

Homer2 Protein Regulates Plasma Membrane Ca^{2+} -ATPase-mediated Ca^{2+} Signaling in Mouse Parotid Gland Acinar Cells*

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Background: Homer proteins bind multiple Ca^{2+} -signaling proteins to shape the Ca^{2+} signal by poorly understood mechanisms.

Results: Homer2 regulates PMCA expression and activity in parotid acinar cells.

Conclusion: Homer2 acts as a regulator of PMCA-mediated Ca^{2+} clearance.

Significance: Inhibition of Ca^{2+} clearance by Homer2 further clarifies its role in Ca^{2+} signaling.

Homer proteins are scaffold molecules with a domain structure consisting of an N-terminal Ena/VASP homology 1 protein-binding domain and a C-terminal leucine zipper/coiled-coil domain. The Ena/VASP homology 1 domain recognizes proline-rich motifs and binds multiple Ca^{2+} -signaling proteins, including G protein-coupled receptors, inositol 1,4,5-triphosphate receptors, ryanodine receptors, and transient receptor potential channels. However, their role in Ca^{2+} signaling in nonexcitable cells is not well understood. In this study, we investigated the role of Homer2 on Ca^{2+} signaling in parotid gland acinar cells using *Homer2*-deficient (*Homer2*^{-/-}) mice. Homer2 is localized at the apical pole in acinar cells. Deletion of *Homer2* did not affect inositol 1,4,5-triphosphate receptor localization or channel activity and did not affect the expression and activity of sarco/endoplasmic reticulum Ca^{2+} -ATPase pumps. In contrast, *Homer2* deletion markedly increased expression of plasma membrane Ca^{2+} -ATPase (PMCA) pumps, in particular PMCA4, at the apical pole. Accordingly, *Homer2* deficiency increased Ca^{2+} extrusion by acinar cells. These findings were supported by co-immunoprecipitation of Homer2 and PMCA in wild-type parotid cells and transfected human embryonic kidney 293 (HEK293) cells. We identified a Homer-binding PPXXF-like motif in the N terminus of PMCA that is required for interaction with Homer2. Mutation of the PPXXF-like motif did not affect the interaction of PMCA with Homer1 but inhibited its interaction with Homer2 and increased Ca^{2+} clearance by PMCA. These findings reveal an important regulation of PMCA by Homer2 that has a central role on PMCA-mediated Ca^{2+} signaling in parotid acinar cells.

Ca^{2+} is a common second messenger with roles in fertilization, muscle contraction, neurotransmitter release, exocytosis, learning, and memory. In addition, it regulates critical functions of polarized secretory gland cells (1, 2). Secretory cell Ca^{2+} signaling is evoked primarily by activation of G protein-coupled receptors (GPCRs),² which are coupled to G_q or G_i . These cells are an excellent model system to study Ca^{2+} signaling and the role of scaffolding proteins in the function of various components of the Ca^{2+} signal, including plasma membrane Ca^{2+} -ATPase (PMCA) and sarco/endoplasmic reticulum Ca^{2+} -ATPase (SERCA) pumps, the inositol 1,4,5-triphosphate (IP_3) receptors (IP_3Rs), Ca^{2+} release channels in the endoplasmic reticulum (ER), and Ca^{2+} influx channels in the plasma membrane (3).

Several GPCRs possess binding motifs that associate with members of the Homer family of scaffolding proteins. The Homer family, which includes Homer1, Homer2, and Homer3, as well as several splice variants, was discovered when *homer1a* was cloned as an immediate early gene product expressed in neurons upon prolonged stimulation (4–7). With the exception of Homer1a, all Homer proteins (*i.e.* Homer1b/c, 2, and 3; referred to as “long Homers”) are expressed constitutively throughout the central nervous system (CNS). These proteins are composed of an N-terminal Ena/VASP homology 1 protein-binding domain and a C terminus that folds into a coiled-coil multimerization domain and leucine zipper (7, 8). The Ena/VASP homology 1 domain is a protein-protein binding module that recognizes the proline-rich motifs PPXXF, PPXF, and LPSSP (9–12) and binds to several GPCRs, canonical transient receptor potential channels, IP_3Rs , ryanodine receptors, and the Shank family of scaffolding proteins (10, 12–16). Homer proteins play a central role in Ca^{2+} signaling via regulation of

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² The abbreviations used are: GPCR, G protein-coupled receptor; SERCA, sarco/endoplasmic reticulum Ca^{2+} -ATPase; PMCA, plasma membrane Ca^{2+} -ATPase; IP_3R , inositol 1,4,5-triphosphate receptors; IP_3 , inositol 1,4,5-trisphosphate; ER, endoplasmic reticulum; SLO, streptolysin O; IP, immunoprecipitation; CPA, cyclopiazonic acid.

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neuronal transcription activity and thereby regulate dendritic spine morphogenesis, synapse remodeling, and synaptic clustering of CNS neurons (9, 17–19). Previous research found that Homer1 regulates Ca^{2+} influx by associating IP_3 R_s with transient receptor potential channels (12). Homer2 tunes GPCR stimulus intensity by regulating the regulator of G protein signaling proteins and phospholipase $C\beta$ -promoting guanosine triphosphatase by $G\alpha$ in pancreatic acinar cells. Moreover, Homer2 and Homer3 bind nuclear factor of activated T cells by competing with calcineurin in T lymphocytes (14, 20). However, the role of Homer proteins in Ca^{2+} signaling in nonexcitable cells remains poorly characterized.

In this study, the role of Homer2 on Ca^{2+} signaling in parotid gland acinar cells was investigated using *Homer2* knock-out (*Homer2*^{-/-}) mice. We report that Homer2 interacts with PMCA (particularly PMCA isoform 4) in model systems and *in vivo* and that this interaction regulates PMCA activity. The findings suggest a mechanism by which Homer proteins can regulate PMCA expression and PMCA-mediated Ca^{2+} efflux in parotid acinar cells. We suggest that by inhibiting transient receptor potential channel-mediated Ca^{2+} influx (12) and by differentially modulating Ca^{2+} extrusion by PMCA (present data), the Homers serve to protect the cells from Ca^{2+} toxicity by facilitating cytosolic Ca^{2+} clearance to limit the Ca^{2+} signal duration.

