



Ca²⁺-CaMKK β pathway is required for adiponectin-induced secretion in rat submandibular gland

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Abstract

Adiponectin functions as a promoter of saliva secretion in rat submandibular gland via activation of adenosine monophosphate-activated protein kinase (AMPK) and increased paracellular permeability. Ca²⁺ mobilization is the primary signal for fluid secretion in salivary acinar cells. However, whether intracellular Ca²⁺ mobilization is involved in adiponectin-induced salivary secretion is unknown. Here, we found that full-length adiponectin (fAd) increased intracellular Ca²⁺ and saliva secretion in submandibular glands. Pre-perfusion with ethylene glycol-bis (2-aminoethylether)-*N,N,N',N'*-tetraacetic acid (EGTA) combined with thapsigargin (TG), an endoplasmic reticulum Ca²⁺-ATPase inhibitor, abolished fAd-induced salivary secretion, AMPK phosphorylation, and enlarged tight junction (TJ) width. Furthermore, in cultured SMG-C6 cells, co-pretreatment with EGTA and TG suppressed fAd-decreased transepithelial electrical resistance and increased 4-kDa FITC-dextran flux responses. Moreover, fAd increased phosphorylation of calcium/calmodulin-dependent protein kinase (CaMKK β), a major kinase that is activated by elevated levels of intracellular Ca²⁺, but not liver kinase B1 phosphorylation. Pre-perfusion of the isolated gland with STO-609, an inhibitor of CaMKK β , abolished fAd-induced salivary secretion, AMPK activation, and enlarged TJ width. CaMKK β shRNA suppressed, whereas CaMKK β re-expression rescued fAd-increased paracellular permeability. Taken together, these results indicate that adiponectin induced Ca²⁺ modulation in rat submandibular gland acinar cells. Ca²⁺-CaMKK β pathway is required for adiponectin-induced secretion through mediating AMPK activation and increase in paracellular permeability in rat submandibular glands.

Keywords Adiponectin · Calcium · Calcium/calmodulin-dependent protein kinase · Adenosine monophosphate-activated protein kinase · Submandibular gland · Secretion

Introduction

Saliva is important for the initiation of digestion and the maintenance of oral health. Severe hyposalivation leads to dental caries, oral pain and mucosal infections (Kaplan and Baum 1993). Saliva secretion is accomplished through the interaction of autonomic neurotransmitters with their respective receptors (i.e., muscarinic acetylcholine cholinergic, α -adrenergic, and β -adrenergic receptors) on gland acinar cells (Baum 1993). Besides, there is increasing evidence that salivary secretion is also evoked by non-cholinergic or non-adrenergic pathways that utilize neuropeptides, such as substance P and neuropeptide Y, as well as other molecules released by nerve terminals (Garrett et al. 1999). In the previous study, we reported that adiponectin, an adipokine, promotes salivary secretion in rat submandibular glands via activation of adenosine monophosphate activated protein

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kinase (AMPK) and increased paracellular permeability (Ding et al. 2013).

Ca²⁺ mobilization plays a crucial role in saliva secretion. For example, activation of muscarinic cholinergic receptors (mAChRs) rapidly triggers the release of intracellular Ca²⁺ ([Ca²⁺]_i) from endoplasmic reticulum (Coronado et al. 1994; Joseph 1996), which subsequently evokes influx of Ca²⁺ from the extracellular medium, resulting in a sustained increase in [Ca²⁺]_i (Liu et al. 1998). The increase in [Ca²⁺]_i in turn causes the opening of Ca²⁺-gated K⁺ and Cl⁻ channels (Giovannucci et al. 2002; Nauntofte and Poulsen 1986; Nehrke et al. 2003), and up-regulates Na⁺/K⁺/2Cl⁻ cotransporter (Evans and Turner 1997), Na⁺/H⁺, and Cl⁻/HCO₃⁻ exchangers (Manganel and Turner 1990; Nguyen et al. 2004). Furthermore, increased [Ca²⁺]_i induces aquaporin 5 trafficking, resulting in the forming of water pores that facilitates the rapid increase in transcellular water permeability (Ishikawa et al. 1998). However, adiponectin-promoted saliva secretion in rat submandibular glands is independent of mAChRs (Ding et al. 2013). Whether Ca²⁺ mobilization is involved in adiponectin-induced salivary secretion is unknown.

AMPK plays a key role in adiponectin-mediated metabolic modulation and cardiovascular protection (Shibata et al. 2005; Yamauchi et al. 2002). Activation of AMPK is controlled by two upstream kinases, liver kinase B1 (LKB1) and calcium/calmodulin-dependent protein kinase (CaMKKβ) (Hawley et al. 2005; Woods et al. 2003). LKB1 has been considered as a constitutively serine/threonine protein kinase that is ubiquitously expressed in mammalian cells, and phosphorylates the catalytic subunit of AMPK by increasing cellular AMP:ATP ratio (Hawley et al. 2003). In contrast, the activation of AMPK by CaMKKβ is initiated by an increase in [Ca²⁺]_i, but not affected by changes in the ATP:AMP ratio (Woods et al. 2005). To date, whether LKB1 and/or CaMKKβ are involved in the adiponectin-mediated AMPK activation and saliva secretion in submandibular gland is not clarified.

Thus, the present study was designed to explore the role of adiponectin-induced intracellular Ca²⁺ mobilization in salivary secretion and its underlying mechanism. These findings will improve our understanding of the biological roles of adiponectin and Ca²⁺ in saliva secretion, and provide a potential therapeutic strategy for the treatment of submandibular gland dysfunction.

Materials and methods

Animal ethics

Healthy male Sprague Dawley (SD) rats weighing 250–270 g each were obtained from the Peking University Health

Science Center. All experimental procedures and procedures for the care and use of the animals were approved by the Peking University Institutional Review Board. All surgical procedures were performed under chloral hydrate anesthesia (400 mg/kg body weight), and all efforts were made to minimize the animals' suffering.

Reagents and antibodies

Rat recombinant full length adiponectin (fAd), ethylene glycol-bis (2-aminoethylether)-*N,N,N',N'*-tetraacetic acid (EGTA), thapsigargin (TG), and STO-609 were purchased from Sigma-Aldrich (St. Louis, MO, USA). Antibodies against CaMKKβ, p-LKB1, and LKB1 (Cat. Nos. sc-271924, sc-32245, and sc-50341, respectively) were from Santa Cruz Biotechnology (Carlsbad, CA, USA). Antibodies against p-CaMKKβ, p-AMPK, and AMPK (Cat. #12818, #2531, and #2532, respectively) were from Cell Signaling Technology (Danvers, MA, USA). Antibody against glyceraldehyde-3-phosphate dehydrogenase (GAPDH) (Cat. No. M2006) was from Abmart (Shanghai, China). Other chemicals and reagents were of analytical grade.

Rat submandibular gland acinar cell preparation

Primary isolated rat submandibular gland acinar cells were prepared by enzymatic digestion according to the method (Zhang et al. 1996, 2006) with minor modifications. Briefly, rat submandibular gland tissues were excised and dissected free of connective tissue, rinsed twice with ice-cold phosphate-buffered saline (PBS), cut into small pieces, and digested in medium containing 100 units/ml of collagenase (Worthington, Lakewood, UK) and 1% BSA for 60 min. The digestion was terminated and washed twice with Dulbecco's modified Eagle's media (DMEM) containing 5% fetal bovine serum (FBS) and centrifuged at 1000g for 5 min. The cells were then resuspended in DMEM containing 15% FBS and filtered through a single layer of nylon bolting cloth (150 mesh).

