable in soma or axons of ORNs. When the antibodies were pre-absorbed with purified hippocalcin proteins, the immunoreactivity of hippocalcin in the ciliary layer was greatly reduced (Fig. 3B, right panel). Immunoblot analysis showed that hippocalcin is also

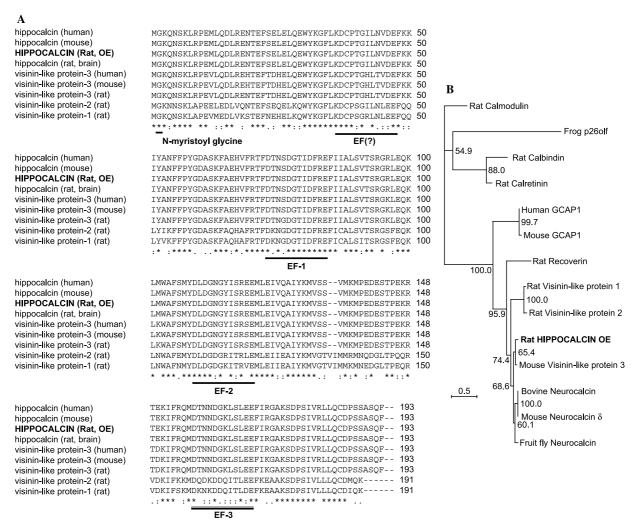


Fig. 2. (A) Multiple sequence alignment of rat, mouse, and human hippocalcins and VILIPs. The alignment was performed using the ClustalW program (http://www.ebi.ac.uk/clustalw/). (*) indicates that the residues in that column are identical in all sequences in the alignment. (:) and (.) indicate conserved substitutions and semi-conserved substitutions of the residues in those columns, respectively. N-myristoyl glycine, three Ca²⁺-binding motifs, and a probable Ca²⁺-binding motif are indicated; see text for details. The accession numbers for the amino acid sequences are: human hippocalcin (*Homo sapiens*), NP_002134; mouse hippocalcin (*Mus musculus*), NP_034601; rat hippocalcin (*Rattus norvegicus*), rat brain hippocalcin (R. norvegicus) [68,69], AAR14053; human VILIP-3 (H. sapiens), NP_002140; mouse VILIP-3 (M. musculus), P35333; rat VILIP-3 (R. norvegicus), and NP_059052; rat VILIP-2 (R. norvegicus), NP_059053; and rat VILIP-1 (R. norvegicus), NP_036818. (B) Phylogenetic analysis of Ca²⁺-binding proteins expressed in the olfactory system, calculated from amino acid sequences. Numbers at the nodes indicate clustering percentage obtained from 1000 bootstrap re-samplings. Sequence data used in the present analyses were taken from GenBank, EMBL, SWISS-PROT, and NCBI databases. The accession numbers for the amino acid sequences are: rat calretinin (R. norvegicus), NP_446440; rat calmodulin (R. norvegicus), NP_114175; rat calbindin (R. norvegicus), NP_114190; fruit fly neurocalcin (Drosophila melanogaster), A55666; rat VILIP 2 (R. norvegicus), NP_059053; bovine neurocalcin (Bos taurus), BAA01706; frog p26olf (Rana catesbeiana), BAA34388; rat recoverin (R. norvegicus), NP_543177; mouse VILIP 3 (M. musculus), P35333; rat VILIP 1 (R. norvegicus), NP_036818; rat hippocalcin olf (R. norvegicus), AAR14053; human GCAP1 (H. sapiens), P43080; mouse GCAP1 (M. musculus), P43081; and mouse neurocalcin olf (M. musculus), NP_598855.

expressed in the olfactory bulb though at lower levels than in the OE (Fig. 3C), indicating additional roles for hippocalcin in olfactory processing.