EXPERIMENTAL PROCEDURES

Antibodies and DNA Constructs—Anti-Homer1a (M-13), anti-Homer1 (D-3), anti-Homer2, and anti-PMCA (5F10) antibodies were obtained from Santa Cruz Biotechnology, Inc. (Santa Cruz, CA). Anti- IP_3 R3 antibodies were purchased from BD Transduction Laboratories (San Jose, CA). Anti-PMCA4 (JA9) antibodies were obtained from Affinity BioReagents (Golden, CO). Anti-FLAG M2 antibodies were obtained from Sigma. PMM2-hPMCA1b and PMM2-hPMCA4b were generous gifts from Dr. Emanuel E. Strehler (Mayo Clinic College of Medicine, Rochester, MN). pRK5-HA-Homer1a, pRK5-HA-Homer1c, and pRK5-HA-Homer2 were generously provided by Dr. Paul Worley (The Johns Hopkins University School of Medicine, Baltimore, MD).

Animals and Preparation of Parotid Acinar Cells—Wild-type (WT) and *Homer2*^{-/-} mice have been described previously (14). The life span of *Homer2*^{-/-} mice is similar to that of WT littermates. All animal protocols were performed according to institution guidelines. Mice were sacrificed by cervical dislocation. The cells were prepared from the parotids of WT and *Homer2*^{-/-} mice by limited collagenase digestion as described previously (21). After isolation, the acinar cells were resuspended in an extracellular physiologic salt solution, the composition of which was as follows (in mM): 140 NaCl, 5 KCl, 1 MgCl₂, 1 CaCl₂, 10 HEPES, and 10 glucose, adjusted to pH 7.4 and 310 mosm.

Cell Culture and DNA Transfection—HEK293 cells (Korean Cell Line Bank, South Korea) were maintained in Dulbecco's modified Eagle's medium (Invitrogen) containing 10% fetal bovine serum (Invitrogen) and 100 units/ml penicillin and streptomycin in a 5% CO₂ incubator. Approximately 1–5 × 10⁵ cells were seeded onto coverslips in 60-mm dishes and then

incubated in antibiotic-free medium. The next day, DNA was mixed with Lipofectamine reagent (Invitrogen) and Opti-MEM, incubated for 20 min at room temperature, and then added to the cell culture media. The transfected cells were assayed at 38–48 h after post-transfection.

Measurement of Intracellular Ca^{2+} Concentration ($[Ca^{2+}]_i$)—Parotid acinar cells from WT and *Homer2*^{-/-} mice were loaded with 5 μM Fura-2/AM (Teflabs Inc., Austin, TX) and 0.05% pluronic F-127 for 60 min in physiologic salt solution. Fura-2 fluorescence was measured at the appropriate excitation wavelengths (340/380 nm) and emission at 510 nm wavelengths (ratio = F_{340}/F_{380}) using a Molecular Devices (Downingtown, PA) imaging system. The emitted fluorescence was monitored with a charge-coupled device camera (Photometrics, Tucson, AZ) attached to an inverted microscope. Fluorescence images were obtained at 2-s intervals. All data were analyzed using MetaFluor software (Molecular Devices).

Immunocytochemistry—The immunostaining procedure was described previously (14). In brief, parotid acinar cells from WT and *Homer2*^{-/-} mice were stained with antibodies against Homer2 (1:50); IP_3 R1, -2, or -3 (1:100); SERCA2, PMCA, or PMCA4 (1:200). Staining was detected using goat anti-rabbit or anti-mouse IgG conjugated to fluorescein or rhodamine. Images were collected with confocal LSM 510 and LSM 710 laser scanning microscopes (Zeiss, Göttingen, Germany).

Western Blotting—Protein extracts were prepared from parotid, submandibular gland, and pancreatic acini from WT and *Homer2*^{-/-} mice. Acinar cells were lysed in a buffer containing 150 mM NaCl, 10 mM Tris (pH 7.8 with HCl), 1 mM EDTA, 1% Nonidet P-40, 0.1% SDS, and protease inhibitors (*i.e.* 2 mM Na₃VO₄, 10 mM NaF, 10 μg/ml aprotinin, 10 μg/ml leupeptin, and 10 μg/ml PMSF). The extracts were spun at 13,000 × g for 15 min, and protein concentration was measured using a Bio-Rad protein assay. The samples (50 μg of protein/well) were separated to 6–8% SDS-polyacrylamide gel. Proteins were transferred to a nitrocellulose membrane, blocked with 5% skimmed milk, and incubated overnight with the specific primary antibody, an anti-SERCA2b (1:500), anti-PMCA4 (JA9, 1:500), and anti-PMCA (5F10, 1:500), at 4 °C. The blots were exposed to horseradish peroxidase-conjugated secondary antibodies for 1 h and detected by chemiluminescence (Amersham Biosciences).

Immunoprecipitation—The immunoprecipitation procedure was modified from Shin *et al.* (22) and Kim *et al.* (23). In brief, parotid and HEK293 microsomes were prepared by homogenizing a minced parotid and harvested HEK293 in a buffer containing 20 mM Mops (pH 6.7 with KOH), 250 mM sucrose, 1 mM EDTA, 1 mM MgCl₂, 10 mM benzamidine, and 0.2 mM PMSF. The homogenized samples were centrifuged at 400 × g for 10 min. The supernatants were collected and centrifuged at 900 × g for 10 min at 4 °C. To avoid protein degradation by digestive enzymes, immunoprecipitation was initiated immediately after completion of microsomal preparation. Microsomes were lysed in a buffer containing 50 mM Tris (pH 6.8 with HCl), 150 mM NaCl, 3 mM EDTA, 2 mM EGTA, and 0.5% Triton X-100 supplemented with protease inhibitors. The lysates were cleared by centrifugation at 14,000 × g for 15 min. About 150 μl of the extract (300 μg of protein/sample) was