Measurement of [Ca²⁺]_i

Primary isolated acinar cells were loaded with Ca²⁺-sensitive fluorescent probe fluo-2/AM (Thermo Fisher Scientific, MA, USA) for 30 min at 37 °C. Excitation was performed at 488 nm and the emission signals were collected through a 515 nm barrier filter. Images were taken every 10 s, and quantitated by average fluorescence intensities in randomly selected three to five cells in each time point from five submandibular gland cells (Carl Zeiss LSM710, Gottingen, Germany).

Perfusion of isolated rat submandibular glands

The effect of adiponectin on the secretory function of submandibular glands was measured according to the methods described previously (Ding et al. 2013). Briefly, after anesthesia, the submandibular glands were isolated and perfused through a polyethylene cannula placed in the external carotid artery. The main excretory duct was cannulated for saliva collection. Krebs-Ringer-HEPES (KRH, 116 mM NaCl, 5.4 mM KCl, 1.25 mM CaCl₂, 0.4 mM MgSO₄, 20 mM HEPES, 0.9 mM Na₂HPO₄, and 5.6 mM glucose, pH 7.4) buffer was warmed to 37 °C, bubbled with 95% O₂ and 5% CO₂, and perfused through the glands at a rate of 1.8 ml/min using a Gilson Minipuls rotary pump. After equilibration for at least 30 min, the glands were perfused with various stimulators or inhibitors (n = 8 for each group) for 10 min. Secretion by the glands was measured as the length of a column of moisture on a piece of filter paper (35 mm × 5 mm). In each group, salivary flow rates of perfused glands were measured. Then gland tissues were collected and analyzed by Western blot and transmission electron microscopy.

Cell culture

The rat submandibular gland cell line SMG-C6 (a generous gift from Dr. David O. Quissell without commercial purpose) was routinely grown at 37 °C in a humidified 5% CO₂ atmosphere in DMEM/F12 (1:1 mixture) medium containing 2.5% FBS, 5 mg/ml transferrin, 1.1 mM hydrocortisone, 0.1 mM retinoic acid, 2 nM thyronine T3, 5 mg/ml insulin, 80 ng/ml epidermal growth factor, 50 mg/ml gentamicin sulfate, 5 mM glutamine, 100 U/ml penicillin, and 100 mg/ml streptomycin (Quissell et al. 1997). All constituents used in culturing SMG-C6 cells were purchased from Sigma-Aldrich Co.

Western blot analysis

The cultured cells were homogenized in lysis buffer (50 mM Tris-HCl, 150 mM NaCl, 1 mM EDTA, 1 mM phenylmethylsulfonyl fluoride, 1% Triton X-100, 0.1% SDS, and 0.1% sodium deoxycholate, pH 7.2) using a polytron homogenizer as previously described (Ding et al. 2013). The homogenates were centrifuged at 12,000g for 10 min at 4 °C. The protein concentration of the supernatant was measured by the Bradford method. Equal amounts of proteins (20 µg) from each sample were separated on 12% SDS-PAGE and electroblotted on polyvinylidene fluoride membranes. The blocked membranes were incubated with antibodies against p-CaMKKβ (1:1000), CaMKKβ (1:400), p-LKB1 (1:600), LKB1 (1:600), p-MAPK (1:1000), AMPK (1:1000), or GAPDH (1:4000), respectively. The blots were then probed with horseradish peroxidase-conjugated

secondary antibodies (ZSGBBIO, Beijing, China), and the target proteins were detected using enhanced chemiluminescence reagent (Pierce Biotechnology, Rockford, IL, USA). GAPDH was used as a loading control.

Transmission electron microscopy

The gland specimens were fixed in 2% paraformaldehyde-1.25% glutaraldehyde. Ultrathin sections were stained with uranyl acetate and lead citrate and examined with a transmission electron microscope (H-7000, HITACHI, Tokyo, Japan). Each image was obtained under the same conditions of brightness and contrast to permit comparison of tight junction (TJ) density among the different groups. For morphometric analysis, the distances between neighboring TJs (shown as the width of the apical TJs) were measured and averaged in ten randomly selected fields in each of four sections by two blinded examiners using ImageJ software (NIH, MD, USA) as previously described (Ding et al. 2013).

Knockdown and re-expression of CaMKKβ

SMG-C6 cells were cultured to 80% confluence and transfected with shRNA of interest using MegeTran 1.0 (Origene, MD, USA) according to the manufacturer's instructions. For knockdown of CaMKKβ, CaMKKβ shRNA (CAGCGACGCCTTGCTGTCTAACACCGTGG) and a scrambled control were constructed in pGFP-V-RS vectors and synthesized by Origene Technologies.

CaMKKβ re-expression ('rescue') was conducted by generating a Myc-tagged cDNA clone of CaMKKβ in a pCMV6 vector (Origene). Plasmid transfection was performed using MegeTran 1.0 at a transfection reagent: DNA ratio which was 3:1 as described in the manufacturer's instructions. The cells were collected 24 h post-transfection.

Transepithelial electrical resistance measurement and paracellular tracer flux assay

Confluent monolayers of SMG-C6 cells were grown in 24-well Corning Transwell™ chambers (polycarbonate membrane, filter pore size 0.4 µm, area 0.33 cm²; Costar) for 7 days, transepithelial electrical resistance (TER) was then measured at 37 °C using an epithelial volt ohm meter (EVOM; WPI, FL, USA). TER values were calculated by subtracting the blank filter (90 Ω) and by multiplying by the surface area of the filter. All measurements were performed on a minimum of three wells.

For paracellular tracer flux assay, 1 mg/ml 4-kDa FITC-dextran was added to the medium at the apical sides of the chambers, and the samples were collected from the basal sides of the chambers after incubation for 3 h. The apparent permeability coefficient (*P*_{app}) was determined as the

increase in the amount of tracer per time per filter area by using a fluorometer (BioTek, VT).

Statistical analysis

Data are presented as the mean \pm SD. Statistical analysis among multiple groups was performed by one-way ANOVA followed by Bonferroni's test using GraphPad software (GraphPad Prism, CA, USA). $P < 0.05$ was considered statistically significant.

Results

Effect of adiponectin on Ca^{2+} mobilization in isolated submandibular gland cells

Ca^{2+} was an important molecule in gland secretion. To reveal whether adiponectin could modulate Ca^{2+} in rat submandibular gland, we isolated rat submandibular glands acinar cells. As shown in Fig. 1a, b $\mu\text{g/ml}$ fAd caused an increase in $[\text{Ca}^{2+}]_i$ of isolated submandibular gland acinar cells, and the increased $[\text{Ca}^{2+}]_i$ could last for more than 5 min.

