Uroplakins play conserved roles in egg fertilization and acquired additional urothelial functions during mammalian divergence

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ABSTRACT Uroplakin (UP) tetraspanins and their associated proteins are major mammalian urothelial differentiation products that form unique two-dimensional crystals of 16-nm particles (“urothelial plaques”) covering the apical urothelial surface. Although uroplakins are highly expressed only in mammalian urothelium and are often referred to as being urothelium specific, they are also expressed in several mouse nonurothelial cell types in stomach, kidney, prostate, epididymis, testis/sperms, and ovary/oocytes. In oocytes, uroplakins colocalize with CD9 on cell-surface and multivesicular body-derived exosomes, and the cytoplasmic tail of UPIIIa undergoes a conserved fertilization-dependent, Fyn-mediated tyrosine phosphorylation that also occurs in Xenopus laevis eggs. Uroplakin knockout and antibody blocking reduce mouse eggs’ fertilization rate in in vitro fertilization assays, and UPII/IIIa double-knockout mice have a smaller litter size. Phylogenetic analyses showed that uroplakin sequences underwent significant mammal-specific changes. These results suggest that, by mediating signal transduction and modulating membrane stability that do not require two-dimensional-crystal formation, uroplakins can perform conserved and more ancestral fertilization functions in mouse and frog eggs. Uroplakins acquired the ability to form two-dimensional-crystalline plaques during mammalian divergence, enabling them to perform additional functions, including umbrella cell enlargement and the formation of permeability and mechanical barriers, to protect/modify the apical surface of the modern-day mammalian urothelium.

INTRODUCTION Urothelium is a multilayered epithelium, also known as transitional epithelium, that covers the luminal side of the lower urinary tract, including the renal pelvis, ureter, urinary bladder, and upper urethra. This epithelium is one of the most frequent sources of tumor formation and is the attachment site of the type 1–fimbriated Escherichia coli, which causes urinary tract infection. The terminally differentiated, superficial urothelial cell, known as the umbrella cell, is highly specialized, as it provides an exceptionally effective permeability barrier...
(Negrete et al., 1996). To perform this function, the tight junction-sealed and stretched umbrella cell is highly enlarged, being able to reach a diameter of 75–100 µm, and its apical surface is completely covered by rigid-looking plaques consisting of naturally formed two-dimensional crystals of 16-nm plaques (“urothelial plaques”; 0.5- to 1-µm diameter) interconnected by the so-called hinge areas (Hicks, 1965; Kachar et al., 1999). We showed that highly purified urothelial plaques contain four major integral membrane proteins, which we named uropakins (urothelial plaques; abbreviated UP) (Wu et al., 1990, 1994; Yu et al., 1990). These are UPIa (28-kDa), UPIb (27-kDa), UPII (15-kDa), and UPIIIa (47-kDa); and several minor UPIII isoforms; see below), which are major mammalian urothelial differentiation products (Wu et al., 2009; Sun et al., 2013). UPIa and UPIb are tetraspanins, which interact specifically with single-spanned partner proteins uropakins UPI and Ila, respectively, to progressively form heterodimer (Ia/II and Ia/Illa), heterotetramer (Ia/IIb/Illa), and 16-nm particle (six heterotetramers) that are then hexagonally packed into two-dimensional crystalline arrays (Wu et al., 1995; Liang et al., 2001; Tu et al., 2002; Hu et al., 2005, 2008; Min et al., 2006). Our knockout (KO) studies showed that uropakins play important roles in the permeability barrier function and the enlargement of the terminally differentiated mouse urothelial umbrella cells (Hu et al., 2000; Kong et al., 2004).

Uropakins have several paradoxical features. First, although uropakins are highly expressed only in the plaque-forming mammalian urothelium and are often referred to as being urothelium specific (Wu et al., 1994; Romih et al., 2002; Olsburgh et al., 2003; Sun et al., 2013), they have been detected, as mRNAs or proteins, in the bladder, fat cells, and oocytes of Xenopus laevis (Sakakibara et al., 2005; García-Espana et al., 2006), pronephric tubules of zebrafish (Mitá et al., 2012), and several mammalian nonurothelial tissues and tumors (see Discussion). The significance of nonurothelial cells expressing a low amount of uropakins is unclear. Second, Fukami, Sato, and coworkers found that in Xenopus eggs the tyrosine (Tyr)-phosphorylation of a UPIlla-like protein may play an integral role in egg fertilization (Mahbub Hasan et al., 2005; Sakakibara et al., 2005; Sato, 2008). However, since uropakin-II and Ila-knockout mice are fertile (Hu et al., 2000; Kong et al., 2004), it is unclear whether this is a Xenopus-specific phenomenon. Third, our phylogenetic studies showed that uropakin genes are highly conserved throughout vertebrate evolution including nonmammals that either have a bladder epithelium not covered by urothelial plaques (e.g., Xenopus laevis) or do not even have a urinary bladder at all (e.g., fishes) (García-Espana et al., 2006; García-Espana et al., 2008; Desalle et al., 2014). Taken together, these considerations raise the possibility that mammalian uropakins may play some additional, highly conserved functions in nonurothelial cells.