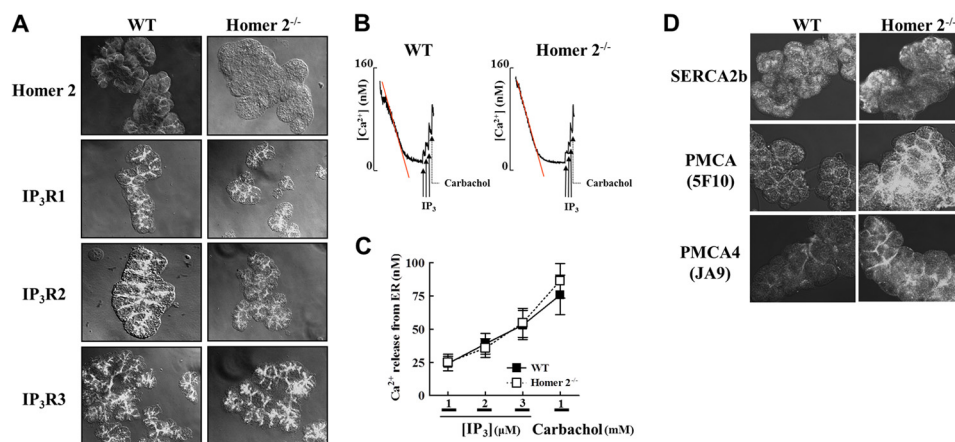


FIGURE 1. Localization of Homer2, IP₃Rs, Ca²⁺ pumps, and IP₃-mediated Ca²⁺ release in WT and *Homer2*^{-/-} cells. *A*, parotid acini from WT and *Homer2*^{-/-} mice were stained for Homer2, IP₃R1, IP₃R2, and IP₃R3. Deletion of *Homer2* did not affect the expression and localization of the IP₃Rs. *B*, cells from WT and *Homer2*^{-/-} mice were permeabilized with SLO and allowed to reduce [Ca²⁺]_i in the incubation media to approximately 75 nM. Next, Ca²⁺ release was measured by adding increasing concentrations of IP₃ (solid arrows) and 1 mM carbachol (dotted arrows). *C*, quantitation of the results shown in *B*. *D*, parotid acini were stained for SERCA2b, total PMCA, and PMCA isoform 4 (PMCA4). Expression of PMCA increased in the apical membrane of *Homer2*^{-/-} cells. Data are presented as the mean ± S.E.

incubated with 10 μl of anti-PMCA (5F10) or 10 μl of anti-FLAG antibodies for 2 h by rocking at 4 °C. Protein A/G-agarose (Thermo Fisher Scientific Inc., Waltham, MA) was added to each mixture, and rocking was continued overnight at 4 °C. Protein A/G-agarose was pelleted at 1,000 $\times g$ for 10 s, and the beads were quickly washed with cold PBS. The immunoprecipitated proteins were separated by SDS-PAGE and probed with anti-FLAG (M2, 1:1,000) by overnight incubation at 4 °C.

Measurement of [Ca²⁺] Efflux—To measure directly the rate of Ca²⁺ efflux by PMCA, we measured the appearance of Ca²⁺ in the external medium using the procedure published by Zhao *et al.* (24) with slight modifications. Intact parotid acini from WT and *Homer2*^{-/-} mice were washed once and then suspended in 1 ml of medium containing 140 mM NaCl, 5 mM KCl, 10 mM glucose, and 10 mM HEPES, pH 7.4, with NaOH. 100 μl of cell suspension were added to 1.5 ml of a similar medium containing 2 μM free acid Fura-2 in a cuvette. After initiation of fluorescence recording, 10 μM EGTA was added to reduce the extracellular Ca²⁺ concentration ([Ca²⁺]_o) to ~100 nM. After establishing a baseline leak for ~1 min, the cells were stimulated with 1 mM carbachol. At the end of the experiment, the signals were calibrated by adding 1 mM CaCl₂ and then 1 mM MnCl₂ to the medium as described previously (24). The design of experiments for PMCA stimulation while measuring cytoplasmic Ca²⁺ was as follows. The cells were stimulated with 1 mM carbachol and 100 μM CPA for about 15–20 s to release most of the Ca²⁺ from internal stores and inhibit the SERCA Ca²⁺ pumps. Efflux by PMCA was initiated by inhibition of the muscarinic receptors with the antagonist 10 μM atropine and removal of external Ca²⁺ to prevent Ca²⁺ influx. CPA was maintained to inhibit the SERCA pumps.

Measurement of Ca²⁺ Uptake and Release from Internal Stores—IP₃-mediated Ca²⁺ release from internal stores was measured in SLO-permeabilized cells as described previously (24). Cells were washed with a high K⁺ (120 mM KCl, 20 mM NaCl, 10 mM glucose, and 10 mM HEPES, pH 7.4, with NaOH), Chelex-treated medium and added to Chelex-treated medium containing an ATP regeneration system (composed of 3 mM

ATP, 5 mM MgCl₂, 10 mM creatine phosphate, and 5 units/ml creatine kinase), a mixture of mitochondrial inhibitors, 2 μM Fluo-3, and 3 mg/ml SLO (Difco). After the addition of cells, the concentration of free Ca²⁺ in this medium was ~350–400 nM. In this medium, the cells were permeabilized almost instantaneously so that Ca²⁺ uptake into the ER could be measured immediately. Uptake of Ca²⁺ into the ER was allowed to continue until [Ca²⁺] in the medium was stabilized. Then IP₃ was added in increasing concentrations to measure the extent of Ca²⁺ release and the potency of IP₃ in mobilizing Ca²⁺ from the ER. Subsequent addition of 1 mM carbachol was used to monitor the receptor-evoked Ca²⁺ release.

Data Analysis and Statistics—All numeric values are represented as the mean ± S.E. Statistical significance, determined to be $p < 0.05$, was calculated using the Student's unpaired *t* test.

