Roles of PKC and phospho-adducin in transepithelial fluid secretion by Malpighian tubules of the yellow fever mosquito

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The diuretic hormone aedeskinin-III is known to increase the paracellular Cl⁻ conductance in Malpighian (renal) tubules of the mosquito Aedes aegypti via a G protein-coupled receptor. The increase serves the blood-meal-initiated diuresis and is associated with elevated levels of Ca²⁺ and phosphorylated adducin in the cytosol of tubule. In the present study we have cloned adducin in Aedes Malpighian tubules and investigated its physiological roles. Immunolabeling experiments are consistent with the association of adducin with the cortical cytoskeleton, especially near the apical brush border of the tubule. An antibody against phosphorylated adducin revealed the transient phosphorylation of adducin 2 min after stimulating tubules with aedeskinin-III. The PKC inhibitor bisindolyImaleimide-I blocked the phosphorylation of adducin as well as the electrophysiological and diuretic effects of aedeskinin-III. Bisindolylmaleimide-I also inhibited fluid secretion in control tubules. Phorbol 12-myristate 13-acetate increased phosphorylated adducin levels in Malpighian tubules, but it inhibited fluid secretion. Thus, the phosphorylation of adducin by PKC alone is insufficient to trigger diuretic rates of fluid secretion; elevated levels of intracellular Ca²⁺ may also be required. The above results suggest that the phosphorylation of adducin, which is known to destabilize the cytoskeleton, may (1) facilitate the traffic of transporters into the apical brush border supporting diuretic rates of cation secretion and (2) destabilize proteins in the septate junction thereby enabling paracellular anion (CI⁻) secretion at diuretic rates. Moreover, PKC and the phosphorylation of adducin play a central role in control and diuretic tubules, consistent with the dynamic behavior of both transcellular and paracellular transport pathways.

Introduction

Insect kinins are fastacting diuretic hormones that increase electrolyte and fluid secretion in renal (Malpighian) tubules of insects.¹⁻⁴ The kinins trigger the excretion of excess electrolytes and water which insects ingest, gorging on the blood of vertebrates or the sap of plants.^{5,6} In Malpighian tubules of the yellow fever mosquito *Aedes aegypti*, aedeskinin-III (AK-III) short-circuits the epithelium with switchlike speed by increasing the paracellular Cl⁻ conductance.^{7,8} The increase accelerates the transepithelial secretion of KCl and NaCl and consequently water.⁹⁻¹¹

It is known that kinins bind to G protein-coupled receptors in the plasma membrane of principal cells and stellate cells, leading to elevated levels of $Ins(1,4,5)P_3$ presumably via elevated phospholipase C activity.¹¹⁻¹⁶ In turn, the elevated levels of $Ins(1,4,5)P_3$ raise intracellular Ca²⁺ concentrations thereby depleting intracellular Ca²⁺ stores. Store depletion triggers the opening of plasma membrane L-type Ca²⁺ channels which is the essential step in signal transduction.¹⁷ In the absence of extracellular Ca²⁺, the physiological effects of kinins are markedly blunted and transient in *Aedes* Malpighian tubules.¹⁷ How extracellular Ca²⁺ entering cells brings about the increase in paracellular Cl⁻ conductance remains unknown, but the switch like response to kinin diuretic peptides indicates a post translational mechanism.^{7,8}

In an attempt to identify the proteins of the AK-III signaling pathway in *Aedes* Malpighian tubules, we have analyzed the cytosolic proteome of *Aedes* Malpighian tubules before and after treatment with AK-III for only 1 min.¹⁸ In this previous study we observed prominent changes to the cytosolic abundance and phosphorylation state of proteins associated with the cytoskeleton. Of interest in the present study was the significant increase in cytosolic adducin in phosphorylated form after stimulating Malpighian tubules with AK-III.

Adducin was first observed as a 200 kDa protein of the spectrin cytoskeleton of red blood cells and characterized as a calmodulin-binding protein.¹⁹ It is associated with regions of cell-cell contact in other cells.²⁰ Adducin promotes the binding of spectrin to actin, it binds actin, and it bundles actin filaments.

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Figure 1. Aedes adducin gene. (**A**) map of Aedes adducin gene as predicted by the Aedes genome.⁶⁹ Numbered, vertical green bars are exons, horizontal black lines are introns. For more detail see **Table 2**. (**B**) fractured exon map of adducin cDNAs cloned from Aedes Malpighian tubules. Numbers refer to the exons in panel A. Orange and blue bars correspond respectively to the 5' and 3' untranslated regions which flank the open reading frame (gray and black). Vertical red bars indicate poly-adenylation sites. Note the presence of exon 7 in AeAdd-B and its absence in AeAdd-A.

Binding to actin and spectrin stabilizes the spectrin cytoskeleton.²¹⁻²⁴ Significantly, adducin is strongly expressed along the lateral membranes of epithelial cells where it stabilizes epithelial junctions.²⁵⁻²⁷

The goal of the present study was to clone the cDNAs encoding adducins in Malpighian tubules of Aedes aegypti and to elucidate the function of the corresponding proteins in the tubule. We identified two splice variants of the adducin gene and found adducin localized primarily to the subapical region of principal cells. Treating isolated Malpighian tubules with AK-III caused a transient increase in the phosphorylation of the COOH-terminal MARCKS domain of adducin in a time course that parallels the electrophysiological effects of AK-III on Malpighian tubules. The PKC agonist, phorbol myristate acetate (PMA), increased the abundance of phosphorylated adducin (phospho-adducin) in isolated Malpighian tubules, whereas the PKC antagonists staurosporine and bisindolylmaleimide-I decreased the abundance of phospho-adducin. Bisindolylmaleimide-I also blocked the effect of AK-III on (1) tubule electrophysiology and (2) the stimulation of fluid secretion in isolated Malpighian tubules. Thus, PKC and adducin are key mediators of the diuresis triggered by AK-III in Aedes Malpighian tubules.

Results

Molecular cloning of adducin transcripts. The *Aedes* genome contains a single gene that encodes a putative adducin, AAEL011105 (www.vectorbase.org). The gene consists of 13 predicted exons distributed along 50 kb of Supercontig 1.541 at the nucleotide positions 304004-253709 (Fig. 1A). The exact genomic position of each exon is listed in Table 2. As shown in Figure 1B, our RT-PCR studies of *Aedes* Malpighian tubules

detected the expression of two distinct adducin cDNAs derived from gene AAEL011105 that we designate as *Ae*Add-A and *Ae*Add-B. The transcripts are identical except for the presence of the 24 bp of exon 7 in *Ae*Add-B (**Fig. 1B**). The length of the 3'untranslated region (UTR) in each splice variant is highly variable (red vertical bars in **Fig. 1B**) which may reflect the several poly-adenylation sites within exon 13.