Another important question has to do with the relationship among uropakin la, lb, and other tetraspanins. The tetraspanin family consists of tetra-spanned integral membrane proteins that can be divided into two groups (García-Espana et al., 2008; Charrin et al., 2014). The first group consists of several “general” tetraspanins, including CD9, CD37, CD53, CD63, CD81, CD87, CD91, and CD151. They are widely distributed, can interact with one another to form tetraspanin-enriched microdomains, and are involved in cellular migration, interaction, fusion, infection, and fertilization (Berditchevski and Rubinstein, 2013; Termini and Gillette, 2017; van Deventer et al., 2017). The second group consists of highly “specialized” and tissue-specific tetraspanins, that is, the retina-associated peripherin/RDS and Rom-1 (Goldberg et al., 2016; Stuck et al., 2016) and the urothelium-associated uropakins UPIa and UPIb (Wu et al., 2009; Wankel et al., 2016). However, whether the tetraspanins of this second group of “specialized” tetraspanins are so sharply tissue restricted and whether they can interact with other archetypal tetraspanins of the general group is unclear.

We report here that uropakins are made not only by mouse urothelium but also by several nonurothelial cell types, many but not all of urogenital origin, in the stomach, kidney, prostate, epididymis, testes/sperms, and ovary/oocytes. We show that uropakin knockout and antibody blocking diminish mouse eggs’ fertilization function in vitro fertilization assays; that UPII/Ila double-knockout mice have a smaller litter size; and that on fertilization the cytoplasmic tail of mouse uroplakin Ila undergoes a conserved Fyn-mediated tyrosine phosphorylation that plays an integral role in Xenopus laevis egg fertilization (Mahbub Hasan et al., 2005; Sakakibara et al., 2005). Together with our phylogenetic data, these results indicate that uropakins, which were first described and have been studied predominantly in mammalian urothelium, can actually perform conserved and more ancestral functions in egg fertilization and that they acquired only later during mammalian divergence their ability to form two-dimensional crystals of 16-nm particles (urothelial plaques) to perform additional functions, including umbrella cell expansion and the formation of a highly efficient permeability and mechanical barrier, to protect and functionally modify the apical surface of the modern-day mammalian urinary bladder epithelium. Our results also indicate that the “specialized” uropakin tetraspanins can colocalize and may interact with “general” tetraspanins such as CD9, thus blurring the artificial boundary between these two classes of tetraspanins.

RESULTS
Uropakins can be detected in nonurothelial lineages
Existing data showed that uropakins are highly expressed in the mammalian urothelium that covers the entire lower urinary tract, including the bladder, ureter, and renal pelvis (for a discussion of urothelium nomenclature, see Liang et al. [2005]). Relatively little is known about whether uropakins are expressed in nonurothelial tissues, because some of the existing antibodies to uropakins, although sufficiently specific for the detection of uropakins in urothelial cells, have a relatively low signal-background ratio in nonurothelial cells (unpublished data). We improved the sensitivity and specificity of uropakin detection in this study by using affinity-purified rabbit polyclonal antibodies against synthetic peptides or recombinant uropakin domains, plus several newly generated mouse monoclonal antibodies listed in Table 1 (for a complete list of antibodies to uropakins, see Supplemental Table S1). When these antibodies were used to immunoblot the proteins of purified plaques or the total urothelial extracts, they recognized their correct antigen as a single band, thus establishing their monospecificity. All the experiments were performed using multiple antibodies, with similar results. Under these conditions, we can unambiguously detect uropakins not only in mouse urothelial upper cells (Figure 1, A1–A5, showing the detection of UPIa, II, Ib, Ila, and Ilib) but also in the keratin 8 (K8) keratin-positive principal cells of renal medullary collecting ductal epithelium (Figure 1B), some of the K8-positive secretory cells of the anterior prostate epithelium (Figure 1C), the H, K(+)ATPase-positive, acid-secreting parietal cells of the gastric epithelium (Figure 1D), and the principal cells of cauda epididymal epithelium (Figure 1E), corneal epithelium (Figure 1F) (Adachi et al., 2000), ovary (Figure 1, G–J, see below), and testis (Figure 1L and Figure 6 later in this article and Supplemental Figure S5; see below). In the following experiments, we will focus on an in-depth analysis of the ovary and testis uropakins, which were confirmed by RT-PCR (Figure 1M).
Uroplakins are surface-expressed as heterotetramers in mouse oocytes/eggs