Ca^{2+} is required for adiponectin-induced secretion in submandibular glands

To identify the role of Ca^{2+} in the secretion induced by adiponectin, we performed *ex vivo* perfusion of isolated rat submandibular glands. The basal saliva flow during KRH perfusion was 3.41 ± 1.29 mm/5 min, which was consistent with previous study (Ding et al. 2013). $1 \mu\text{g/ml}$ fAd significantly increased salivary flow rate by 289.62%. Pre-perfusion with EGTA (1 mM), a Ca^{2+} chelator, or TG (1 μM), an endoplasmic reticulum Ca^{2+} -ATPase inhibitor, alone did not reduce fAd-induced increase in salivary secretion. However, pre-perfusion with EGTA combined with TG abolished fAd-induced secretion. EGTA and/or TG alone had no effect on the basal saliva flow (Fig. 1c).

Next, we measured CaMKK β and AMPK phosphorylation in these perfused glands (Fig. 1d). Compared with the KRH perfused glands, the levels of p-CaMKK β in fAd-perfused glands with or without EGTA or TG were increased by 162.83, 143.61, and 173.46%, respectively. However, pre-perfusion with EGTA combined with TG abolished fAd-induced increased CaMKK β phosphorylation. Perfusion with EGTA and/or TG alone did not affect the basal level of p-CaMKK β (Fig. 1e). With the same tissues, the levels of p-AMPK in fAd-perfused glands with or without EGTA or TG were increased by 230.41, 214.73, and 208.39%, respectively, as compared with the KRH perfused gland. Pre-perfusion with EGTA combined with TG abolished

fAd-induced increase in AMPK phosphorylation, whereas perfusion with EGTA and/or TG alone did not affect the basal level of p-AMPK (Fig. 1f). These results suggest that Ca^{2+} mobilization is required for adiponectin-induced salivary secretion, CaMKK β and AMPK activation in rat submandibular glands. Both influx of extracellular Ca^{2+} and release of Ca^{2+} from endoplasmic reticulum might involve in these processes.

Ca^{2+} is involved in adiponectin-modulated tight junction ultrastructure

TJs are specialized structures located in the apical regions of lateral membranes between neighboring cells, establishing a barrier to the diffusion of solutes through paracellular pathway (Tsukita et al. 2001). To explore the possible role of Ca^{2+} in the paracellular pathway regulated by adiponectin, we examined the morphology of TJs in perfused glands under transmitted electron microscope. TJs were located in the apical portions of lateral membranes in acini, forming a slightly opened paracellular channel in unperfused submandibular glands (Fig. 2a) and were not affected by KRH perfusion (Fig. 2b). Perfusion with fAd increased TJ distance between neighboring epithelial cells (Fig. 2c), consistent with our previous findings (Ding et al. 2013). Pre-treatment with EGTA or TG did not affect the increase in TJ width induced by fAd (Fig. 2d, e). However, pre-treatment with EGTA combined with TG suppressed fAd-induced increase in TJ width (Fig. 2f). EGTA and/or TG alone had no influence on TJ ultrastructure (Fig. 2g–i).

Quantitative analysis showed that the average width of apical TJs was 11.34 ± 0.04 nm in KRH-perfused glands, similar with that in unperfused glands. The width of apical TJs was increased by 105.35% in fAd-perfused glands, by 113.64% in glands pre-perfused with EGTA, and by 102.48% in glands pre-treated with TG, compared with the KRH perfused glands. Pre-perfusion with EGTA combined with TG suppressed the increase in TJ width induced by fAd, whereas EGTA and/or TG alone had no effect on the basal TJ width (Fig. 2j). These results suggest that Ca^{2+} , including both extracellular Ca^{2+} influx and Ca^{2+} released from endoplasmic reticulum, are involved in the "opening" of TJs regulated by adiponectin in submandibular glands.

Ca^{2+} contributes to the adiponectin-modulated paracellular permeability

We previously found that the adiponectin-induced saliva secretion in rat submandibular glands was accomplished by an increase in epithelial paracellular permeability (Ding et al. 2013). To determine whether Ca^{2+} is involved in the regulation of paracellular permeability by adiponectin, we performed TER measurement on monolayers of polarized