The nucleotide sequences of the adducin cDNAs cloned in our laboratory are identical to the corresponding regions of the *Aedes* genome with one notable exception. According to the genome, residue 276,127 (in exon 4) is "G," but in 18 of our 25 sequenced RT-PCR products this residue is "A" (the other 7 products indicated "G"). The identity of this residue affects the coding of the 333rd amino acid of the adducin

protein. A Gly³³³ results when the residue is "G," whereas Ser³³³ results when the residue is "A." Given our sequencing results, we presume that the majority of the adducin cDNAs encode Ser³³³ rather than Gly³³³.

Amino acid sequence of *Aedes* adducin. The *Ae*Add-A transcript, which lacks exon 7, encodes a protein of 710 amino acids (78.7 kDa), and the *Ae*Add-B transcript, which includes exon 7, encodes a protein of 718 amino acids (79.6 kDa). The 8 amino acids encoded by exon 7 are part of the so-called "neck" domain of the protein (red box, Fig. 2). The *Aedes* adducins share with other adducins a putative MARCKS domain in the COOH terminus of the protein (blue box, Fig. 2). The MARCKS domain includes a highly-conserved serine (green oval, Fig. 2) that is known to be phosphorylated by PKC and PKA.³⁷⁻³⁹

Figure 3 illustrates the phylogenetic relationship between the amino-acid sequence of AeAdd-B and those of adducins from other organisms. In brief, AeAdd-B is most closely related to the adducin of Drosophila (DrAdd) and is part of the larger branch that includes adducins from other invertebrates (Caenorhabditis elegans, CaAdd; Schistosoma mansoni, ScAdd). The adducins of humans (HoAdd) cluster in an independent branch of the tree.

Adducin expression and localization in the Malpighian tubules. Immunoblots were performed to characterize the expression of adducin protein in Malpighian tubules of adult female *Aedes* mosquitoes. For the sake of comparison we also examined the expression of adducin immunoreactivity in the midgut. Crude lysates of both tissues yield a protein band of ~100 kDa that exhibits adducin immunoreactivity (Fig. 4). Although the band is larger than the expected size of adducin (~79 kDa) based on the cloned cDNAs, it is known that adducins run slightly higher than expected on SDS-PAGE because of the highlycharged COOH-terminal MARCKS domain.^{29,40}



Figure 2. Aedes adducin. (**A**) Amino-acid sequence alignment of AeAdd-A and AeAdd-B. The red box highlights the eight amino acids present in AeAdd-B that are encoded by exon 7. The blue box highlights the C-terminal MARCKS domain with a serine (green oval) that is predicted to be phosphorylated. (**B**) Adducin protein with phosphorylation sites modeled after mammalian β -adducin.⁷⁰

A protein band larger than 250 kDa also exhibited adducin immunoreactivity in the Malpighian tubules and midgut (Fig. 4). It is likely that this band represents nondenatured adducin in complex with other cytoskeletal elements, such as spectrin, which is a protein of ~280 kDa in *Aedes aegypti* (www.vectorbase.org). Significantly, the antibody against phospho-adducin detected only the ~100 kDa band of adducin (Fig. 4).

Immunolabeling of sections of paraffin embedded *Aedes* Malpighian tubules revealed strong adducin immunoreactivity along the base of the brush border in principal cells (Fig. 5). Weak adducin immunoreactivity was observed near the basal membrane of principal cells consistent with the presence of adducin in the cortical cytoskeleton. Immunoreactivity was diffuse in the cytoplasm of principal cells. Immunolabeling of stellate cells was also observed, but a precise localization was not possible in view of the small size of these cells.

Time-dependent changes in phospho-adducin. In a previous proteomic study we observed that adducin appears in the cytosol of Malpighian tubules in phosphorylated form after treatment with the diuretic peptide AK-III for only 2 min.¹⁸ It was therefore

of interest to examine the time course of the adducin phosphorylation in western blots of tubule lysates. As shown in the representative western blot of **Figure 6A**, the strongest immunoreactivity to phospho-adducin was observed 2 min after adding AK-III (10^{-7} M) to the peritubular medium of Malpighian tubules. Thereafter, the immunoreactivity to phospho-adducin progressively diminished in spite of the presence of AK-III. Notably, the immunoreactivity to total adducin did not change significantly with time (**Fig. 6A**).

Figure 6B summarizes the effects of AK-III on the immunoreactivity of phospho-adducin normalized to that of total adducin in four separate trials. Again, after adding AK-III to the peritubular bath of tubules, phospho-adducin peaked 2 min. After a 10 min exposure to AK-III, phospho-adducin levels are not significantly different from control tubules and after a 20 min exposure to AK-III, phospho-adducin levels have clearly returned to control levels. Thus, the AK-III mediated phosphorylation of adducin is transient.

The phosphorylation of *Aedes* adducin by protein kinase C. The signal transduction of AK-III includes the essential role



Figure 3. Neighbor-joining tree of adducin amino-acid sequences from *Aedes aegypti (Ae), Drosophila melanogaster (Dr), Caenorhabditis elegans (Ca), Schistosoma mansoni (Sc),* and *Homo sapiens (Ho).* The tree is rooted to an adducin-like protein of *Dictyostelium discoideum (Di)*. Filled circles indicate the nodes of branches for which bootstrap scores are provided (from 1,000 replicates). The total branch length between two proteins represents the proportion of amino acids that differ between them. The scale bar corresponds to a proportional difference (branch length) of 0.1 (i.e., a 10% difference in amino acids). The tree was constructed with MEGA 4 software⁷¹ using Poisson corrected distance estimates. Accession numbers are as follows: *Ae*Add, F705874; *Dr*Add, NP_001188977; *Ca*Add, AAD49856; *ScAdd*, XP_002578303; *Ho*Add α , NP_001110; *Ho*Add β , NP_001608; *Ho*Add γ , NP_058432; *Di*Add, XP_640404.



Figure 4. Representative western blots of the immunoreactivity to adducin and phospho-adducin in lysates of *Aedes* midgut and Malpighian tubule. Numbers correspond to molecular weight markers in kDa.

of Ca²⁺ as second messenger.^{11,17,41} For this reason we explored the role of protein kinase C (PKC) in the phosphorylation of adducin. Furthermore, PKC is known to phosphorylate Ser⁷²⁶ in the MARCKS domain of human α adducin,^{22,38,39} and the

homologous Ser is present in *Aedes* adducin (Fig. 2). Isolated Malpighian tubules were treated with known stimulators and inhibitors of PKC. Lysates of the tubules were then examined for phospho-adducin immunoreactivity in western blots. Equal protein loading was verified in this series of experiments with an antibody to β -tubulin (E7) because we had exhausted our supply of the adducin antibody.