Our results indicate that oocyte is the uroplakin-expressing cells in the ovary and that all four major uroplakins (Ia, Ib, II, IIIa) as well as UPIib, a minor UPII isoform (Deng et al., 2002), are associated with the cell periphery and cytoplasmic vesicles of mouse oocytes (Figure 1G, 1–5). Although immunoblotting did not work because the tremendous dilution of the oocyte uroplakins by other uroplakin-negative ovarian cellular proteins, the staining specificity was established using at least two independent antibodies as well as by PCR (Figure 1M). Importantly, staining of the UPII-knockout mouse oocytes showed not only the expected loss of UPII but also its partner UPIa (Figure 1H, 1–5) (Kong et al., 2004). Moreover, consistent with our earlier data (Deng et al., 2002; Hu et al., 2002), the UPIIa-null oocytes lacked UPIIa but retained its partner UPIb that can be stabilized by UPIIib (Figure 1J, 1–5) (Deng et al., 2002). As expected, the UP-staining pattern of the UPII/UPIIa-double-knockout oocytes was the sum of the two single knockouts (Figure 1K, 1–5). These results confirmed the specificity of the UP staining. Moreover, we found that UPIa and II, which form the UPIa/II pair, colocalized precisely in intact mouse eggs (Supplemental Figure S1A), as did UPIb and IIIa that form the UPIb/IIIa pair (Supplemental Figure S1B). Even Ia and IIIa (members from the two pairs) colocalized (Supplemental Figure S1C), suggesting the formation of the UPIa/IIa/Ib/IIb heterotetramer (Hu et al., 2005). A control experiment showed that even though UPIa and CD9 colocalized precisely in some areas (see below), their global distributions were distinguishable (Supplemental Figure S1D). Taken together, these results suggest that the four major uroplakins are present in mouse eggs, which are devoid of two-dimensional uroplakin crystals, as a heterotetramer complex (UPIa/IIa/Ib/IIb) (Hu et al., 2005). Uroplakin staining was also seen in human ovary (Supplemental Figure S2).

Like several other egg markers, including CD9, GM1 ganglioside (a lipid raft marker), and lens culinaris agglutinin (LCA, which recognizes cortical granules; Figure 2, A, B, and E), uroplakins became polarized to the microvilli-enriched membranes during egg maturation (Figure 2, C and D), and UPIIa knockdown did not affect polarization (Supplemental Figure S3). Moreover, staining of intact oocytes and eggs showed that the UPIIa is exposed on the cell surface (Figure 2C, 1 and 2). The use of paraffin-sections showed that although in immature oocytes UPIIa was predominantly associated with intracellular vesicles (Figure 2D1), it became mostly surface membrane associated in mature eggs (Figure 2D2), indicating major maturation-associated translocation. In addition, developmental studies show that uroplakins were first detected in oocytes of the preantral follicles, where uroplakins were present at cell peripheries as well as cytoplasmic vesicles (Supplemental Figure S4).

In mouse eggs, uroplakins are associated with multivesicular bodies that give rise to the uroplakin-, CD9-, and CD81-positive exosomes

To determine the subcellular localization of uroplakins, we costained uroplakins with markers for various oocyte organelles. Consistent with uroplakins’ integral membrane nature, they partially colocalized with an endoplasmic reticulum (ER) marker (Sec23A; Figure 2F), a Golgi marker (GM130; Figure 2G), and a late endosome marker (lysosomal associated membrane protein 1 or LAMP1; Figure 2H). They did not colocalize at all with cortical granule (Figure 2K). However, they partially colocalized with two independent markers for multivesicular bodies (MVB), that is, Alix (Figure 2M; an adaptor protein involved in intraluminal vesicle formation) (Morita et al., 2007) and Hsp90 (Figure 2L2 and unpublished data). Taken together, these results indicate that uroplakins are associated with, in addition to the cell surface, the cytoplasmic MVBs.

High-resolution confocal microscopy revealed that, in many areas of the egg cell surface, uroplakins colocalized extremely well with CD9 (Figure 3A) but only partially with CD81 (Figure 3B). They also partially colocalized, in the perivitelline space (PVS) and zona pellucida (ZP), on some extracellular organelles that appeared to be exosomes based on their size, location, and antigenic properties (Figure 3, C and D). The colocalization of uroplakins with CD9 and CD81 is of interest because these two CD molecules are egg exosome markers and they are known to play key roles in egg fertilization (Miyado et al., 2000; Le Naour et al., 2006; Rubinstein et al., 2006). The fact that uroplakins colocalized partially with CD9 and CD81 in exosomes indicates that these extracellular organelles are heterogeneous with respect to their UP, CD9, and CD81 composition (Figure 3D).