Olfactory tissues express a variety of Ca²⁺-binding proteins in distinct subcellular locations. Calmodulin is expressed in olfactory cilia, cytoplasm, and axon fibers [18,48]. Calretinin is expressed in rat apical dendrites and axon fibers of ORNs [18]; p26olf in frog olfactory cilia [16]; calbindin-D28k in external fiber bundles of ORNs [15]; neurocalcin in the cell body, cytoplasm, and axon fibers of ORNs [17,18,49]; VILIP in rat

olfactory cilia and dendritic knobs [19]; and GCAP1 in rat olfactory cilia [13]. The cellular localization of these Ca²⁺-binding proteins may be closely related to their functions in the ORNs. Thus, our immunohistochemistry data suggested that hippocalcin may be involved in initial olfactory signal transduction or adaptation, as proposed for other Ca²⁺-binding proteins localized to the olfactory cilia.

We tested whether hippocalcin could affect second messenger signaling in olfactory cilia. Since N-terminal myristoylation is necessary for the interaction of

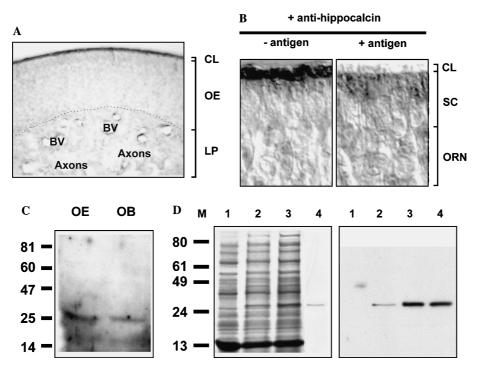


Fig. 3. Expression of hippocalcin proteins in the OE. (A) Immunohistochemical analysis of hippocalcin in the rat OE. Axons, BV, CL, OE, and LP represent axon bundle, blood vessel, cilia layer, OE, and lamina propria, respectively. (B) Pre-absorption of the antibody against hippocalcin. Immunoreactivity was blocked when the antibodies were pre-absorbed with purified hippocalcin protein (1.5µg) (right panel). CL, SC, and ORN represent ciliary layer, supporting cell layer, and ORN layer, respectively. (C) Immunoblot analysis in the OE and OB. Protein sizes are represented on the left side. OB and OE represent the olfactory bulb and the OE, respectively. (D) Purification of hippocalcin protein. Bacterially expressed hippocalcin was purified using a T7 Tag Purification Kit (Novagen) (left panel). 1, 2, 3, and 4 indicate un-induced samples, insoluble fractions, and purified hippocalcin proteins, respectively. Immunoblot analysis confirmed expression and purification of hippocalcin proteins (right panel). Protein sizes (M) are represented on the left side in kiloDaltons.

hippocalcin with other membrane proteins [20], we expressed both hippocalcin and N-myristoyl transferase in bacteria. After over-expression in bacteria, hippocalcin was purified to apparent homogeneity (Fig. 3D). Using purified hippocalcin and isolated olfactory cilia, we examined the effects of hippocalcin on cyclase activities. We first determined the dose–response of AC activity to hippocalcin at a free Ca²⁺ concentration of 10 nM (Fig. 4A). At this free Ca²⁺ concentration, AC activity was increased as the concentration of hippocalcin protein was raised. Under these conditions, the maximum effect of hippocalcin was achieved at 40 µM. Next, we examined the effect of hippocalcin on AC activity at various free Ca²⁺ concentrations. Incubation with hippocalcin modulated ciliary AC activity in a Ca²⁺-dependent manner. Specifically, hippocalcin increased AC activity at low free Ca²⁺ concentration and this activity decreased as free Ca²⁺ concentrations were increased (Fig. 4B). The normal intracellular Ca²⁺ concentration in the cilium is 40 nM at rest and increases up to 300 nM upon odorant stimulation [50]. Hippocalcin increased AC activity significantly at concentrations as low as 1 nM (Fig. 4B). When free Ca²⁺ concentrations were raised above 300 nM, which mimics the intracellular Ca2+ concentration upon odorant stimulation, hippocalcin no longer had an effect on

AC activity. Thus, hippocalcin mediated activity of ACs at physiological Ca²⁺ concentrations. An opposite effect has been reported for VILIP-1; in vitro recombinant VI-LIP-1 attenuated odorant-induced AC activation in a Ca²⁺-dependent manner [19]. Thus, VILIP-1 and hippocalcin may fine-tune the AC activity in the adaptation of ORNs to odorant stimulation.