RESULTS

***Homer2* Deletion Does Not Affect the Polarized Expression of IP₃Rs, IP₃-mediated Ca²⁺ Release, or SERCA Activity**—Comparison of the receptor-evoked Ca²⁺ signaling in wild-type (WT) and *Homer2*^{-/-} acini revealed the altered signaling in the *Homer2*^{-/-} acini. To further examine this phenomenon, we analyzed the expression and activity of key Ca²⁺ transporters. IP₃Rs are established binding partners of Homer proteins (10, 12). Therefore, we first examined the localization and expression of Homer2 and IP₃Rs in parotid acinar cells from WT and *Homer2*^{-/-} mice. In WT parotid acinar cells, Homer2, and IP₃Rs, positive staining was primarily observed in the apical pole. As expected, Homer2 staining was not detected in *Homer2*^{-/-} acini, and the expression and localization of all IP₃R isoforms remained unchanged in *Homer2*^{-/-} acini (Fig. 1A).

We next examined whether *Homer2* deletion affects Ca²⁺ uptake into the ER, as well as the activity of IP₃Rs and their response to IP₃. Parotid acinar cells from WT and *Homer2*^{-/-} mice were permeabilized with SLO within 10–15 s, and the [Ca²⁺] in the incubation media was reduced to 50–80 nM within 2 min at 37 °C by SERCA-mediated Ca²⁺ uptake into the IP₃-mobilizable pool, similar to the method used with pancre-

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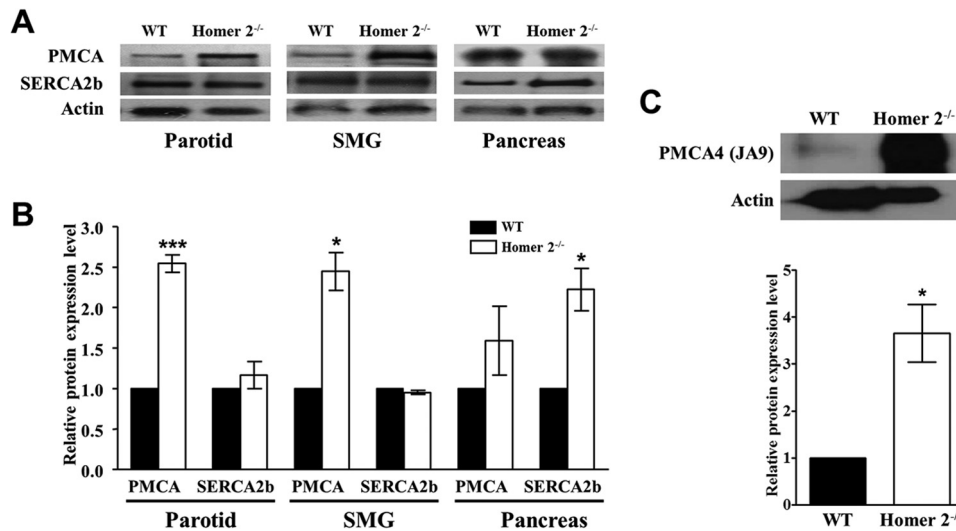


FIGURE 2. **Expression of Ca^{2+} pumps in WT and *Homer2*^{-/-} cells.** *A*, total PMCA and SERCA2b levels were analyzed by Western blot prepared from parotid, submandibular gland, and pancreatic acini from four WT and four *Homer2*^{-/-} mice. Band intensity was analyzed by densitometry. *B*, quantitation of Western blot data in *A*. *C*, PMCA4 expression increased in the parotid of *Homer2*^{-/-} mice. Quantitation is shown in the *bar graph*. Data were normalized to expression levels in WT cells and are presented as the mean \pm S.E. *, $p < 0.05$; ***, $p < 0.001$ (compared with WT).