These data are confirmed and extended by the immunogold EM-labeling studies, which established the association of uroplakins with plasma membrane (Figure 3E), microvilli (Figure 3F), intraluminal vesicles of MVBs (Figure 3, G and H) (Vieira et al., 2014), as well as extracellular vesicles in the PVS that are 30 to 150 nm in diameter and morphologically similar to exosomes (Figure 3I) (Meldolesi, 2018). Finally, antibodies to ovastacin produced expected staining for ovastacin.
of cortical granules (Figure 3J) that are morphologically distinct from the UP-positive multivesicular bodies (Figure 3, G and H) that, in UPII-knockout eggs, are replaced by abnormal multilobular structures (Figure 3K).

Mouse uroplakin IIIa undergoes conserved tyrosine-phosphorylation on egg fertilization

Of the four major uroplakins (Ia/II and Ib/IIda), UPIIa is the only one that has a long cytoplasmic tail that is known to play a signaling function (Thumbikat et al., 2009). Fukami, Sato, and coworkers showed that in Xenopus eggs a UPIIa-like protein undergoes Src-mediated Tyr-phosphorylation in a Y[S/T] motif and that antibody to the extracellular domain of this uroplakin abolishes Xenopus egg fertilization (Mahbub Hasan et al., 2005; Sakakibara et al., 2005; Mahbub Hasan et al., 2007). Since the molecular players involved in sperm–egg interaction/fertilization can vary even among mammalian species (Evans, 2012), and since a similar (S/A)Y[T/S] motif is conserved in all mammalian UPIIls (Supplemental Figure S6D), we wanted to see whether this fertilization-dependent event occurs in the evolutionarily distant mouse eggs.

To confirm this phosphorylation event in Xenopus oocytes, we made antibodies against all five frog uroplakins, that is, xUPIa, xUPIb, xUPII, xUPIIIa, and xUPIIIb, which we described earlier as cDNAs (x stands for Xenopus) (Garcia-Espana et al., 2006); of these, only xUPIb and xUPIIIb have been studied as proteins (Mahbub Hasan et al., 2005; Sakakibara et al., 2005). We also made sequence-specific antibodies against the nonphosphorylated and Tyr249-phosphorylated peptide of xUPIIIa (see Supplemental Table S1 for a list of antibodies to uroplakin proteins and peptides). Antibodies to all five uroplakins stained the superficial cells of Xenopus bladder epithelium (unpublished data), as well as the cell surface of Xenopus oocytes and mature eggs (Figure 4, A–F); none of them known to have two-dimensional crystalline plaques.

Although the untreated eggs had no detectable Tyr-phosphorylation of UPIIIa and Src (Figure 4, M and N), sperm-fertilized and H2O2-activated Xenopus eggs showed the phosphorylation of Tyr249 of xUPIIIa (Figure 4, P and V) as well as Tyr416 of Src (Figure 4, Q and W). Control experiments confirmed that tyrosine phosphorylation of both xUPIIIa and Src in sperm-fertilized eggs was blocked by a rabbit antibody to the extracellular domain of xIIIa (Figure 4, S and T) (Sakakibara et al., 2005), and showed that the reactions in peroxide-activated eggs