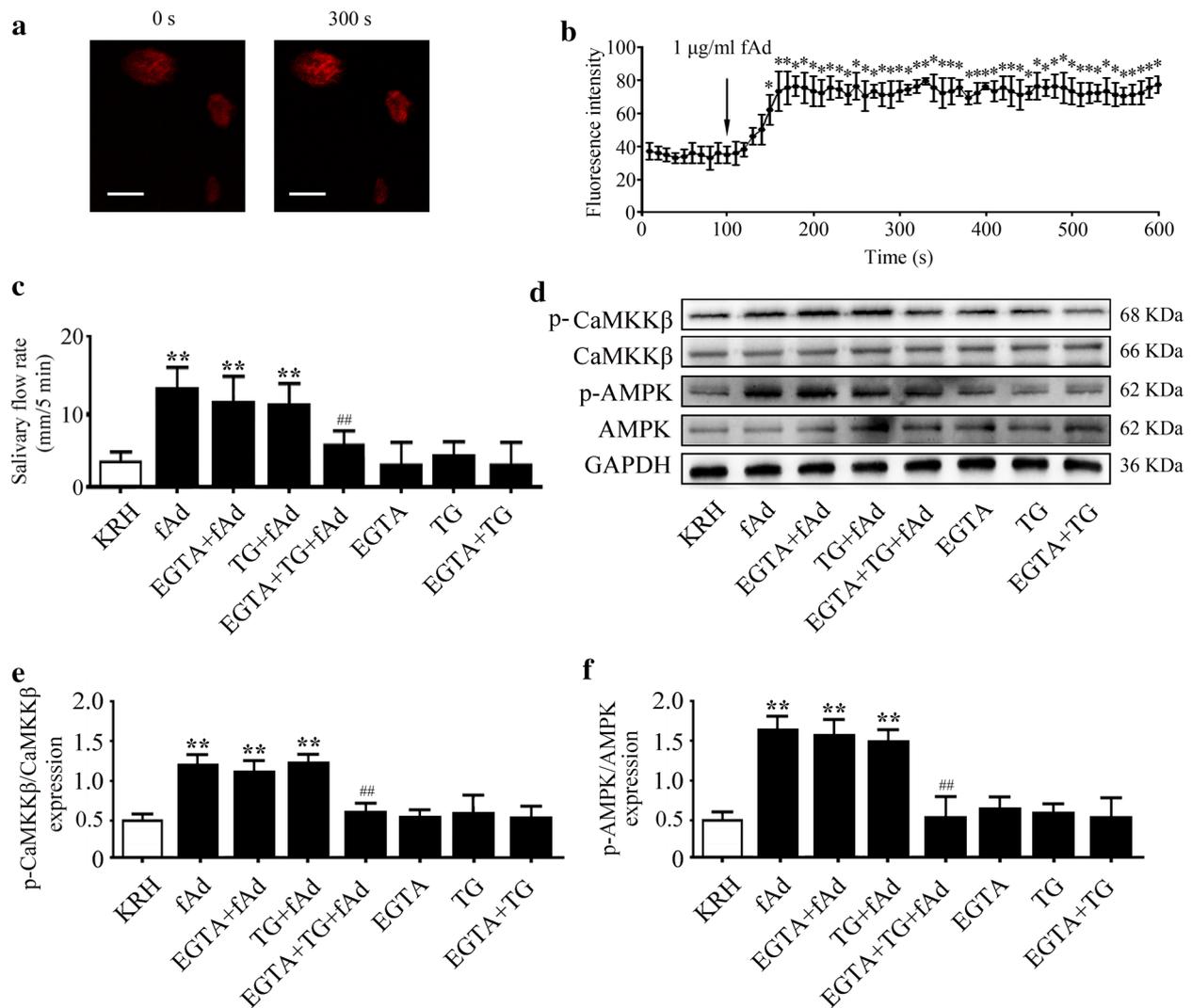


Fig. 1 Ca²⁺ mobilization is required for adiponectin-induced secretion in submandibular glands. **a** Representative fluorescence image of furo-2/AM-loaded cells before (left) and after (right) stimulation with 1 μg/ml fAd. Bar: 20 μm. **b** Quantitative analysis fluorescence intensities of submandibular cells stimulated by 1 μg/ml fAd and the time-course. Arrows denote the time of fAd application. The values shown are means ± SD from five independent experiments. **P* < 0.05 compared with the average intensities of untreated cells. Isolated rat submandibular glands were perfused with Krebs–Ringer–HEPES (KRH), and salivary flow rates were measured after equilibration for 30 min, perfusion with stimulators or inhibitors for 10 min. **c**

The salivary flow rate in glands perfused with KRH for 10 min, perfused with 1 μg/ml fAd for 10 min, pre-perfused with 1 μM EGTA for 30 min, and then fAd for 10 min, pre-perfused with 1 μM TG for 30 min, and then fAd for 10 min, pre-perfused with EGTA combined with TG for 30 min, and then fAd for 10 min, EGTA alone, TG alone, EGTA and TG. **d** p-CaMKKβ and p-AMPK expression in the above eight groups. **e** Quantitative analysis of p-CaMKKβ expression normalized to CaMKKβ. **f** Quantitative analysis of p-AMPK expression normalized to AMPK. The values shown are means ± SD from four independent experiments. **P* < 0.05 and ***P* < 0.01 compared with KRH group. ##*P* < 0.01 compared with fAd group

SMG-C6 cells. The basal TER value of untreated monolayers was $678 \pm 54.62 \Omega \text{ cm}^2$, consistent with previous studies (Ding et al. 2013; Kawedia et al. 2008). fAd caused a rapid and significant decrease in TER value (Fig. 3a, b). Pre-treatment with either EGTA or TG did not affect fAd-induced decrease in TER value (Fig. 3c, d). However, pre-treatment with EGTA combined with TG abolished fAd decreased TER value (Fig. 3e). EGTA and/or TG alone did not affect the basal TER values (Fig. 3f–h).

Paracellular permeability can also be evaluated using 4-kDa FITC-dextran as a non-charged paracellular tracer. Results showed that the *P*_{app} for 4-kDa FITC-dextran was greatly increased by fAd with or without EGTA/TG pre-treatment (Fig. 4a–d), whereas pre-treatment with EGTA combined with TG abolished fAd-induced increase in *P*_{app} (Fig. 4e). EGTA and/or TG alone did not affect the basal *P*_{app} (Fig. 4f–h). These data indicate that adiponectin regulates paracellular permeability in a Ca²⁺-dependent manner,

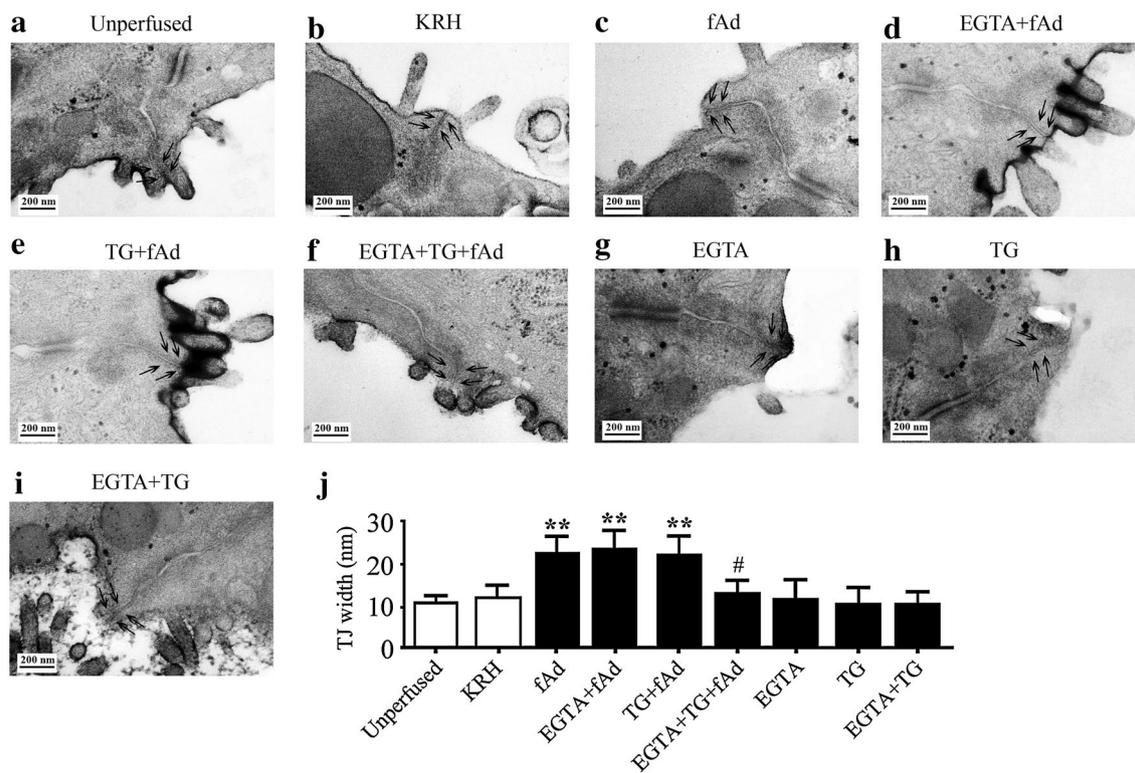


Fig. 2 Ca^{2+} is involved in adiponectin-modulated tight junction ultrastructure. Representative images of TJ structure in unperfused glands (a), in glands perfused with KRH (b), 1 $\mu\text{g}/\text{ml}$ fAd for 10 min (c), pre-perfused with 1 μM EGTA for 30 min, and then fAd for 10 min (d), pre-perfused with 1 μM TG for 30 min, and then fAd for 10 min (e), pre-perfused with EGTA combined with TG for 30 min, and then

fAd for 10 min (f), EGTA alone (g), TG alone (h), EGTA and TG (i). TJs are indicated by arrows. Bar: 200 μm . **j** The width of apical TJs was measured in ten randomly selected fields in four sections from each gland. The values shown are means \pm SD from four independent experiments. ** $P < 0.01$ compared with the KRH group. # $P < 0.05$ compared with fAd group

and both extracellular Ca^{2+} influx and intracellular Ca^{2+} release contribute to this effect.

CaMKK β is an upstream kinase for adiponectin-induced AMPK activation

LKB1 and CaMKK β have been identified as upstream kinases that activate AMPK (Woods et al. 2003, 2005). To determine whether LKB1 or CaMKK β are involved in adiponectin-induced AMPK activation in rat submandibular glands, we stimulated SMG-C6 cells with fAd for 10 min. As shown in Fig. 5a–d, fAd significantly increased the phosphorylation of CaMKK β and AMPK, whereas LKB1 phosphorylation was not changed. These results suggest that CaMKK β , but not LKB1, is responsible for adiponectin-induced activation of AMPK in submandibular glands.