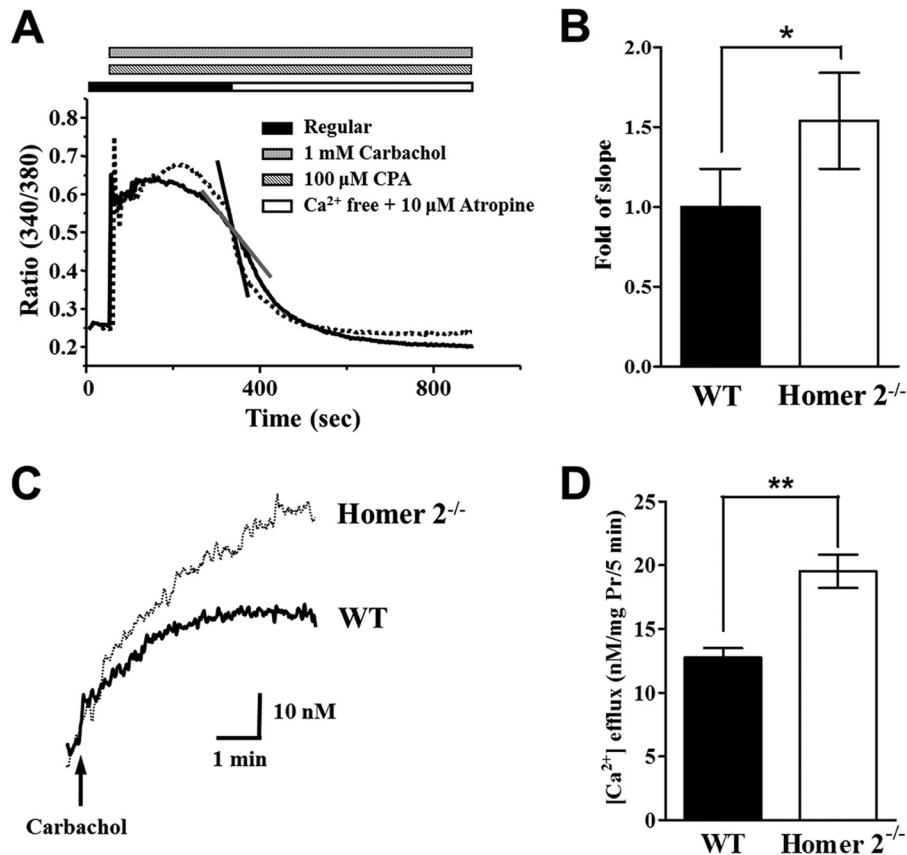


FIGURE 3. **Characterization of Ca^{2+} signaling in WT and *Homer2*^{-/-} cells.** *A*, parotid acini from WT and *Homer2*^{-/-} mice were stimulated with 1 mM carbachol while inhibiting SERCA with 100 μM CPA to elevate $[\text{Ca}^{2+}]_i$ and prevent Ca^{2+} uptake by the ER. When $[\text{Ca}^{2+}]_i$ stabilized at a steady state representing a balance of Ca^{2+} efflux and influx, Ca^{2+} clearance was initiated by removing external Ca^{2+} and adding of 10 μM atropine. *B*, average rate of Ca^{2+} clearance was determined from the slope of $[\text{Ca}^{2+}]_i$ decline as a measure of PMCA activity. Results were normalized to the slope obtained from WT cells and are presented as the mean \pm S.E. The rate of Ca^{2+} clearance increased significantly in *Homer2*^{-/-} cells. *C*, WT and *Homer2*^{-/-} parotid acinar cells in lightly Ca^{2+} -buffered media were exposed to 1 mM carbachol while measuring $[\text{Ca}^{2+}]_o$. *Homer2*^{-/-} cells exhibited an increased rate of $[\text{Ca}^{2+}]_o$, indicating a higher rate of PMCA in *Homer2*^{-/-} than cells from WT mice. *D*, quantitation of multiple experiments similar to those shown in *C*. The average rate of increased $[\text{Ca}^{2+}]_o$ was significantly higher in *Homer2*^{-/-} cells. Data are depicted as the mean \pm S.E. *, $p < 0.05$; **, $p < 0.01$ (compared with WT).

atic acinar cells (14). The rate and extent of Ca^{2+} uptake into the ER was similar in WT and *Homer2*^{-/-} cells, providing the first indication that, unlike in the pancreas (14), *Homer2* dele-

tion did not affect parotid acinar cell SERCA activity. The addition of increasing concentrations of IP_3 and the muscarinic agonist carbachol resulted in a similar increase in $[\text{Ca}^{2+}]_o$ due to

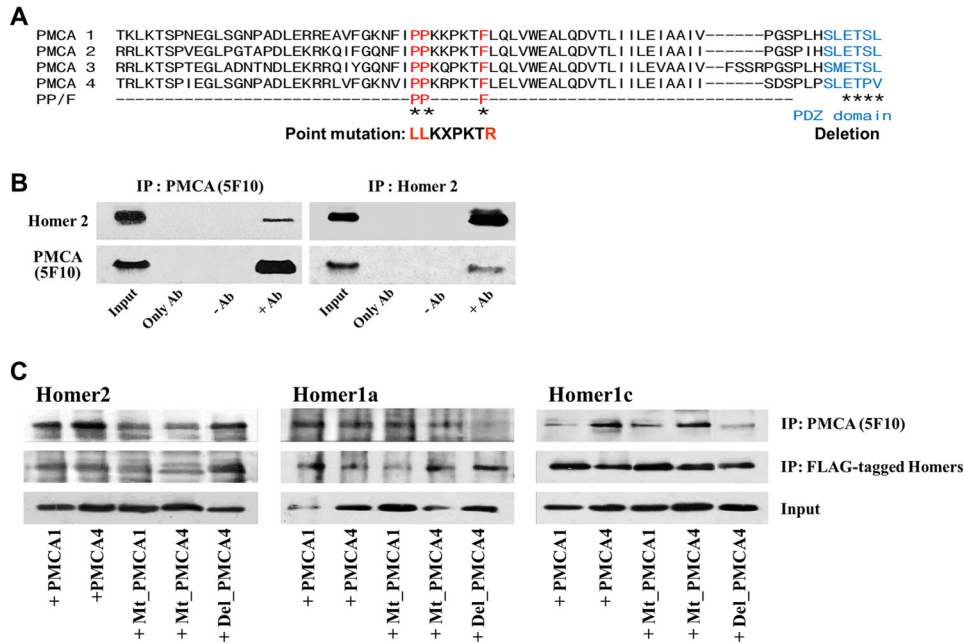


FIGURE 4. Interaction of Homer proteins with the PPXXF-like motif and PDZ-binding domain of PMCA. *A*, amino acid sequences of mouse PMCA isoforms were aligned using BLAST. Alignments show the mutations in the PPXXF-like motif (red) and the PDZ-binding motif of PMCA (blue). *B*, PMCA and Homer2 co-immunoprecipitate from cell lysates prepared from parotid acini of WT mice. In immunoprecipitation (IP) experiments, Input denotes extract samples used for Western blot; Only Ab denotes control IP using antibodies without extract in the IP assay; -Ab denotes control IP using extract without antibodies in the IP assay; and +Ab denotes IP using extract and antibodies in the IP assay. *C*, co-IP of Homer proteins (FLAG-tagged) and PMCA (wild-type, PPXXF-like motif mutants (Mt_PMCA), or PMCA4 with deleted PDZ-binding motif (Del_PMCA4) co-transfected in HEK293 cells. Extracts prepared from the cells co-transfected with the respective FLAG-Homers and the indicated PMCA constructs were used to IP either PMCA with the 5F10 antibodies or the Homers with anti-FLAG antibodies, and the precipitates were probed with anti-FLAG antibodies to detect the Homers. Note that Mt_PMCAs reduced the interaction of PMCA with Homer2 but not the interaction of PMCA with Homer1a and Homer1c. However, the interaction of PMCA with Homer1a and Homer1c was decreased when the Del_PMCA4 mutant was used.