Phorbol myristate acetate (PMA) is an activator of PKC.^{42,43} Isolated Malpighian tubules were treated with 10⁻⁶ M PMA for 20 min to allow sufficient time for the entry of PMA into the cells of the tubule. As shown in **Figure 7A**, we consistently observed higher levels of phospho-adducin after treating tubules with PMA. Standardized to β -tubulin, the increase in phosphoadducin immunoreactivity is statistically significant (p < 0.05, **Fig. 7A**).

Next we investigated the effects of PKC inhibitors on the phosphorylation of adducin. One group of Malpighian tubules served as the control group, a second group was treated with AK-III (10^{-7} M) for 2 min, and the third group of tubules was preincubated with the PKC inhibitor of interest for 20 min before they were treated with AK-III. The treatment of Malpighian tubules with AK-III for 2 min significantly increased the phospho-adducin immunoreactivity in all tubules studied (**Fig.** 7**B**–**D**). However, when the tubules were first incubated with staurosporine (10^{-7} M) or bisindolylmaleimide-I (BIM-1, 10^{-5} M) for 20 min, treatment with AK-III failed to stimulate the phosphorylation of adducin (**Fig.** 7**B** and **C**). In contrast, the pre-incubation of Malpighian tubules with chelerythrine (5×10^{-6} M) did not inhibit the AK-III mediated phosphorylation of adducin (**Fig.** 7**D**).

Physiological studies in intact Malpighian tubules. To evaluate the effects of PKC activators and inhibitors on the physiological performance of Malpighian tubules we conducted two-electrode voltage clamping experiments and fluid secretion assays in isolated Malpighian tubules. The effect of the PKC activator phorbol myristate acetate (PMA) was of interest first. In the typical experiment a Malpighian tubule was bathed in Ringer solution, and a principal cell of the tubule was impaled with voltage and current electrodes for the measurement of the basolateral membrane voltage and the input resistance of the cell. Before applying PMA, the tubule was prepulsed with AK-III to determine its responsiveness to kinin stimulation.⁹

As shown in the representative experiment of Figure 8A, the addition of AK-III (10^{-7} M) to the peritubular bath immediately hyperpolarized the basolateral membrane voltage (V_{bl}) from an oscillating voltage in the vicinity of -60 mV to a stable voltage of -88 mV. In parallel, the input resistance of the principal cell (R_{in}) dropped from 456 to 358 k Ω . The sudden hyperpolarization of V_{bl} together with the decrease of R_{in} reflect the well-known post-translational diuretic effects of AK-III in *Aedes* Malpighian tubules which include (1) the increase in the paracellular Cl⁻ conductance and (2) the activation of Ca²⁺ channels in the basolateral membrane of principal cells.^{8,17} The washout of AK-III returned the tubule to oscillating membrane voltages and previous cell input resistances. The subsequent addition of PMA (10^{-6} M) to the peritubular medium caused a slow

depolarization of V_{bl} together with an increase in R_{in} (Fig. 8A). In 13 tubule experiments, V_{bl} significantly (p < 0.02) depolarized from -69.5 \pm 1.8 mV to -59.3 ± 2.4 mV after an average of 10 min in the presence of 10⁻⁶ M PMA. At the same time the cell input resistance R_{in} significantly increased from 439.7 \pm 18.4 k Ω to 557.9 \pm 19.3 k Ω (p < 0.05). The gradual change of V_{μ} and R_{in} suggest the gradual decline in the rate of transepithelial ion secretion. Nevertheless, the tubule still responded to AK-III in the presence of PMA. After a more than 20 min exposure to PMA, the addition of AK-III reversibly lead to the prompt hyperpolarization of V_{bl} from -65 to -86 mV in parallel with the reduction of R_{in} from 455 to 405 k (Fig. 8A). Thus, the gradual electrophysiological changes in the presence of PMA did not preclude the usual electrophysiological response of the tubule to AK-III.

In 13 Malpighian tubules studied by the method of Ramsay, the addition of PMA (10^{-6} M) to the peritubular Ringer bath significantly (p < 0.02) reduced the rate of fluid secretion from 0.92 ± 0.06 nl/min to 0.33 ± 0.03 nl/ min (**Fig. 9A**).

The effects of the PKC inhibitor bisindolylmaleimide-I was of interest next. Figure 8B illustrates a typical experiment that shows the electrophysiological effects of BIM-I. Again, it was important to first confirm that the tubule responded to the diuretic peptide AK-III. The addition of AK-III (10^{-6} M) to the peritubular bath hyperpolarized V_{bl} from -80 to -100 mV in parallel with the drop of R_{in} from 323 to 278 k Ω (Fig. 8B). The hyperpolarization of V_{bl} together with the reduction of R_{in} is consistently observed when tubules do respond to aedeskinins. In a previous study, AK-III significantly (p < 0.001) hyperpolarized

Α subapical brush border nucleus, principal cell basolateral membrane tubule stellate cell lumen 50 µm В

Figure 5. Representative immunolocalization of adducin in consecutive sections of a Malpighian tubule of *Aedes aegypti*. (**A**) Adducin immunoreactivity is indicated by the red staining. Nuclei are stained blue by hematoxylin. (**B**) Negative control. Tubules undergoing the same adducin staining procedure but without antibody served as negative control.

 V_{bl} from -64.3 to -87.8 mV in parallel with the significant (p < 0.01) reduction of R_{in} from 343.3 to 265.2 k Ω .⁹ The washout of AK-III recovered prestimulation values (Fig. 8B). Subsequently, the addition of BIM-I (10⁻⁵ M) to the peritubular bath triggered the gradual decay of V_{bl} together with a large increase of R_{in} . By the time V_{bl} had decayed to -23 mV, R_{in} had increased to 872 k Ω (Fig. 8B). In the presence of BIM-I, AK-III (10⁻⁶ M) had no significant effect on V_{bl} and R_{in} (Fig. 8B). Thus, BIM-I blocks the

electrophysiological effects of AK-III as it blocks the phosphorylation of adducin (**Fig.** 7). The washout of both AK-III and BIM-I repolarized V_{bl} to -56 mV and decreased R_{in} to 483 k Ω . In nine tubules experiments, BIM-I significantly (p < 0.01) depolarized V_{bl} from -63.6 ± 6.0 mV to -42.7 ± 4.6 mV while significantly (p < 0.01) increasing R_{in} from 418.1 ± 53.1 k Ω to 739.0 ± 118.9 k Ω . In the presence of BIMI, the effects of AK-III on V_{bl} and R_{in} were consistently blocked.