CaMKK β is required for adiponectin-induced secretion in submandibular glands

To further reveal the role of CaMKK β in adiponectin-induced saliva secretion, the isolated submandibular glands

were perfused with STO-609, a CaMKK β antagonist. As shown in Fig. 5e, pre-perfusion with STO-609 (1 mM) abolished increased secretion induced by fAd, whereas STO-609 alone had no influence. Moreover, pre-perfusion with STO-609 abolished fAd-induced AMPK phosphorylation, whereas STO-609 alone did not change the level of p-AMPK (Fig. 5f, g). These results indicate that CaMKK β might be an up-stream kinase to modulate adiponectin-promoted AMPK phosphorylation.

CaMKK β is involved in adiponectin-modulated tight junction ultrastructure and paracellular permeability

We next examined whether CaMKK β is involved in the modulation of TJ ultrastructure and paracellular permeability induced by adiponectin. Perfusion with fAd increased TJ width. Pre-perfusion with STO-609 abolished fAd-induced increase in TJ width, while STO-609 alone had no effect on the ultrastructure of TJs (Fig. 6a–e). These results suggest that CaMKK β is involved in the TJ “opening” in response to adiponectin.

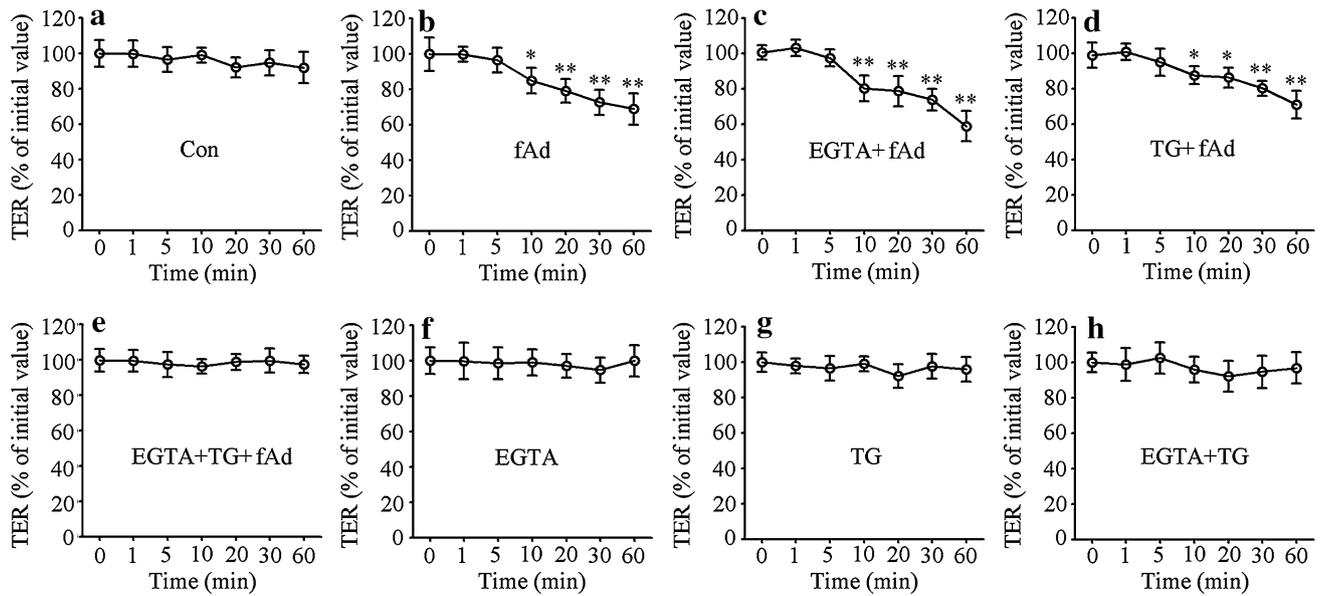


Fig. 3 Ca^{2+} is required for adiponectin-decreased TER values. The time course of transepithelial electrical resistance (TER) in SMG-C6 cells was measured using an epithelial volt ohm meter. The value of TER in untreated SMG-C6 cells (a), in cells incubated with 1 $\mu\text{g}/\text{ml}$ fAd for 60 min (b), pre-incubated with 1 μM EGTA for 30 min, and then fAd for 60 min (c), pre-incubated with 1 μM TG, and then

fAd for 60 min (d), pre-incubated with EGTA combined with TG, and then fAd for 60 min (e), EGTA alone (f), TG alone (g), EGTA and TG (h). The values shown are means \pm SD from four independent experiments. * $P < 0.05$ and ** $P < 0.01$ compared with untreated control cells

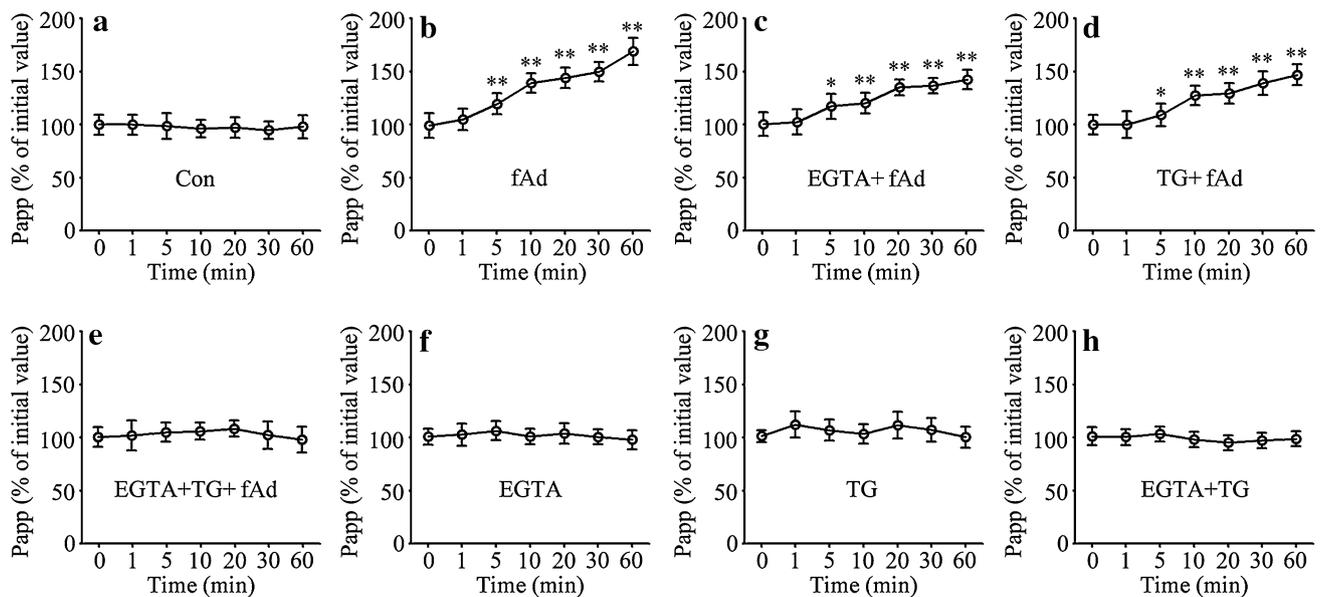


Fig. 4 Ca^{2+} is required for adiponectin-increased permeability of FITC-dextran. The apparent permeability coefficient (P_{app}) of 4-kDa FITC-dextran was determined as the increase in the amount of tracer per time per filter area. The P_{app} value in untreated SMG-C6 cells (a), in cells incubated with 1 $\mu\text{g}/\text{ml}$ fAd (b), pre-incubated with 1 μM EGTA, and then fAd for 60 min (c), pre-incubated with 1 μM TG,

and then fAd for 60 min (d), pre-incubated with EGTA combined with TG, and then fAd for 60 min (e), EGTA alone (f), TG alone (g), EGTA and TG (h). The values shown are means \pm SD from four independent experiments. * $P < 0.05$ and ** $P < 0.01$ compared with untreated control cells