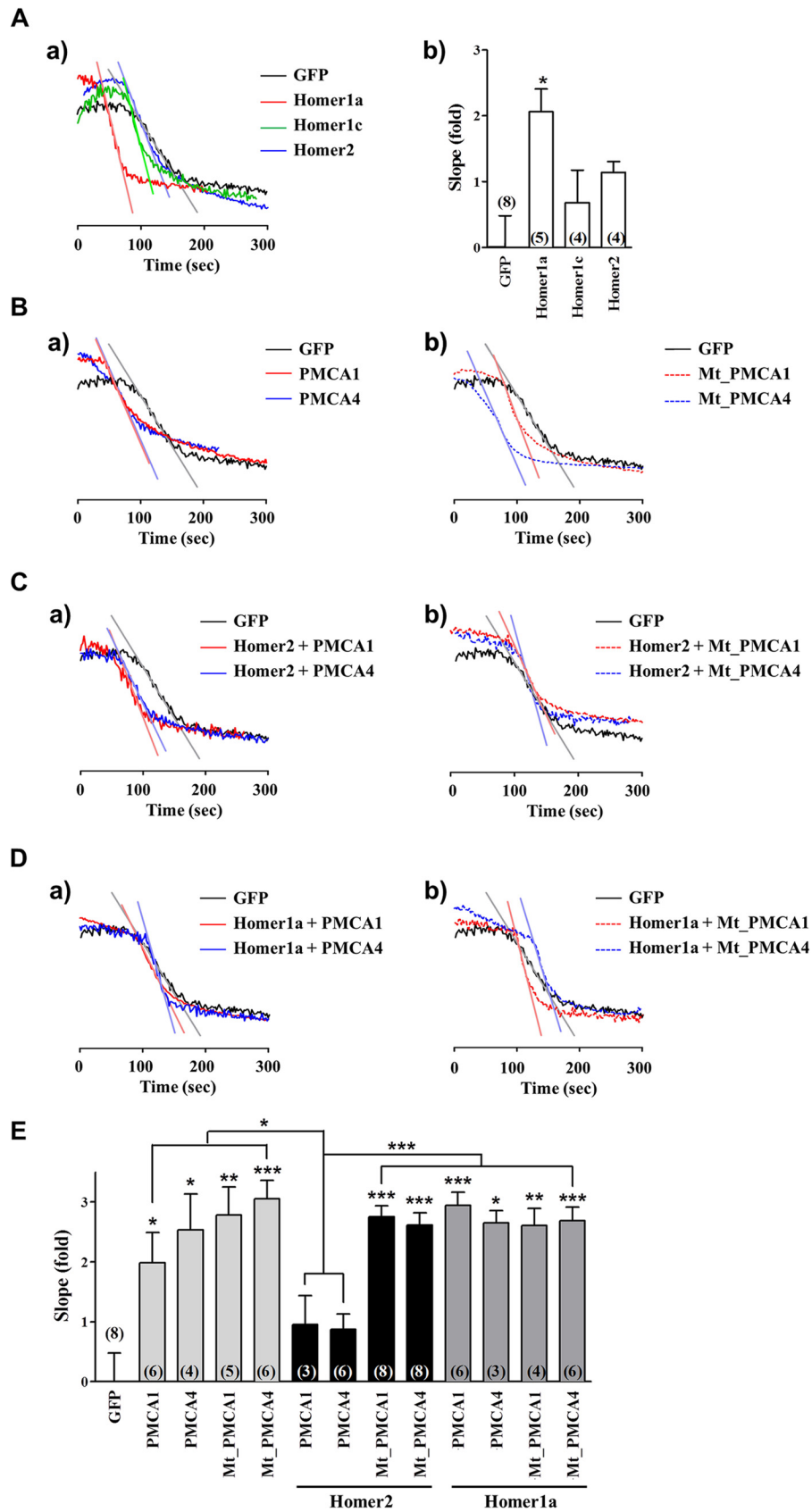
Ca^{2+} release from stores of SLO-permeabilized WT and *Homer2*^{-/-} cells (Fig. 1, *B* and *C*). Thus, no compensatory effects in expression, localization, and IP₃R activity were observed in *Homer2*^{-/-} mice.

PMCA Expression Is Selectively Increased in Specific Tissues from Homer2^{-/-} Mice—Next, we examined the expression of the SERCA and PMCA pumps. Because both PMCA1 and PMCA4 are expressed in salivary gland cells, we used a pan anti-PMCA antibody, 5F10, which detects both isoforms, to determine PMCA expression. To verify this PMCA expression, we used the anti-PMCA antibody JA9, which is specific for PMCA4 (14, 24). Notably, immunostaining experiments suggested increased PMCA expression in the apical region of *Homer2*^{-/-} parotid acinar cells, whereas the SERCA2 expression remained unaffected in these mice (Fig. 1*D*). To further analyze how Ca^{2+} pump expression was affected by *Homer2* deletion, we examined protein expression using Western blotting analyses and parotid membranes prepared from WT and *Homer2*^{-/-} parotid acinar cells. As shown in Fig. 2, *A* and *B*, PMCA expression in parotid acinar cells from *Homer2*^{-/-} mice increased significantly to 2.5 ± 0.1-fold greater than parotid acinar cells from WT mice (*n* = 4, *p* < 0.001). However, the expression of SERCA2b (1.2 ± 0.2-fold of WT, *n* = 4) remained unchanged in the parotid membranes. Similar results were obtained with submandibular gland membranes. Interestingly, however, opposite results were observed in pancreas membranes, suggesting a tissue-specific adaptive response to *Homer2* deletion. Furthermore, the protein levels of the PMCA4 isoform were higher in *Homer2*^{-/-} parotid mem-

branes compared with WT (3.7 ± 0.6-fold of WT, *n* = 4, *p* < 0.05, Fig. 2*C*).

Rate of [Ca²⁺]_{efflux} Is Increased in Homer2^{-/-} Parotid Cells—The major routes for Ca^{2+} clearance in nonexcitable cells, such as parotid acinar cells, are Ca^{2+} uptake into the ER by the SERCA pumps and Ca^{2+} efflux across the plasma membrane by PMCA (25–27). The results shown in Figs. 1*B* and 2 suggest that *Homer2* deletion does not affect SERCA expression and activity but increases the PMCA expression. To determine whether increased PMCA protein expression translates to increased PMCA activity in intact cells, we examined PMCA-mediated Ca^{2+} clearance in WT and *Homer2*^{-/-} parotid acini. In the first protocol, cells were stimulated with a high concentration of carbachol and treated with the SERCA inhibitor CPA to release ER Ca^{2+} and maximally activate Ca^{2+} influx to cause a large increase in cytoplasmic Ca^{2+} . Ca^{2+} clearance was then initiated by terminating cell stimulation with atropine while simultaneously inhibiting SERCA activity. Fig. 3*A* shows that the addition of 10 μM atropine in a Ca^{2+} -free solution resulted in an immediate clearance of Ca^{2+} primarily by PMCA (Fig. 3*A*). Comparing the slope of Ca^{2+} clearance revealed that Ca^{2+} clearance in *Homer2*^{-/-} cells is 1.5-fold faster compared with WT cells (1.5 ± 0.3-fold, *n* = 4, *p* < 0.05, Fig. 3*B*). In a second protocol, we assayed PMCA activity by measuring the change in $[Ca^{2+}]_o$ in cells incubated in media with low external Ca^{2+} concentration and stimulated with 1 mM carbachol. As shown in Fig. 3*C*, $[Ca^{2+}]_o$ increased significantly in response to carbachol stimulation in both cell types. Importantly, the change in

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[Ca²⁺]_o was ~1.5-fold higher in *Homer2*^{-/-} cells compared with WT cells (*n* = 4, *p* < 0.01, Fig. 3D).