![FIGURE 1: Detection of uroplakins in nonurothelial cells. Immunofluorescence staining of paraffin sections of mouse bladder urothelium (A), as well as those of nonurothelial tissues, including the (B) renal medulla, (C) anterior prostate, (D) stomach, (E) cauda epididymis, (F) cornea, (G–K) ovary, and (L) testis. Ovary samples are from (G) wild-type mouse, (H) UPII knockout, (J) UPIIIa knockout, and (K) UPII/UPIIIa double knockout. Antibodies to individual uroplakins include (A1) Ia 4867; (A2 and L) Ib 7472; (B, C, E) 7727; (A3) II 160; (A4) IIIa 182; (D) 35804; and (F) IIIb 6177; for the description of these antibodies, see Table 1 and Supplemental Table S1). Other antibodies include alpha-tubulin (α-Tub), K8, V-ATPase, and H+;K+-ATPase, as noted. (M) Detection of uroplakins mRNAs by RT-PCR in bladder (lane 1; B), testis (2; T), ovary (3; O), or water control after 30 (low, L) or 35 (high, H) cycles Glyceraldehyde 3-phosphate dehydrogenase (GAPDH) control. Note the detection of uroplakins in B the apical surface of the K8-positive renal collecting duct, (C) the upper parts of some of the K8-positive secretory prostatic epithelial cells, (D) the H+;K+-ATPase-positive gastric parietal cells, (E) the apical surface of the V-ATPase-negative epididymal epithelial cells, (F) the corneal epithelium, (G–K) oocytes, and (L) spermatids in testis. Also note in G to K the absence of UPIII (and its partner UPII) and UPIIIa in the UPII- and UPIIIa-KO oocytes, respectively, thus confirming the specificity of the uroplakin staining. L = lumen. Nuclear staining: DAPI (A–F) or To-Pro-3 (G–K). Bars equal to 10 µm (A, C, F, and L), 50 µm (B, E), 20 µm (D), or 30 µm (G–J).]
The specificity of the phosphorylation was shown in a control, indicating that the amount of total phosphorylated tyrosine (detected by a pan anti-phosphotyrosine antibody) rose tremendously on egg activation (Figure 5, M and N), and this was also SU6656 sensitive (Figure 5O). High-resolution confocal microscopy revealed that the UPII, Fyn, and CD9 colocalized in discrete patches (Figure 5P) and that, after fertilization, the Tyr-phosphorylated UPIIIa and Fyn remained colocalized (Figure 5Q). Notably, UPIIa barely colocalized with GM1, a raft marker, indicating that the uroplakin-containing tetraspanin-enriched microdomains, like that of other tetraspanins (Claes et al., 2001; Le Naour et al., 2006), are distinct and K), and this event was also blocked by SU6656 (Figure 5L).

Interestingly, egg activation also led to the phosphorylation of Tyr (Figure 5E), which was blocked by Src-inhibitor SU6656 (Figure 5F).

Ethanol activation of mouse eggs led to Tyr phosphorylation (Mahbub Hasan et al., 2006), confirmed that on fertilization Tyr phosphorylation (Figure 5D), while control mouse eggs had no detectable Tyr phosphorylation (Figure 5C). While fertilized mouse eggs have an in vitro fertilization rate of ∼82%, those of the UPIIIa KO, and double KO dropped to 47, 41, and 39%, respectively. The specificity of the phosphorylation was shown in a control, indicating that the amount of total phosphorylated tyrosine (detected by a pan anti-phosphotyrosine antibody) rose tremendously on egg activation (Figure 5, M and N), and this was also SU6656 sensitive (Figure 5O).

We next examined the possible phosphorylation of Tyr of mouse UPIIIa (in the same motif as Tyr of Xenopus). While control mouse eggs had no detectable Tyr-phosphorylation (Figure 5D), ethanol activation of mouse eggs led to Tyr-phosphorylation (Figure 5E), which was blocked by Src-inhibitor SU6656 (Figure 5F). Interestingly, egg activation also led to the phosphorylation of Tyr in Fyn, the major Src-related kinase in mouse oocytes (Figure 5, J and K), and this event was also blocked by SU6656 (Figure 5L).

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in all these cases the blastocyst formation rates were not significantly affected (Figure 7A). To assess the effects of UP deficiency on male vs. female gametes, we repeated the IVF assays using gametes from normal mice in combination with gametes from the UP-null mice of the opposite sex. The results showed that eggs from the UPII KO and UPIIa KO mice had reduced fertilization rates of 42 and 34%, respectively, indicating that uroplakins play an essential role in egg fertilization (Figure 7B). However, sperms from the UPII KO, UPIIa KO, and double-KO mice had normal fertilization rates of 87, 89, and 77%, respectively, indicating that uroplakin-null sperms function normally in the setting of this in vitro assay (Figure 7C; see Discussion). The importance of uroplakins in egg fertilization was supported by the fact that preincubation of normal eggs with antibodies against the extracellular epitopes of CD9, UPla, and UPlb reduced the in vitro fertilization from 68% to 40, 32, and 7%, respectively (Figure 7D). Although an antibody to UPIIa failed to block fertilization, it is possible that this was due to the inaccessibility of its particular epitope. Finally, we assessed the litter size of the UPII-, UPIIa- and double-knockout mice. Although the litter sizes of the UPII KO and UPIIa KO mice were in the normal range (Hu et al., 2000; Kong et al., 2004), those of the double-KO mice were significantly reduced from an average of 11–6 (Figure 7E).