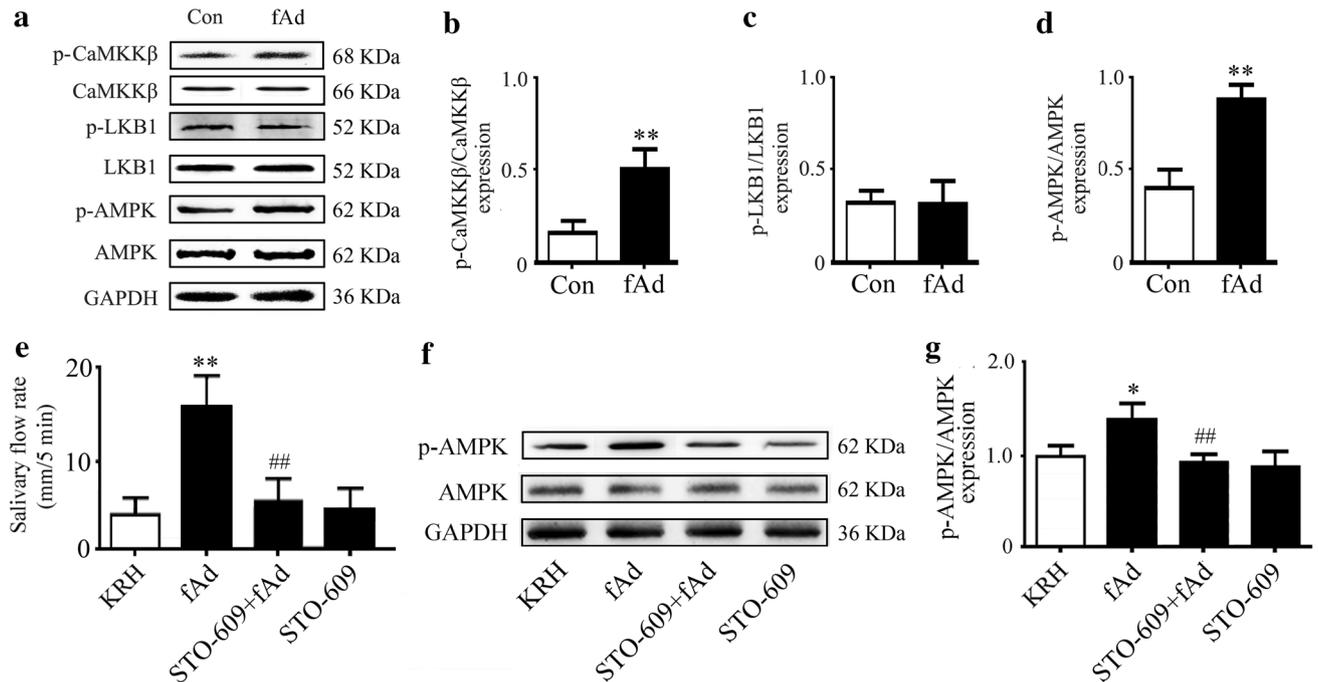
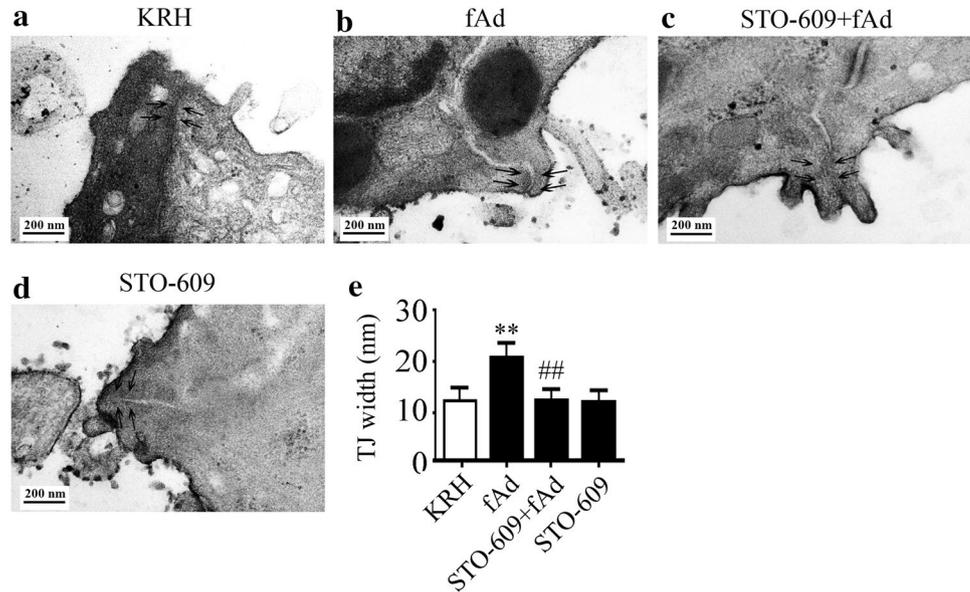


Fig. 5 CaMKK β is required for adiponectin-induced AMPK activation and secretion in submandibular glands. **a** Expression of p-CaMKK β , CaMKK β , p-LKB1, LKB1, p-AMPK, and AMPK in SMG-C6 cells stimulated with 1 μ g/ml fAd for 10 min. Quantitative analysis of p-CaMKK β expression normalized to CaMKK β (**b**), p-LKB expression normalized to LKB (**c**), p-AMPK expression normalized to AMPK (**d**). The values shown are means \pm SD from four independent experiments. ** P < 0.01 compared with untreated control cells. **e** The salivary flow rate in glands perfused with KRH

for 10 min, perfused with 1 μ g/ml fAd for 10 min, pre-perfused with 1 mM STO-609 for 30 min, and then fAd for 10 min, STO-609 alone. The values shown are means \pm SD from four independent experiments. ** P < 0.01 compared with KRH group. ## P < 0.05 compared with fAd group. **f** p-AMPK expression in the above four groups. **g** Quantitative analysis of p-AMPK expression normalized to AMPK. The values shown are means \pm SD from four independent experiments. * P < 0.05 compared with KRH group. ## P < 0.05 compared with fAd group