Numerous reports have demonstrated that basal levels of phosphorylation are essential to maintain protein function [51-53]. Thus, we determined whether the phosphorylation level of ciliary proteins affected the modulation of AC by hippocalcin. At low free Ca²⁺ levels, protein kinase treatment inhibited the effect of hippocalcin on AC activity, whereas either native cilia or protein phosphatase-treated cilia showed increases in hippocalcin-mediated AC activity (Fig. 4C, left panel). At the high free Ca²⁺ conditions, only protein phosphatase-treated cilia showed significant increases in hippocalcin-mediated AC activity (Fig. 4C, left panel). Compared to native cilia, protein phosphatase treatment also increased the basal level of AC activity significantly, whereas protein kinase treatment significantly decreased it (Fig. 4C, left panel). Our data suggest that the dynamic status of phosphorylation of ciliary ACs may play a critical role in hippocalcin dependency; i.e.,

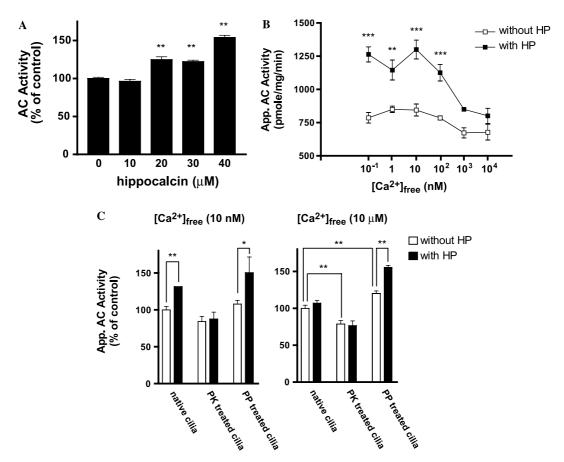


Fig. 4. Effects of hippocalcin on AC activity. (A) Dose–response of hippocalcin on AC activities in rat olfactory cilia. Apparent AC activities are determined at the free Ca^{2+} concentration of 10nM. (B) Effects of hippocalcin on AC activity in rat olfactory cilia. Rat olfactory cilia were isolated and assayed for AC activity. Apparent AC activities are determined under various Ca^{2+} concentrations with and without hippocalcin. (C) Effects of protein kinases and protein phosphatases on hippocalcin-mediated AC activity in rat olfactory cilia. Rat olfactory cilia were treated with protein kinases (or protein phosphatases) and assayed for AC activity. Apparent AC activities are determined under two free Ca^{2+} concentrations with and without hippocalcin. Statistical significances are indicated (one-way ANOVA, *p < 0.05, **p < 0.01).

hyper-phosphorylation appears to desensitize the hippocalcin-mediated AC activity. Specifically, Ca²⁺ influx by odorant stimulation induces phosphorylation of AC3 that in turn causes AC3 desensitization [54,55]. Our data also showed that there is some degree of basal phosphorylation of AC3 [55]. AC3 is the most prominent isoform of AC found in olfactory cilia [56–60], and phosphorylation-dependent regulation of AC by hippocalcin can be one of the mechanisms for odor adaptation. Other ACs, such as AC2 or AC4, are also present in the OE [61], and thus represent other potential targets for regulation by hippocalcin.