Emergence of mammal-specific uroplakin sequences

On the basis of the fully sequenced genomes of a broad array of chordates (cephalochordats, urochordates, and vertebrates), we have previously analyzed the evolution of uroplakin genes and showed that they originate in the common ancestor of vertebrates (Garcia-Esparza et al., 2006; Desalle et al., 2014; Chicote et al., 2017). Analyses of the vertebrate uroplakin sequences revealed many mammal-specific residues dispersed by residues that are common to all (mammalian and nonmammalian) uroplakins (Figure 8, A and B, and Supplemental Figure S6) (Desalle et al., 2014). It is likely that the amino acid residues that are common to all vertebrates are involved in forming the basic tetraspanin structure and playing ancestral functions in nonurothelial cells, including egg fertilization, whereas some of the mammal-specific residues may contribute to two-dimensional crystallization and other mammal-specific functions (Figure 8, A and B, and Supplemental Figure S6). It is likely that the amino acid residues that are common to all vertebrates are involved in forming the basic tetraspanin structure and playing ancestral functions in nonurothelial cells, including egg fertilization, whereas some of the mammal-specific residues may contribute to two-dimensional crystallization and other mammal-specific functions (Figure 8, A and B, and Supplemental Figure S6). Finally, of all the uroplakins, UPlb stood out as having the smallest number of...
Xenopus Oocyte

Egg

xUPIIIa-Y249p  Src-Y416p  P-Tyr

Control

Sperm

α-xIIla Sperm

H₂O₂

H₂O₂  SKI-606

FIGURE 4: Fertilization of Xenopus laevis (Xl) eggs led to tyrosine phosphorylation of UPIIIa and Src. Xenopus oocytes (A–I) were stained using (A) normal IgG (negative control), or antibodies to (B) Xenopus UPIa or xUPIa (19228), (C) xUPII (13641), (D) xUPIb (13638), (E) xIIla (19230), (F) xIIlb (4865), (G) xIIla-Y249 (nonphosphorylated peptide, 35761), (H) xIIla-Y249P (Tyr-phosphorylated peptide, 35760), or (I) Src-Y416P (Tyr-phosphorylated peptide). Alpha-tubulin was colocalized in A–F as a control. Note the detection of all five Xenopus homologues of mammalian uroplakins on the egg surface. (J–L) Xenopus eggs were stained using antibodies to (J) xIIIa-Y249, (K) normal mouse IgG, or (L) normal rabbit IgG. (M–O) Control Xl eggs, (P–R) sperm-fertilized eggs, (S–U) eggs pretreated with an antibody to xUPIIla before sperm fertilization, (V–X) hydrogen peroxide-activated eggs, and (Y–ZZ) SKI-606–pretreated eggs before hydrogen peroxide activation. In this series of experiments, each type of egg was stained using antibodies to xIIla Y249P (first column), Src-Y416P (second), and pan phosphorylated-tyrosine (third). Note the increased Y-phosphorylation of xIIla and Src in both sperm-fertilized (P, Q) and peroxide-activated (V, W) eggs and the blockage of this reaction by an anti-xUPIIIa antibody (S, T) and Src inhibitor SKI-606 (Y, Z). Bars equal to 20 µm.

mammal-specific sequences (13%; vs. 55% common sequence), compared with all other uroplakins: UPla (33%, 40%), UPlI (49%; 21%), UPIIIa (36%; 27%), and UPIIIb (27%; 25%) (Figure 8B).

e.g., Carpenter et al. [2016] and Human-Protein-Atlas [2018]. Thus, for most purposes of cell/tumor identification it is in general more reliable to detect the uroplakin proteins.
Our finding that uroplakin expression is less urothelium-restricted than previously thought raises a cautionary note about the use of uroplakin promoters to drive urothelial expression of transgenes for developing mouse models of bladder tumorigenesis and other urinary tract diseases (see, e.g., Lin et al., 1995; Wu, 2009; Wu et al., 2009; Schnegelsberg et al., 2010). Although in most of these studies the transgenes targeted mainly urothelium and generated useful models, some studies showed unexpected targeting of nonurothelial tissues such as the lung (Ahmad et al., 2011; Ayala de la Pena et al., 2014), which may be explained by this work. In future studies, the urothelial specificity of uroplakin promoters can be further improved, if needed, by using an inducible system coupled with the intravesicle delivery of the doxycycline or tamoxifen inducer (Zhou et al., 2010; Van Batavia et al., 2014).