Fig. 6 CaMKK β is involved in adiponectin-modulated tight junction ultrastructure. Representative images of TJ in glands perfused with KRH (**a**), perfused with 1 μ g/ml fAd for 10 min (**b**), pre-perfused with 1 mM STO-609 for 30 min, and then fAd for 10 min (**c**), perfused with STO-609 alone (**d**). TJs are indicated by arrows. Bar: 200 μ m. **e** The width of apical TJs was measured in ten randomly selected fields in four sections from each gland. The values shown are means \pm SD from four independent experiments. ** P < 0.01 compared with KRH group. ## P < 0.05 compared with KRH group



To confirm the role of CaMKK β in adiponectin-modulated paracellular permeability, we knockdown CaMKK β expression by transfection with CaMKK β shRNA and then rescued its expression by CaMKK β cDNA into SMG-C6 cells. As shown in Fig. 7a, the expression of CaMKK β protein was decreased in CaMKK β shRNA-transfected cells, and increased in CaMKK β cDNA-transfected cells. As shown in Fig. 7b, transfection with CaMKK β shRNA did not affect the basal TER value. fAd-decreased TER values

(Fig. 7c) were abolished in CaMKK β -knockdown cells (Fig. 7d). Furthermore, fAd-mediated decrease in TER values was reappeared in CaMKK β rescued cells (Fig. 7e). Paracellular permeability assay was also performed using FITC-dextran, and the results showed that knockdown of CaMKK β abolished and rescue of CaMKK β reappeared fAd-induced Papp increase (Fig. 8a–e). These results indicate that adiponectin increases paracellular permeability in a CaMKK β -dependent manner.

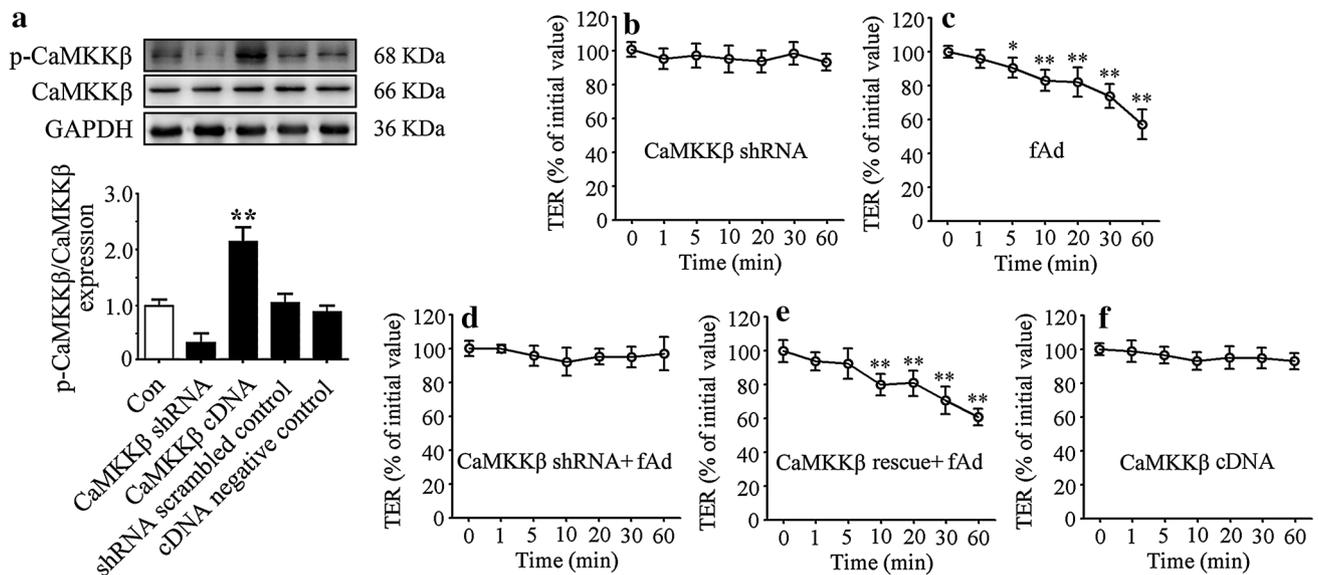
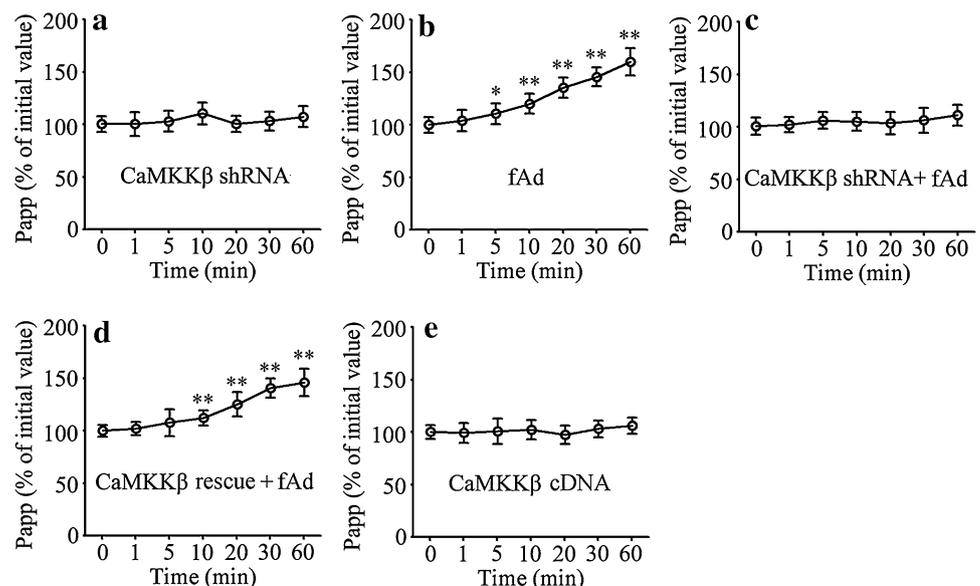


Fig. 7 CaMKK β is required for adiponectin-decreased TER values. **a** Expressions of p-CaMKK β and CaMKK β in SMG-C6 cells transfected with CaMKK β shRNA and CaMKK β cDNA. ** P < 0.01 compared with untransfected control cells. The basal TER value in SMG-C6 cells transfected with CaMKK β shRNA alone (**b**), in cells

incubated with 1 μ g/ml fAd for 60 min (**c**), transfected with CaMKK β shRNA, and then fAd for 60 min (**d**), rescued with CaMKK β cDNA, and then fAd for 60 min (**e**), CaMKK β cDNA alone (**f**). The values shown are means \pm SD from four independent experiments. * P < 0.05 and ** P < 0.01 compared with untreated control cells

Fig. 8 CaMKK β is required for adiponectin-increased permeability of FITC-dextran. The basal Papp value in SMG-C6 cells transfected with CaMKK β shRNA alone (**a**), in cells incubated with 1 μ g/ml fAd for 60 min (**b**), transfected with CaMKK β shRNA, and then fAd for 60 min (**c**), rescued with CaMKK β cDNA, and then fAd for 60 min (**d**), CaMKK β cDNA alone (**e**). The values shown are means \pm SD from four independent experiments. * P < 0.05 and ** P < 0.01 compared with untreated control cells



Discussion

In the present study, we demonstrated that adiponectin increased Ca^{2+} modulation in rat submandibular glands acinar cells. Both extracellular Ca^{2+} influx and release of Ca^{2+} from endoplasmic reticulum were involved in adiponectin-induced salivary secretion, activation of AMPK, and increase in paracellular permeability. Moreover, CaMKK β activated by increased $[\text{Ca}^{2+}]_i$, but not by LKB1, acted as an upstream kinase for adiponectin-mediated AMPK activation. Inhibition of CaMKK β by pharmacological reagent or CaMKK β shRNA abolished, whereas CaMKK β re-expression retained, the increase in TJ width and paracellular permeability induced by fAd. These results reveal that the Ca^{2+} -CaMKK β pathway is responsible for adiponectin-induced salivary secretion via activation of AMPK and increase in paracellular permeability in rat submandibular glands.