Figure 6. Transient phosphorylation of adducin in the presence of the diuretic peptide AK-III (10⁻⁷ M) in isolated Malpighian tubules of *Aedes aegypti*. (**A**) Representative western blots of phospho-adducin and adducin in Malpighian tubules that were incubated without (time 0) or with AK-III for the times indicated. (**B**) Phospho-adducin immunoreactivity standardized to total adducin immunoreactivity determined in 4 independent western blots. Values are mean \pm SE; **p < 0.01, *p < 0.05, one-way ANOVA with a Newman-Keuls post-test.

BIM-I also reduced rates of fluid secretion in isolated Malpighian tubules. In seven tubule experiments, the addition of BIM-I (10⁻⁵ M) to the peritubular bath of unstimulated, control Malpighian tubules significantly (p < 0.05) reduced the spontaneous rate of fluid secretion from 0.48 ± 0.07 nl/min to 0.17 ± 0.04 nl/min. (**Fig. 9B**). Tubules pretreated with BIM-I for 30 min failed to increase the rate of fluid secretion after the addition of AK-III to the peritubular bath (**Fig. 9C**). Instead, the average rate of fluid secretion tended to decrease upon the addition of AK-III from 0.44 ± 0.14 nl/min to 0.24 ± 0.11 nl/min in the presence of BIMI, though not significantly. The usual response of tubules to AK-III is the significant (p < 0.001) increase of the fluid secretion rate from, for example, 0.52 ± 0.06 nl/min to 1.04 ± 0.13 nl/min in 21 tubules (Schepel et al. 2010). Thus, the pre-incubation with BIM-I blocked the diuretic effects of AK-III.

Discussion

Adducin in Malpighian tubules of *Aedes aegypti*. We have cloned two alternativelyspliced variants of adducin cDNAs from *Aedes* Malpighian tubules that differ by the presence or absence of exon 7 (Fig. 1B). The 3'untranslated region (UTR) of these cDNAs, which is encoded by exon 13, is characterized by variable lengths. Different lengths of the 3'UTR region in the adducin of Drosophila ovaries are thought to influence the trafficking of adducin transcripts within the cell according to specific localization signals found in the UTR sequence.⁴⁴ Thus, differences in the 3'UTR length of *Ae*Add transcripts in Malpighian tubules may direct the posttranscriptional trafficking of adducin mRNA.

As shown in Figure 2B, adducin proteins form three broad structural/functional regions: a globular NH_2 -terminal head domain, a neck domain, and a COOH-terminal tail.^{22,40} The NH_2 -terminal head region of *Aedes* adducin contains an aldolase II superfamily domain similar to human and Drosophila adducin. The head region has been implicated in the interaction of adducin with clathrin coated vesicles through a motif that is present in *Ae*Add-A and *Ae*Add-B (Fig. 2). Thus, *Aedes* adducin may be involved in endocytosis.^{45,46}

The neck region in mammalian adducin is necessary for recruiting spectrin and actin²² and for forming heterooligomers of adducin isoforms.²¹ Exon 7 in *Ae*Add-B encodes the 8 amino-acid residues "WIDANVDE" in the neck domain of the mosquito protein. Thus, the alternative splicing of exon 7 may result in adducins with differential abilities to recruit spectrin and actin and/or to form oligomers.

The COOH-terminal tail of adducin is thought to directly interact with actin and spectrin thereby regulating the assembly of the spectrin/actin cytoskeleton. The tail includes the putative calmodulin binding site as well as the MARCKS domain (**Fig. 2**). The latter is the target of protein kinase C (PKC) and protein kinase A (PKA) for regulating the activity of adducin.^{37,39} The MARCKS domain is well conserved among *Aedes*, Drosophila and Homo and includes the serine residue (red box and red highlighted Ser

in Fig. 2). The phosphorylation of the MARCKS domain causes adducin to dissociate from spectrin and actin, promoting the disassembly of the spectrin cytoskeleton. As a result, proteins of tight and adherens junctions may change conformation, position, or be internalized.^{38,39,47,48}

Immunolocalization of adducin in Aedes Malpighian tubules. In histological sections of Aedes Malpighian tubules, adducin immunoreactivity is observed in both principal and stellate cells (Fig. 5). Notably, prominent immunolabeling occurs along the base of the apical brush border of principal cells, which is strikingly similar to the localization of actin and β spectrin in Aedes Malpighian tubules.⁴⁹ The presence of these three proteins at the apical base of principal cells indicates an actin/spectrin cytoskeleton that may serve the structure and function of the tall brush border extending into the lumen of the tubule.3 Since adducin caps F-actin and recruits spectrin to actin,²² adducin may stabilize actin filaments extending into the microvilli of principal cells (Fig. 4). Moreover, the rapid increase in phospho-adducin immunoreactivity after stimulation with the diuretic peptide AK-III (Fig. 6) may reflect the destabilization of the apical cytoskeleton, thereby allowing traffic into and out of microvilli.^{46,50,51} Bradley and Satir have observed in Rhodnius Malpighian tubules that mitochondria move from a position below the cell cortex to one inside the microvilli within 10 min after stimulating tubules with the diuretic hormone serotonin.52,53 The progressive loss of phospho-adducin 10 min after AK-IIIstimulation (Fig. 6) suggests similar cytoskeletal dynamics in Aedes Malpighian tubules and reflect the re-assembly of the cytoskeleton after solute transporters and/or mitochondria have moved into the brush border of Aedes Malpighian tubules.

Physiology of adducin. The present study was prompted by our previous proteomic study that indicated modifications to the cytoskeleton as one mechanism for regulating the rate of electrolyte and fluid secretion in Malpighian tubules.¹⁸ The cytoskeletal protein adducin, actin and actin depolymerizing factor appeared in the cytosol of Malpighian tubules at elevated levels after stimulating Malpighian tubules with aedeskinin-III for only 1 min. A role of protein kinase C was further implicated by the requirement of Ca²⁺ for aedeskinin signaling.^{3,17} Among the first steps in kinin signaling is the activation of Ca²⁺ channels in the basolateral membrane of principal cells.^{17,41}

One function of Ca2+ is the activation of protein kinase C, as documented in electrophysiological studies and in measurements of fluid secretion (Figs. 8 and 9). Malpighian tubules respond to kinin diuretic peptides (aedeskinins and leucokinins) with the hyperpolarization of the basolateral membrane voltage of principal cells. The hyperpolarization reflects the short circuit of the transepithelial voltage as kinins trigger the sudden increase of the paracellular Cl⁻ conductance.^{8,9,17,54} In parallel, the cell input resistance decreases due to the activation of basolateral membrane Ca2+ channels (Fig. 8). These electrophysiological effects are absent when tubules have first been treated with the PKC inhibitor bisindolylmaleimide-I (Fig. 8B). Accordingly, the inhibition of PKC prevents aedeskinin-III from increasing the paracellular Cl⁻ conductance, thereby blocking the key ionic event of the diuresis triggered by kinin diuretic peptides. Studies of fluid secretion confirm this conclusion. Tubules normally respond to aedeskinin or leucokinin by immediately doubling the rate of fluid secretion,^{55,56} but bisindolylmaleimide-I prevents the aedeskinin-mediated increase in fluid secretion (Fig. 9C). These findings implicate PKC as playing an integral role in the increase of the paracellular secretion of Cl- that leads to a corresponding increase in the transepithelial secretion of cations and water.