VILIPs are reported to modulate intracellular cGMP level [62,63]. We tested whether hippocalcin affected the activity of GCs in cilia. First, we determined the doseresponse of GC to various concentrations of hippocalcin at a free Ca²⁺ concentration of 10 nM (Fig. 5A). GC activity was significantly reduced as the concentration of hippocalcin protein increased. Next, we examined the effect of hippocalcin on GC activity at various free Ca²⁺ concentrations. Hippocalcin significantly inhibited

the activity of particulate GCs over a limited range of free Ca²⁺ concentrations, specifically in the range of 1– 10 nM (Fig. 4C). The inhibitory effect of hippocalcin on GC activity decreased as intracellular Ca2+ concentrations increased. These results suggest that in the olfactory system, hippocalcin may play a role in keeping the level of cGMP low in ORNs at rest, lowering the activity of cGMP-dependent protein kinases (PKGs). PKGs are reported to inhibit AC activity in olfactory cilia [13]. Hippocalcin did not affect the activity of GCs at higher Ca²⁺ concentrations, suggesting that it is not involved in regulation of particulate GCs in the cilia upon odorant stimulation and subsequent increases in intracellular Ca²⁺. GCAP1 has been proposed to function in that role [13]. VILIP-1 can also stimulate a type of particulate GC, GC-B [63], which is abundantly expressed in the OE [64].

Hippocalcin has been also reported to perform other functions, such as inhibiting G-protein-linked receptor kinase 1 (GRK1) in a Ca²⁺-dependent manner [65] and protecting neurons against Ca²⁺-induced cell death

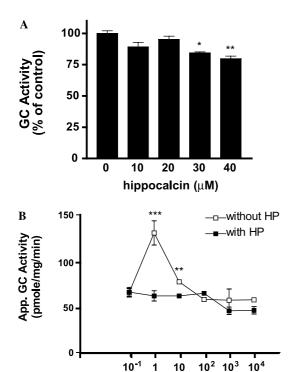


Fig. 5. Effects of hippocalcin on GC activity. (A) Dose–response of hippocalcin on GC activities in rat olfactory cilia. Apparent GC activities are determined at the free Ca^{2+} concentration of 10 nM. (B) Effects of hippocalcin on GC activity in rat olfactory cilia. Apparent GC activities were determined under various Ca^{2+} concentrations with and without hippocalcin. Statistical significances are indicated (oneway ANOVA, *p < 0.05, **p < 0.01, and ***p < 0.001).

[Ca²⁺]_{free} (nM)

by interacting with neuronal apoptosis inhibitory protein [66]. In cerebellar Purkinje cells in the brain, VI-LIP-3, which is the Ca²⁺ sensor displaying the highest sequence homology to hippocalcin, has been implicated in complex brain functions such as synaptic plasticity [67]. Thus, hippocalcin may play multiple roles in ORNs in addition to modulating second messenger signaling.

In summary, multiple Ca²⁺-binding proteins are expressed in ORNs. One of their roles may be the fine-tuning of the response to odorant stimulation and adaptation as changes in intracellular Ca²⁺ level occur. Here, we have identified the Ca²⁺-binding protein hippocalcin in cilia, and have demonstrated its ability to modulate ACs and GCs, which are involved in stimulus detection and adaptation, respectively. Hippocalcin may play a similar dual role in other systems, modulating signal transduction in response to fluctuations in intracellular Ca²⁺ concentration.

Acknowledgments

The authors wish to thank Drs. M. Kobayashi for his kind gift of hippocalcin antibody and Andrei Alekseev for pBB131 and his technical advice. The NINDS

(GVR) and NIDCD (G.V.R., J.P.S., and C.M.) and Howard Hughes Medical Institute (A.M.) supported this work. This research was also supported by NIH Grant EY09339, a grant from Research to Prevent Blindness, Inc. (RPB) to the Departments of Ophthalmology at the University of Washington, and a grant from the E.K. Bishop Foundation. K.P. is the RPB Senior Investigator.

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