Homer2 Interacts with and Regulates PMCA Expression—To identify potential interacting sites between PMCA and the Homer proteins, we searched for a PPXXF-like motif in PMCA that may interact with the Ena/VASP homology 1 domain of Homer proteins (16). The only such potential motif is present in the N-terminal 91–98 residues of several PMCA subtypes; however, proline and phenylalanine are separated by more than two residues (Fig. 4A). To determine whether this region functions as a Homer-binding motif to mediate the PMCA-Homer2 interaction, we first examined whether Homer2 selectively binds PMCA in parotid acinar cells from WT mice using co-immunoprecipitation assays. As shown in Fig. 4B, PMCA co-immunoprecipitates with endogenous Homer2 suggesting that endogenous Homer2 associates with PMCA.

To examine whether Homer proteins bind PMCA1 and/or PMCA4 in cell lysates and whether mutations in the potential Homer-binding motif of PMCA affect the interaction, we performed co-immunoprecipitation assays using HEK293 cells that transiently express Homer2 and PMCA or PMCA mutants. As shown in Fig. 4C, Homer2 co-immunoprecipitated with PMCA1b and PMCA4b but did not co-immunoprecipitate with the PMCA mutants (Mt_PMCA), which changes the PP/LL and F/R of the PPXXF-like motif. These results indicate that the PPXXF-containing region functions as a Homer-binding motif and mediates Homer2 interaction with PMCA. To determine whether these mutants interact with other Homer proteins, we examined the association between Homer1 and PMCA in transfected HEK293 cells. Mutation of PMCA did not affect the interaction with short Homer1a, which lacks the coiled-coil domain, or long Homer1c. This result suggests significant specificity of the Homer-binding motif on PMCA toward Homer2.

PMCA isoforms 2b and 4b have PDZ domain-binding ligands in their C terminus that interact with various scaffolding proteins such as membrane-associated guanylate kinase, NHERF2 (Na⁺/H⁺ exchanger regulatory factor-2), NOS-I (nitric-oxide synthase I), and Homer1 (Ania-3) (28–31), some of which interact with the Homers. Therefore, we examined whether mutation of the PDZ ligand of PMCA4 (marked Del_PMCA4) affected the interaction of Homers with PMCA. Co-expression of Homers and Del_PMCA4 showed reduced interaction of Homer1a and Homer1c with Del_PMCA4 compared with their interaction with WT PMCA4. However, interaction of Homer2 was unaffected (Fig. 4C). These results indicate that the PDZ ligand of PMCA4b may function to mediate Homer1 variant interaction with PMCA.

A previous study suggests that exogenous expression of Homer1a, Homer1c, or Homer2a had no effect on endogenous PMCA function, whereas knockdown of Homer1 slowed PMCA-mediated Ca²⁺ clearance in neuronal cells (44). We re-

examined these effects in nonexcitable cells by expressing the Homers in HEK293 cells and measured native PMCA activity. Transfection of Homer1a enhanced native PMCA activity, whereas transfection of the long Homers, Homer1c or Homer2, had no effect (Fig. 5A). These findings suggest that the level of endogenous long Homers are saturating with respect to PMCA in HEK293 cells, and thus further expression had no inhibitory effect. Accordingly, Homer1a likely relieved the tonic inhibition by the long Homers to activate the native PMCA activity. Therefore, to examine the relationship between the Homer-binding motif and PMCA activity, we measured the effect of Homer proteins on Ca²⁺ extrusion by expressed WT and mutant PMCA pumps. Overexpression of WT PMCA pumps increased Ca²⁺ clearance, and expression of the PPXXF-like mutants additionally accelerated Ca²⁺ clearance (Fig. 5, B and E). Co-transfection of Homer2 and WT PMCA pumps significantly slowed Ca²⁺ clearance. This inhibitory effect was abolished when Homer2 was co-expressed with the PMCA mutants (Fig. 5, C and E). However, Homer1a did not increase PMCA activity when cells were co-transfected with WT and mutant PMCA pumps compared with the overexpression of Homer1a or WT and mutant PMCA pumps (Fig. 5, D and E). These results indicate that Homer2 interaction with the PPXXF-like motif on PMCA is essential to regulate Ca²⁺ clearance and Ca²⁺ signaling in nonexcitable cells.

DISCUSSION

In this study, we demonstrate a novel interaction between Homer2 and PMCA in native parotid acinar cells. Furthermore, we provide evidence in support of a critical role for Homer2 in modulating PMCA activity and thus Ca²⁺ signaling. Previous reports that focused on the molecular structure of Homer proteins in the CNS indicated that these proteins bind GPCRs, such as mGluR1/5, and IP₃Rs, as well as act as scaffolding proteins for assembling Ca²⁺-signaling complexes in cellular microdomains (8–10, 13, 32). The apical region of exocrine cells is equivalent to the CNS synaptic region, where signaling complexes are clustered to form a “trigger zone” from which all forms of Ca²⁺ signals, including Ca²⁺ waves, are initiated (33, 34). Consistently, immunocytochemical studies have demonstrated that all IP₃R types are highly enriched in the apical region (14, 35, 36). Accordingly, we found that Homer2 and IP₃Rs are expressed in the apical pole of parotid acinar cells. However, *Homer2* deletion had no effect on the expression or function of any of the IP₃R isoforms, suggesting that Homer2 may have another role on Ca²⁺ signaling in parotid acinar cells.