The pharmacology of the adducin phosphorylation uncovered in the present study illuminates on the type of PKC that is activated by aedeskinin-III. The PKC antagonists staurosporine and bisindolylmaleimide-I block the AK-III induced phosphorylation of adducin, but chelerythrine does not (Fig. 7). Chelerythrine is an inhibitor of the conventional α and β isoforms of PKC by binding to the catalytic domain of kinases.⁵⁷ In contrast, bisindolylmaleimide-I, the most selective inhibitor of conventional α , β , and γ PKC's and staurosporine are structurally similar and block the ATP binding pocket of PKC's.⁵⁸⁻⁶¹ Thus, the PKC that phosphorylates adducin

in Aedes Malpighian tubules may be a variant of the γ PKC.

Our use of an antibody specific to the COOH-terminal MARCKS domain of adducin proves that aedeskinin brings about the phosphorylation of this domain (Figs. 6 and 7). The MARCKS domain of adducin is required for protein-protein interactions with actin and spectrin.^{48,62} Adducin binds to the barbed ends and to the sides of actin filaments thereby enhancing the





association of spectrin with actin filaments that stabilizes the spectrin-actin meshwork.⁶² Upon phosphorylation of the MARCKS domain, phospho-adducin dissociates from spectrin and actin, thereby destabilizing the cytoskeleton. In the present study, the phosphorylation of the MARCKS domain took place within 2 min of AK-III stimulation, confirming the sudden rise in cytosolic phospho-adducin we have observed in our proteomic study of



Figure 8. Representative effects of protein kinase C (PKC) activator and inhibitor on the basolateral membrane voltage (V_{bl}) and the input resistance ($R_{in'}, k\Omega$) of a principal cell in isolated Malpighian tubules of *Aedes aegypti*. The tubules were prepulsed with aedeskinin-III (AK-III, 10⁻⁶ M) to ascertain an active signaling pathway. (**A**) The PKC activator phorbol myristate acetate (PMA, 10⁻⁶ M) had minor effects on V_{bl} and $R_{in'}$. In the presence of PMA the tubule still responded to AK-III. (**B**) The PKC inhibitor bisindolylmaleimide-I (BIMI, 10⁻⁵ M) significantly depolarized V_{bl} and increased $R_{in'}$. In the presence of BIM-I the tubule did not respond to AK-III. PMA and BIM-I were dissolved in DMSO (final bath concentration of 0.1%). Arrows indicate the times $R_{in'}$ was determined.

Aedes Malpighian tubules stimulated with AK-III for only 1 min.¹⁸ Levels of cytosolic phospho-adducin start to return to control levels two min after stimulation with AK-III (**Fig. 6**). The time-dependent changes in the phosphorylation of adducin are consistent with (1) the known switchlike on/off effects of leucokinin and aedeskinin on the paracellular shunt resistance of mosquito Malpighian tubules,^{7,8} (2) the transient actin reorganization of the subapical cytoskeleton in *Aedes* Malpighian tubules after a blood meal⁴⁹ and (3) the switch-like behavior of G proteincoupled receptors.¹²

How the adducin-controlled cytoskeleton may affect the switchlike increase of the paracellular Cl⁻ conductance following treatment with the diuretic hormone AK-III is explored below.^{7,8} Importantly, adducin is found concentrated at sites of cell-cell contact along the lateral membranes of intestinal epithelial cells.²⁵ Here, adducin is a key player in remodeling the tight junction in Ca²⁺- and phosphorylation-dependent ways.^{25,63} Phorbol ester causes adducin to leave epithelial contact sites in keratino-cytes and MDCK cells, consistent with the phosphorylation of adducin by PKC and the destabilization of junctional complexes along the paracellular pathway.²⁵ Similarly, the phosphorylation of adducin in *Aedes* Malpighian tubules may destabilize the cyto-skeleton along the septate junction, thereby increasing the electrical conductance of the paracellular pathway.^{8,13,41}

Taken together, the present study illuminates the post-calcium steps of aedeskinin signaling (Fig. 10). The binding of AK-III to its G protein-coupled receptor triggers the formation of diacyl glycerol (DAG) and the entry of Ca²⁺ into the cell. Calcium binding to PKC frees the catalytic domain of PKC which acquires the property of binding to DAG. The activation of PKC by DAG leads to the phosphorylation of proteins nearby, among them adducin. The phosphorylation of adducin for only a few minutes likely allows changes in the cytoskeleton not only at the base of the brush border, but also at the basal and lateral membranes of epithelial cells. The cytoskeletal dynamics along the brush border may fortify the transport activities of the brush border. The cytoskeletal changes at the basolateral membrane may regulate the trafficking of Ca2+ channels and other cation uptake mechanisms, thereby providing additional substrates for the enhanced transcellular secretion of cations. The cytoskeletal changes along the septate junction may modify the junctional complex extending into the paracellular space with the effect of increasing the paracellular secretion of Cl-.

Role of PKC in spontaneous, basal fluid secretion. After isolation from the mosquito, Malpighian tubules bathed in Ringer solution secrete fluid for hours without stimulation by extracellular agents. Nevertheless, PKC participates in the mechanism for sustaining control, basal fluid secretion rates. In control tubules, the PKC inhibitor BIM-I depolarizes the basolateral membrane voltage of principal cells and increases the cell input resistance (Fig. 8B) consistent with the reduction in electrogenic transepithelial ion secretion (Fig. 8B). In parallel, rates of transepithelial fluid secretion decrease (Fig. 9B). The decrease suggests that as much as 65% of the spontaneous fluid secretion rate is dependent on the activity of PKC.

The spontaneous activity of PKC may also account for the spontaneous oscillations of voltage and resistance observed in control, unstimulated tubules.⁵⁴ The oscillations are dependent on transepithelial Cl- gradients, and they reflect spontaneous fluctuations in the paracellular Cl⁻ conductance.^{54,64} The frequency of these oscillations resembles the frequencies of changes in intracellular free Ca2+ that stem from Ca2+ release and reuptake mechanisms at intracellular Ca2+ stores. Moreover, BIM-I eliminates the oscillations (Fig. 8B), indicating the role of PKC in mediating the spontaneous changes of the paracellular Cl conductance. Supporting this conclusion, we have observed the Ca²⁺ dependence of spontaneous oscillations.¹⁷ Increasing the extracellular Ca²⁺ concentration progressively reduces the frequency and amplitude of oscillations by increasing the duration of high paracellular Cl- conductance.11,17 Moreover, the addition of the Ca²⁺ ionophore A-23187 to the peritubular Ringer of Malpighian tubules eliminates oscillations altogether by inducing a high, steady-state paracellular Cl⁻ conductance similar to that induced by kinin diuretic peptides.⁶⁵ In view of the known activation of PKC by Ca²⁺, it is tempting to conclude that PKC mediates the spontaneous oscillations of the paracellular Cl⁻ conductances via the phosphorylation of adducin and other proteins.