Distinct trafficking patterns of uroplakins in oocytes and urothelial umbrella cells

Our data indicate that the uroplakin trafficking patterns of the eggs and urothelial cells are distinct (see models in Figure 9). Thus, in mouse oocytes where a low level of uroplakins are targeted to the cell-surface and MVB-derived exosomes, these proteins are delivered from the ER/Golgi either via small exocytotic vesicles (Figure 2D1) to the cell surface (Figure 2D2), or to the MVBs (Figures 2M and 3, G and H), which then give rise to the UP-, CD9-, and/or CD81-positive exosomes (Figures 3, D and I, and 9A). In urothelial umbrella cells, where a huge amount of uroplakins is targeted predominantly to the apical surface, they are first assembled in post-Golgi vesicles into two-dimensional crystals of 16-nm particles (with six tetramers), which are delivered via specialized discoidal (DV) and fusiform vesicles (FV) to the apical surface (Wankel et al., 2016; Gallo et al., 2018) and then endocytosed for MVB/lysosome-mediated degradation (Figure 9B) (Khandelwal et al., 2009; Vieira et al., 2014). It should be noted, however, that even though the major uroplakin pathways (thick arrows in Figure 9) differ in these two cell types, they do overlap (Figure 9).

Possible roles of uroplakins in fertilization

An important conclusion of our work is that uroplakins, which were discovered as major mammalian urothelial differentiation products and have been studied predominantly in this epithelium (Wu et al., 2009; Sun et al., 2013; Wankel et al., 2016), perform essential functions in mouse egg fertilization. This conclusion is based on our observations that antibodies to the extracellular domains of UPIIa and Fyn led to the tyrosine-phosphorylation of UPIIIa and Fyn. Permeabilized normal eggs (control; first column), ethanol-activated eggs (second column), and SU6656 (Fyn inhibitor)-pretreated and ethanol-activated eggs (third column) were stained using antibodies to (A–C) UPIIIa-Y266 (35759), (D–F) UPIIIa-Y266P (35758) (synthetic peptide containing the phosphorylated (P)-Tyr 266), (G–I) Fyn, (J–L) Fyn-Y418P (synthetic peptide containing the phosphorylated Tyr 418 of Fyn, and (M–O) phosphorylated Tyr. Note in the second column that ethanol-activation led to the Tyr-phosphorylation of UPIIIa-Y266 (E) and Fyn-Y418 (K), and their blockage by SU6656 (third column), (P–R) Immunostaining of intact mouse eggs with antibodies to (P) UPIII (S3045)/Fyn/CD9, (Q) UPIIIa-Y266P/Fyn-Y418P, and (R) UPIIa/GM1-CTB (a raft marker). Note that in P the substantial colocalization of UPIII/Fyn with CD9; in Q the colocalization of tyrosine-phosphorylated UPIIIa and tyrosine-phosphorylated Fyn; and in R the poor colocalization between UPIIa and the raft marker GM-1. Bars equal to 10 µm (A–O) or 5 µm (P–R).

1He F, Wu XR, Sun T-T (2017). UPII Promoter: an update, http://sun-lab.med.nyu.edu/protocols-reagents/sun-lab-reagents.
FIGURE 6: Association of uroplakins with the mouse sperm head. (A, B) Paraffin-sections of mouse testis were triple-stained using antibodies to uroplakins Ia (4867) or II (160) (green), alpha-tubulin (red) and To-Pro-3 (blue; nuclei). (C–E) Uroplakin staining of intact mouse sperms from (C) wild-type, (D) UPII-knockout, and (E) UPIIIa-knockout mouse using antibodies to specific uroplakins as noted. (F, G) Staining of mouse sperms that have been treated with (F) PBS or (G) 0.1% Triton X-100. (H, I) UPilla staining of a control sperm (H) or a capacitated sperm (I). (J, K) Triple staining of control sperms (J) or those pretreated with 0.1% Triton X-100 (K), using antibodies to UPilla, Sp56 (acrosome marker) and DAPI. (L, M) Double-staining of sperms (that have been heat-treated) using anti-UPilla and PSA. Antibodies to uroplakins are as follows: Ia, AU-Ia-1; II, S3045; Ib, AU-Ib-2; IIIa, 35804; IIIb, 6177 (Table 1 and Supplemental Table S1). Note in H, I, J, and K the UP-staining of the sperm heads, in D the absence of UPII and its partner Ia in the UPII-null sperms, and in E the absence of UPilla in the IIIa-KO sperms. Also note in L that heat treatment had little effects on hook-associated UP-staining, and led to the formation of abnormal PSA stained swelling near the transitional zone between the mid-body and tail. The sperms were collected from cauda epididymis, and nuclear staining was by DAPI. Bars equal to 10 µm (A, B), 5 µm (F, G, L, and M), or 2 µm (C–E, H–K).
These data suggest that uropakins perform important, conserved functions in egg fertilization. Although the precise function of uropakins in egg fertilization is unclear, their association with the plasma membrane and particularly notably the microvilli of eggs (Figures 2, A–C, and 3, E and F), as well as egg-associated exosomes (Figure 3, D, G, and H), suggests that they perform membrane-related fertilization functions, including sperm–egg interactions and the formation and/or modulation of exosomes that play a critical role in egg–sperm fusion (Harada et al., 2013). Regarding the surface-related function, it has been suggested that xUPIIa may serve as a sperm receptor, based on the observation that an antibody to its extracellular domain reduced Xenopus egg fertilization (Mahbub Hasan et al., 2005). In this regard, it is of interest to note that in our studies antibodies to la and lb, but not liia (possibly due to the inaccessibility of the juxta-membrane epitope), reduced mouse egg in vitro fertilization (Figure 7D). This suggests that antibody binding to uropakins la and lb may induce local structural changes of the uropakins complex (see, e.g., Wang et al. [2009]), thus hampering UPIIIa signaling and egg fertilization, and that antibody blocking data per se do not necessarily mean that the protein antigen(s) is a sperm receptor. Overall, existing data suggest that uropakins participate in the following cellular processes in fertilization.