In rat pituitary cells, adiponectin increases $[\text{Ca}^{2+}]_i$ through both the influx of extracellular Ca^{2+} and the release of Ca^{2+} from endoplasmic reticulum resulting in the secretion of growth hormone (Steyn et al. 2009). In C2C12 myocytes and *Xenopus* oocytes, adiponectin induced extracellular Ca^{2+} influx via adiponectin receptor 1. Inhibition of Ca^{2+} influx is associated with insulin resistance and reduced exercise tolerance due to the decreased adiponectin levels (Iwabu et al. 2010). In addition, adiponectin-induced increased Ca^{2+} influx is also involved in adrenal cortical hormone release in pituitary cells (Chen et al. 2014). In human and rabbit submandibular glands, activation of mAChRs and transient receptor potential vanilloid subtype 1 increase salivary secretion through elevated $[\text{Ca}^{2+}]_i$ (Ding et al. 2014; Zhang et al. 2006). Moreover, in human transplanted epiphora submandibular glands, the increased $[\text{Ca}^{2+}]_i$ mobilization induced by mAChR activation contributes to hyperscretion (Ding et al. 2014). However, adiponectin-promoted salivary secretion in rat submandibular glands is independent of cholinergic manner (Ding et al. 2013). In the present study, we found that adiponectin could induce increase in $[\text{Ca}^{2+}]_i$ in isolated rat submandibular gland acinar cells. Either EGTA or TG did not affect the adiponectin-induced salivary secretion, however, co-inhibition of extracellular Ca^{2+} influx and intracellular Ca^{2+} release from endoplasmic reticulum completely abolished adiponectin-induced salivary secretion. These results identify that the elevated $[\text{Ca}^{2+}]_i$, derived from both extracellular Ca^{2+} influx and intracellular Ca^{2+} release, is required for the adiponectin-induced salivary secretion in submandibular glands.

The secretion of saliva across the salivary epithelium can be accomplished via either aquaporin 5-based transcellular route or TJ-based paracellular route (Kawedia

et al. 2007). We previously demonstrated that adiponectin-induced salivary secretion in rat submandibular glands involves modulation of paracellular permeability through TJs, but not mediated by aquaporin 5 (Ding et al. 2013). TJs, which consist of a narrow belt-like structure at the most apical portion of the lateral membranes, serve as an indispensable gate for the transport of materials through paracellular pathway (Tsukita et al. 2001). Secretory stimulators, such as carbachol and capsaicin are reported to induce salivation by altering TJ properties and increasing paracellular permeability (Cong et al. 2013, 2015; Ding et al. 2013, 2017; Yang et al. 2017). Moreover, proinflammatory cytokines change TJ content and disrupt TJ barrier function, which might be the mechanism for the hyposecretion that occurs in Sjögren's syndrome (Abe et al. 2016; Baker et al. 2008; Ewert et al. 2010; Mei et al. 2015; Zhang et al. 2016). In rat submandibular glands, adiponectin increases TJ width and paracellular permeability (Ding et al. 2013). Furthermore, AMPK activation regulates the content and distribution of TJ components, resulting in an increased paracellular permeability and salivary secretion (Xiang et al. 2014). Here, inhibition of either extracellular Ca^{2+} influx or Ca^{2+} released from endoplasmic reticulum did not affect adiponectin-induced increase in TJ width in isolated rat submandibular glands. However, co-inhibition of Ca^{2+} influx and Ca^{2+} release from endoplasmic reticulum abolished the adiponectin-induced increased TJ width. Furthermore, in SMG-C6 cells, EGTA combination with TG abolished increased paracellular permeability induced by adiponectin. These results indicate that both Ca^{2+} influx and Ca^{2+} release from endoplasmic reticulum participate in adiponectin-induced salivary secretion via regulating TJ ultrastructure and function.

AMPK is controlled by two upstream kinases, LKB1 and CaMKK β (Hawley et al. 2005; Woods et al. 2003). In Alzheimer's disease, the CaMKK β -AMPK pathway plays a major role in mediating the early synaptotoxic effects of amyloid- β 1-42 oligomers both in vitro and in vivo, serving as a potential therapeutic target for Alzheimer's disease (Mairet-Coello et al. 2013). In human pancreatic tumor cells, 2-deoxyglucose-induced endoplasmic reticulum stress results in an increased Ca^{2+} leakage from endoplasmic reticulum, which subsequently activates AMPK via CaMKK β and ultimately leads to cell autophagy (Xi et al. 2013). In HeLa cells, the increased $[\text{Ca}^{2+}]_i$ activates CaMKK β -AMPK pathway, which participates in the protective effect of *baicalin* against the development of hepatic steatosis and obesity in vivo (Ma et al. 2012). However, whether LKB1 and/or CaMKK β are involved in adiponectin-mediated AMPK activation in submandibular glands is not determined. Here, we found that adiponectin induced AMPK activation by increasing phosphorylation of CaMKK β , but not LKB1 phosphorylation, suggesting that CaMKK β , but not LKB1, is the

upstream kinase for adiponectin-mediated AMPK activation in rat submandibular glands. Furthermore, co-inhibition of Ca^{2+} influx and Ca^{2+} release from endoplasmic reticulum abolished adiponectin-induced $\text{CaMKK}\beta$ and AMPK activation and salivary secretion. These results suggest that Ca^{2+} - $\text{CaMKK}\beta$ -AMPK signaling pathway is involved in adiponectin-modulated salivary secretion. In SMG-C6 cells, $\text{CaMKK}\beta$ knockdown did not alter basal TER or Papp values, suggesting that the maintenance of basal paracellular permeability does not require $\text{CaMKK}\beta$ involvement. However, increased paracellular permeability induced by adiponectin was inhibited by $\text{CaMKK}\beta$ -knockdown, whereas re-expression of $\text{CaMKK}\beta$ rescued the effect of adiponectin. These results suggest that Ca^{2+} - $\text{CaMKK}\beta$ pathway is involved in adiponectin-modulated paracellular permeability in rat submandibular glands.

In summary, our results provide evidence that Ca^{2+} - and $\text{CaMKK}\beta$ -mediated AMPK activation is required for adiponectin-induced salivary secretion. Our findings will enrich our understanding of the secretory mechanisms that link adiponectin to saliva secretion by regulating Ca^{2+} modulation and paracellular permeability.

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Author contributions CD performed the major experiments and wrote the manuscript. ZHD and SLL participated in data interpretation and manuscript improvement. LLW and GYY designed the study, analyzed the data, and wrote the manuscript. All authors read and approved the final manuscript.

Compliance with ethical standards

Conflict of interest The authors declare that they have no competing interests.

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