Treating Malpighian tubules with the PKC agonist, PMA, results in the phosphorylation of adducin (Fig. 5). PMA is a phorbol ester that mimics the role of DAG in the activation of conventional PKCs.42,43,66 Although PMA activates PKC and phosphorylate adducin in intact tubules, this effect is insufficient to trigger diuresis. Moreover, PMA brings about the inhibition of fluid secretion, and it depolarizes the basolateral membrane voltage (Fig. 8A; Fig. 9). In parallel, the magnitude of the spontaneous voltage oscillations decreases and the cell input resistance increases. These effects indicate the inhibition of transepithelial electrolyte and fluid secretion beyond the phosphorylation of Aedes adducin. PMA is reported to withdraw the Na/K/2Cl cotransporter from the basolateral membrane of principal cells as in T84 cells⁶⁷ which in *Aedes* Malpighian tubules is expected to inhibit fluid secretion.³⁶ PMA is also reported to target RasGRP and Munc-13 which may bring about the inhibition of fluid secretion in the intact tubule independently of PKC.68

In the presence of PMA, the tubules still respond to stimulation by aedeskinin with the usual marked hyperpolarization of the basolateral membrane voltage together with the reduction in cell input resistance (**Fig. 8A**). These responses present the electrophysiological signature of the kinin diuresis that includes signaling by Ca^{2+} . Accordingly, the full-blown diuresis triggered by kinin diuretic peptides requires not only PKC and the phosphorylation of adducin, but also additional post-translational effects of Ca^{2+} that remain to be elucidated.



Figure 9. The effects activators and inhibitors of protein kinase C (PKC) on the rate of transepithelial fluid secretion in isolated Malpighian tubules of *Aedes aegypti*. In view of the variability in spontaneous (control) rates of fluid secretion in Malpighian tubules from different hatches of mosquitoes, each tubule was used as its own control. Accordingly, the paired Student's t-test was used to evaluate the significance of the difference between the control and experimental periods. (**A**) Aedeskinin-III (AK = III, 10⁻⁶ M) significantly increases the rate of fluid secretion. Data are from Schepel et al.⁷² (**B**) The PKC activator phorbol myristate acetate (PMA, 10⁻⁶ M) significantly reduced transepithelial fluid secretion. (**C**) The PKC inhibitor bisindolyImaleimide-I (BIM-I, 10⁻⁵ M) significantly reduced transepithelial fluid secretion. (**m**) The PKC inhibitor BIM-I blocked the diuretic effects of aedeskinin-III (AK-III). Data are mean ± SE; (n = number of Malpighian tubules tubules); *p < 0.05; **p < 0.02; ***p < 0.001; ns, not significant.

Materials and Methods

Mosquito rearing and isolation of Malpighian tubules. The mosquito colony (*Aedes aegypti*) was maintained as described previously.⁸ Malpighian tubules and midgut were isolated from female mosquitoes and transferred to freshly prepared Ringer solution containing in mM: NaCl 150, KCl 3.4, CaCl₂ 1.7, NaHCO₃ 1.8, MgSO₄ 1, glucose 5, and HEPES 25. The pH of the solution was adjusted to 7.1. For molecular studies, the tubules and midgut were immediately frozen in liquid nitrogen after experimental treatments and stored at -80°C.

Molecular cloning. Malpighian tubule cDNA was generated using a Generacer kit (Invitrogen) as described previously.²⁸ In brief, 150 Malpighian tubules were isolated from 30 adult female *Aedes* mosquitoes and placed in a solution of ice-cold Trizol reagent (Invitrogen). The total RNA was extracted and used as a template to create two independent pools of singlestranded cDNA for the 5' and 3'rapid amplification of cDNA ends (RACE). We refer to these pools, respectively, as the 5'cDNA and 3'cDNA.

The *Aedes aegypti* genome (www.vectorbase.org) was referenced to design the primers Ad-1F and Ad-3R (Table 1) which



Figure 10. Hypothetical model of aedeskinin signaling. Aedeskinin-III (AK-III) binds to the kinin GPCR located on the basolateral membrane of principal and/or stellate cells. GTP- α is released and activates phospholipase C (PLC) to produce inositol-triphosphate [Ins(1,4,5) P_3] and diacyl-glycerol (DAG). Ins(1,4,5) P_3 causes the release of intracellular calcium stores. Store depletion triggers the opening of Ca²⁺ channels in the basolateral membrane.⁷³ Elevated intracellular Ca²⁺ levels target cytosolic protein kinase C (PKC) to the cell membrane where binding to DAG activates the kinase. PKC phosphorylates adducin causing its dissociation from the actin/spectrin cytoskeleton along the paracellular pathway and the apical membrane. The destabilization of the cytoskeleton along the paracellular space thereby increasing the paracellular electrical conductance. Along the apical membrane, the destabilization of the cytoskeleton may allow the traffic of transporters and cell organelles into and out of the brush border.

bind to regions of the predicted openreading frame (ORF) of a putative adducin gene (accession number AAEL011105). The 5'RACE was conducted on 0.5 μ l of 5'cDNA using the following: Generacer 5'primer (Invitrogen), reverse primer Ad-3R (**Table 1**), and Platinum PCR Supermix HF (Invitrogen). The 3' RACE was conducted on 0.5 μ l of 3'cDNA using the following: Generacer 3'primer (Invitrogen), forward primer Ad-1F (**Table 1**) and Platinum PCR Supermix HF (Invitrogen). For both the 5'RACE and 3'RACE, the total reaction volume was 25 μ l, and a touchdown thermocycling protocol was employed as described in the Generacer kit (Invitrogen). All RACE products were run on a 1% agarose gel supplemented with ethidium bromide and visualized with a UV transilluminator.

The RACE products were ligated into TOPO TA cloning plasmids (Invitrogen) and transformed into TOP10 One Shot *E. coli* (Invitrogen) following the manufacturer's protocol. Plasmid DNA was isolated using QIA Spin miniprep kits (Qiagen). The purified plasmid DNA was sequenced by the Cornell University Life Sciences Core Laboratory Centers.