Further analyses revealed that Homer2 affected PMCA expression in the apical membranes of parotid acinar cells. Hence, the most interesting finding from our study is the adaptive increase in PMCA protein expression in *Homer2*-deficient parotid and submandibular gland acinar cells. Like SERCA, the PMCA has a crucial role in maintaining Ca²⁺ homeostasis

FIGURE 5. Effects of the Homers on Ca²⁺ clearance by PMCA in HEK293 cells. A, overexpression of Homer1a increases the rate of Ca²⁺ clearance, whereas expression of Homer1c and Homer2 do not (*panel a*). *Panel b*, quantitation of the results in (*panel a*). B, overexpression of WT PMCA (*panel a*) and Mt_PMCA (*panel b*) increase the rate of the Ca²⁺ clearance. C, overexpression of Homer2 significantly decreases the effect of WT PMCA (*panel a*) but not of Mt_PMCA (*panel b*) on Ca²⁺ efflux in co-transfected HEK293 cells. D, overexpression of Homer1a does not activate WT PMCA (*panel a*) or Mt_PMCA (*panel b*) in co-transfected cells. E, quantitation of PMCA activity in B–D. Results were normalized to GFP expression levels and are presented as the mean ± S.E. *, *p* < 0.05; **, *p* < 0.01; ***, *p* < 0.001 (compared with GFP).

Regulation of PMCA-mediated Ca^{2+} Signaling by Homer2

(37). Previous studies suggested that PMCA is expressed in both the lateral and apical regions of pancreatic, submandibular gland, and parotid cells, as well as in the brain (31, 38–41). In addition, PMCA co-localize with mGluR1, $\text{IP}_3\text{R1}$, and Homer proteins, including Homer1a, Homer1b/c, and Homer3, in neurons (31, 40, 41). Here, we discovered that Homer2 interacts with PMCA and decreases the rate of $[\text{Ca}^{2+}]_i$ clearance in native parotid acinar cells, which is absent in *Homer2*-deficient mice. These results are similar to previous findings showing up-regulation of specific PMCA isoforms due to adaptations in Ca^{2+} signaling and Ca^{2+} -dependent cellular functions in pancreatic and submandibular gland acinar cells from *serca2*-deficient mice (24). Therefore, it appears that PMCA expression and function is particularly sensitive to perturbations in Ca^{2+} signaling, and cells modify PMCA activity to adjusting the Ca^{2+} signal.

Adaptations of PMCA expression may be regulated by Ca^{2+} itself. Several studies have shown that Ca^{2+} can alter the expression levels of Ca^{2+} -signaling components, such as pumps and channels, thereby maintaining flexibility in Ca^{2+} -signaling remodeling systems (3). In primary cultured cerebellar granule cells, PMCA2, PMCA3, and $\text{IP}_3\text{R1}$ expression is up-regulated during the activation of Ca^{2+} -dependent events. In contrast, type 2 $\text{Na}^+/\text{Ca}^{2+}$ exchanger and PMCA4 are rapidly down-regulated (1, 42, 43). An interesting aspect of our findings is that *Homer2*-deficient parotid cells exhibit increased expression and activity of PMCA but not of SERCA2. However, *Homer2*-deficient pancreatic acinar cells increase SERCA2 but not PMCA expression, suggesting that PMCA adaptation occurs in a tissue-specific manner, perhaps reflecting specific cellular functions and localization.

PMCA interacts with partner molecules through their PDZ-binding motif in the C terminus (28–30). In this study, we characterized an additional novel binding region (PPXXF-like motif) of PMCA isoforms that participate in the interaction with, and inhibition of, PMCA by Homer2. Thus, mutation of the PPXXF-like motif reduced the interaction of PMCA1 and PMCA4 with Homer2 (Fig. 4C) and prevented inhibition of ectopically expressed PMCA by Homer2 (Fig. 5E). In contrast, the PMCA4 PDZ motif does not appear to be essential for the interaction with Homer2 because its deletion did not prevent the interaction of PMCA4 with Homer2 (Fig. 4C).

Interestingly, the role of the PPXXF-like motif appears to be specific for the interaction of PMCA with Homer2. Thus, mutation of the PPXXF-like motif does not affect the interaction of PMCA with a short or long Homer1 (Fig. 4C) or activation of PMCA activity by Homer1a in co-transfected HEK293 cells (Fig. 5, D and E). These results were unexpected in view of the reports that Homer1a (Ania-3) retains binding to the PDZ-binding motif in the C terminus of PMCA isoforms and activates $[\text{Ca}^{2+}]_i$ clearance by PMCA when expressed with long Homer1 or the N terminus of Homer2 as an analog of Homer1a in neuronal cells (31, 44). Using expression in HEK293 cells, we confirmed interaction of Homer1 with the PDZ motif of PMCA. We found that this interaction is eliminated by deletion of the PDZ motif on PMCA4, as was reported previously (31). However, unlike a previous report, the Ca^{2+} clearance rate by native PMCA was increased by overexpression of short

Homer1a but not by overexpression of long Homers (Fig. 5A) (44). These experiments suggest that the level of long Homers in HEK293 cells (and perhaps other nonexcitable cells) is saturating with respect to PMCA, and thus additional expression of the long Homers caused no further inhibition. A prediction of this interpretation is the deletion of the long Homers, and their inhibition by the short Homer1a should accelerate PMCA activity and Ca^{2+} clearance. This was indeed the case, in which deletion of Homer2 in mice or expression of Homer1a in HEK293 cells increased the native PMCA activity.

In summary, the key findings of our study are as follows: 1) Homer2 binding to the PPXXF-like motif of PMCA inhibited PMCA activity, and 2) expression and binding of Homer1 to the PDZ domain of PMCA increased their activity. Thus, PMCA undergoes dual regulation by Homer proteins, *i.e.* inhibition by the Homer2 and stimulation by Homer1. In this manner, Homer proteins can regulate the duration of the Ca^{2+} signal in parotid acinar cells to either extend or shorten the signal by inhibition or stimulation of PMCA activity, respectively.

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