Once the 5' and 3' ends of the adducin cDNAs were sequenced, two primers were designed to amplify a "full-length" cDNA containing the entire ORF (Ad-FL1F and Ad-FL1R in Table 1). These primers were added to a mixture containing 0.5μ l of Malpighian tubule 3'cDNA and Platinum PCR Supermix HF (Invitrogen) which was subjected to the following amplification protocol: one cycle at 94°C for 2 min; 35 cycles at 94°C for 30 s, 65°C for 30 s, and 68°C for 4 min; and one final cycle at 68°C for 10 min. The PCR products were TOPO cloned and sequenced as described above.

The sequencing data from the RACE experiments and full-length PCR were assembled to form consensus sequences. As described in the Results, two different splice variants of *Aedes* adducin were identified in Malpighian tubules: *Ae*Add-A (GenBank accession number F705874) and *Ae*Add-B (GenBank accession number F705875).

Antibodies. To detect adducin immunoreactivity on western blots and in immunohistochemistry, a polyclonal rabbit antibody—affinity purified against human α and β adducins—was used.^{29,30} We refer to this antibody as the "adducin antibody." To detect phosphorylated adducin in western blotting, a polyclonal rabbit antibody raised against the phosphorylated human γ -adducin was purchased from Millipore. We refer to this antibody as the "phosphoadducin antibody." It targets the COOH-terminal MARCKS domain of adducin. A monoclonal mouse antibody raised against β -tubulin (E7) was also purchased for use in western blots (Developmental Studies Hybridoma Bank, University of Iowa).

Western blotting. Crude lysates of Malpighian tubules and midguts isolated from 10 adult female mosquitoes were prepared in a 10-fold dilution of ice-cold Ringer solution supplemented with 50 μ M EDTA, Halt protease inhibitor cocktail (Thermo

Fisher Scientific) and Halt phosphatase inhibitor cocktail (Thermo Total protein content of samples was assessed using a Pierce bicinchoninic acid (BCA) protein assay (Thermo Fisher Scientific). To denature proteins, the samples were supplemented with an appropriate amount of a 5× Laemmli sample buffer and boiled for 5 min.³¹

An 8% acrylamide gel was prepared and each lane was loaded with 25 µg of total protein derived from Malpighian tubules or midguts. The proteins were separated by electrophoresis and then transferred to a polyvinylidene difluoride (PVDF) membrane (Bio-Rad). The PVDF membrane was blocked for 90 min at room temperature in a blocking solution consisting of Tween-Tris-buffered saline (TTBS; 10 mM TRIS-HCl, 150 mM NaCl, 0.01% Tween 20, pH 7.4) and nonfat dry milk powder (5% w/v). The PVDF was then probed with either the adducin antibody or phospho-adducin antibody (diluted 1:1,000 in blocking solution) overnight at 4°C. On the following day the PVDF membrane was washed in TTBS three times (5 min each) and probed with a goat-anti rabbit secondary antibody conjugated with horseradish peroxidase for 90 min at room temperature. After washing in TTBS three more times, the PVDF was placed in Supersignal West Pico Chemiluminescent Substrate (Thermo Fisher Scientific) for 5 min. The luminescent signal

Table 1. Primers used in the cloning of adducin cDNAs

Primer	Use	Primer Sequence	Genomic position
Ad3R	5' RACE	5' CCG GCG TCG AAG AAT GTG TTG GCG AAG CTT 3'	271811271840
Ad1F	3' RACE	5' CCG GCA CAC CAG ACC CGA AGA AAA T 3'	275653275629
AdFL_1F	Full Length	5' GTA GTT GAC GCC GCC GTG AAA AAA CGT TGA 3'	303973303944
AdFL_1R	Full Length	5' ATC GTC GCT GAG TGC TGT TCA TGT TGT GAT 3'	255310255339

The genomic positions that the primers anneal to are indicated and are relative to "Supercontig 1.541" of the *Aedes aegypti* genome (www.vectorbase. org). All primers were synthesized by Integrated DNA Technologies.

emitted from the PVDF membrane was detected with X-ray film in a dark room.

Time course and pharmacology of adducin phosphorylation. In time course studies of the phosphorylation of adducin after stimulating Malpighian tubules with the diuretic peptide aedeskinin-III (AK-III), 200 Malpighian tubules were isolated from 40 adult female mosquitoes and distributed equally in four 1.5 ml microcentrifuge tubes containing 200 μ l Ringer solution. To three of the tubes, AK-III was added to a final concentration of 10⁻⁷ M. After 2, 10, or 20 min incubation with AK-III, the Ringer solution was aspirated and the tubules were snap frozen in liquid nitrogen stored at -80°C. The fourth tube (control) received no AK-III.

For identifying the kinase(s) that phosphorylate adducin in Aedes Malpighian tubules, 50 tubules from 10 adult female mosquitoes were isolated per treatment. To test the effects of an agonist of protein kinase C (PKC), two groups of tubules were incubated with either phorbol 12-myristate 13-acetate (PMA, 10⁻⁶ M in 0.05% DMSO) or the vehicle (0.05% DMSO) for 20 min before freezing the tubules in liquid nitrogen. To test the effects of PKC antagonists on adducin phosphorylation, two groups of tubules were incubated with the vehicle (0.05 or 0.1%) DMSO) and one group was incubated with either chelerythrine (Chel, 5×10^{-6} M), bisindolylmaleimide-I (BIM-I, 10^{-5} M), or staurosporine (Stau, 10-7 M). After a 20 min incubation, AK-III (10^{-7} M) was added to one of the tubes containing the vehicle and the tube containing the antagonist. After 2 min of incubation with AK-III, the Ringer solution was aspirated and the tubules were frozen in liquid nitrogen as described above.

The frozen tubules were prepared for western blotting experiments as described above. On a given PVDF, the phosphoadducin immunoreactivity was detected first. After detection, the PVDF was stripped of the phospho-adducin antibody using Restore PLUS western blot stripping buffer or Restore western blot stripping buffer (Thermo Fisher Scientific) for 10 min at room temperature. After confirming the removal of the phosphoadducin antibody, the PVDF membrane was incubated with the adducin antibody to normalize the phospho-adducin signal. In some experiments, normalizing the phospho-adducin signal to that of total adducin was not possible, because the supply of the adducin antibody had been depleted. In these cases it was necessary to use an anti β -tubulin antibody (E7) to normalize levels of phospho-adducin immunoreactivity.

The resulting X-ray films from the phosphorylation experiments were digitized with a scanner and the optical densities of

Table 2. Genomic locations	of the exons	of the Aedes	adducin gene
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Exon	Start Position	End Position	Exon Length (base pairs)
1	304004	303945	60
2	280723	280135	589
3	276606	276439	168
4	276376	276078	299
5	276020	275821	200
6	275745	275622	124
7	272652	272629	24
8	271850	271737	114
9	269803	269518	286
10	269439	269329	111
11	269266	269207	60
12	259176	259109	68
13	255447	253709	1739

All positions are relative to Supercontig 1.541 of the Liverpool LVP AaegL1 genomic strain of *Aedes aegypti* (www.vectorbase.org).

the respective immunoreactivities were quantified using ImageJ software (http://rsbweb.nih.gov/ij).

Immunohistochemistry. Sections of paraformaldehydefixed Malpighian tubules embedded in paraffin were rehydrated and treated with methanol/peroxide as described previously.³² Following the methanol/peroxide treatment, the slides were subjected to a "wetautoclave" antigen-retrieval procedure³³ and then washed in phosphate buffered saline (PBS; 145 mM NaCl, 3.2 mM NaH₂PO₄, and 7.2 mM Na₂HPO₄, pH 7.5) for 5 min. The sections were blocked for 20 min at room temperature with a 4:1 mixture of 10% normal goat serum (Zymed) and 10× Casein solution (Vector Laboratories), then incubated overnight at 4°C with the adducin antibody diluted 1:1,000 in PBS supplemented with 1× casein. The following day, the sections were washed with PBS as described above and probed with a 1:100 dilution of a biontylated goat-anti rabbit secondary antibody (Vector Laboratories) in PBS-casein for 20 min at room temperature. The sections were then washed in PBS and incubated with a streptavidin-conjugated horseradish peroxidase (Zymed) for 30 min at room temperature. After washing again with PBS, the sections were stained with Vector AEC (Vector Laboratories) for 15 min at room temperature and then washed with water. A counter-stain was performed with hematoxylin for 10 s. Since a pre-immune serum for the adducin antibody was not available, we consider tubules undergoing the staining procedure without the antibody as negative control.

Electrophysiological studies of isolated Malpighian tubules. After isolation from a 3- to 5-d-old female mosquito, the set of five Malpighian tubules (still attached to the midgut) were transferred to a perfusion bath containing 0.5 ml of Ringer solution. The bottom of the bath was covered with a thin sheet of Parafilm (American National Can). Malpighian tubules cling to stretched Parafilm which stabilizes them during the perfusion of the bath at a rate of 3 ml/min. The midgut served to position the tubules in the center of the perfusion bath for impalement with microelectrodes. Conventional microelectrodes (Omega dot borosilicate glass capillaries, model 30-30-1; Frederick Haer; or model 1B100-F4; World Precision Instruments) were pulled on a programmable puller (Model P-97; Sutter Instruments) to yield resistances between 20 and 40 M Ω when filled with 3 M KCl. The microelectrodes were bridged to the measuring hardware using an Ag/AgCl junction. The Ag/AgCl junction was prepared by first degreasing a silver wire (OD 0.25 mm) with alcohol and then by Cl plating the wire in a concentrated solution of household Clorox for 15 min. The Ag/AgCl wires were inserted into the back of voltage and current microelectrodes. An Ag/AgCl wire (OD 0.5 mm) was inserted into a 4% agar bridge of Ringer solution to serve as the ground electrode. A single principal cell near the blind (distal) end of one Malpighian tubule was chosen for impalement with both voltage and current electrodes as described previously.³⁴ The voltage electrode measured the basolateral membrane voltage $(V_{\rm bl})$ using the Geneclamp 500 electrometer (Axon Instruments, Molecular Devices,). The input resistance (R_{in}) of the impaled principal cell was measured from current-voltage plots generated by voltage clamping the cell in a series of five increasing voltage-clamp steps: 5 mV, 400 ms each, starting at $(V_{\rm bl} - 10 \text{ mV})$. The voltage-stepping protocol and data acquisition were executed digitally by using a Digidata 1440 (Molecular Devices) under control of the Clampex module of the pCLAMP software package (version 10; Molecular Devices). The experiment was discontinued if V_{bl} measured by voltage and current electrodes differed by more than 10 mV. Moreover, the experiment was discontinued if the tubules did not respond to stimulation by AK-III.9 After washout of AK-III, new control values of V_{bl} and R_{in} were recorded. Thereafter, BIM-I (10⁻⁵ M) or PMA (10⁻⁶ M), was added to the bath and values of $V_{\rm bl}$ and $R_{\rm in}$ were taken every few min for the next 20 min. AK-III (10-6 M) was then added to record effects on V_{bl} and R_{in} . The experiments concluded with a final washout and final measurements of V_{bl} and R_{in}. All experiments were done at room temperature.

developed by Ramsay³⁵ and adopted by us.³⁶ In brief, the distal (blind) end of the tubule was bathed in a Ringer droplet of 50 μ l under light mineral oil. The open end of the tubule was pulled into the oil with a glass hook so that secreted fluid exited the tubule lumen as a droplet separate from the bathing Ringer solution. The glass hook was formed on a microforge (Stoelting Co.) using soft glass (R-6, Drummond Scientific Co.). It was then washed in an acid solution of K₂CrO₄ and H₂SO₄. After rinsing with distilled water and drying, the glass hook was exposed to the vapor of 20 μ l dimethyl-dichlorosilan for 90 sec (Fluka) and then baked overnight at 110°C. The silanization prevents fluid secreted by the tubules from spreading out along the glass hook.

Each tubule was used as its own control. The initial 30 min marked the control fluid secretion period. Thereafter, the secreted fluid droplet (nanoliter volumes, 10-9 liter) was removed and 5 µl of Ringer solution was removed from the peritubular bath and replaced with 5 µl of Ringer solution containing the agent of interest. The 30 min experimental period began as soon as the agent had been added to the peritubular bath. In some experiments tubules were first studied under control conditions for 30 min, then in the presence of bisindolylmaleimide-I for another 30 min, and finally in the presence of bisindolylmaleimide plus AK-III for a third 30 min interval. Rates of fluid secretion were determined by plotting cumulative volume (nl) secreted by the tubule as a function of time (30 min), each for the control and the experimental period(s). The plots were usually linear, yielding the rate of fluid secretion as the slope of the line (Microsoft Excel 2007). Using each tubule as its own control allowed the statistical analysis by the paired t-test which tests the significance of the difference between the control and experimental secretion periods. The paired comparison eliminates the variability between the secretion rates in tubules isolated from different mosquitoes.

Statistical analyses. Graphpad Prism (Graphpad Software, www.graphpad.com) was used for the statistical analyses of quantitative phosphorylations; one-way ANOVA with a Newman-Keuls post-test was used to evaluate the significance of multiple treatments, and the paired Student's t-test was used in studies of fluid secretion and electrophysiology where each tubule was used as its own control.

Disclosure of Potential Conflicts of Interest

The manuscript contains no issues that would present conflict of interest for the authors.

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