**Signal transduction**

**Egg fertilization triggers Tyr phosphorylation of UPIIIa.** Of the four major uropakins, UPIIIa is the only one that has a long C-terminal, cytoplasmic tail (∼50 amino acid residues) (Wu et al., 2009; Desalle et al., 2014), which is known to play a signaling role in, for example, urinary tract infection (Thumbikat et al., 2009). We have confirmed the earlier data by Fukami, Sato, and coworkers that Xenopus egg fertilization triggers the phosphorylation of Tyr266 located in the cytoplasmic tail of xUPIIIa (Figure 4) (Sakakibara et al., 2005) and showed for the first time that this fertilization-dependent Tyr-phosphorylation of UPIIIa (Tyr266) also occurs in mouse eggs (Figure 5). These results suggest that Tyr phosphorylation of UPIIIa must play a conserved role in egg fertilization. The importance of UPIIIa as a signaling molecule is supported by our recent finding that this protein is evolved from a protein tyrosine phosphatase receptor, a family of cell-surface receptors that initiate intracellular Try-phosphorylation-dependent signal transduction in response to the binding of extracellular ligands (Chicote et al., 2017; Yao et al., 2017).

**Fyn is responsible for the tyrosine phosphorylation of UPIIIa.** Our data suggesting that Fyn is responsible for Tyr-phosphorylating UPIIIa in mouse egg include the following: 1) uropakins and Fyn, the major murine egg Src-family kinase (Luo et al., 2009; McGinnis et al., 2011), colocalize forming discrete patches scattered on the surface of mouse eggs both before (Figure 5P) and after fertilization (Figure 5Q), suggesting that Tyr phosphorylation of UPIIIa must play a conserved role in egg fertilization. The four major uropakins, UP Ia and UP Ib may induce global structural changes of the uropakins complex (see, e.g., Wang et al. [2009]), thus hampering...
FIGURE 8: Mammal-specific amino acid residues and motifs in uroplakins. (A) Examples of mammal-specific residues (yellow) and amino acid residues that are shared by ALL uroplakins (“all-uroplakin-residues or AUR; red) from UPIa, Ib, II, IIIa, and IIb of ∼20 mammals (17 placental and three nonplacental) and 10 nonmammals (Desalle et al., 2014). Mammal-specific residues are defined as those that are present in >90% of mammals and <30% of nonmammals. The horizontal green line demarcates the sequences of the mammals (above) and nonmammal vertebrates. (B) The overall location of mammal-specific motifs (MSM; red box; greater than three consecutive residues, five with a single interruption, or six with two interruptions) in uroplakins as indicated. TMD (transmembrane domain). (C) Model depicting the evolution of all the known uroplakin genes. Symbols: red arrows (genealogical relationship), two-headed black arrows (protein–protein interaction), asterisk (a strong pattern of significant skew towards dN/dS>1.0 suggesting possible selection that accompanies the duplication events that produced the paralogue group) (Desalle et al., 2014), phosphotyrosine phosphatase receptor (PTPR) (Chicote et al., 2017), tetraspanin precursor (TM4), million years (MY). See main text for details.

**Uroplakin/CD9 interactions.** Our results indicate that on mouse egg surface, uroplakins colocalized closely with CD9, forming discrete patches (Figures 3A and 5P), and partially with CD81 (Figure 3B). These results are interesting because both tetraspanins CD9 and CD81 are known to play important roles in mouse egg fertilization (Le Naour et al., 2000; Miyado et al., 2000; Rubinstein et al., 2006) and because uroplakins Ia and Ib are also tetraspanins, which have the propensity to form the “tetraspanin-enriched microdomains” (TEMs) that can enhance signal transduction (Termini and Gillette, 2017). The poor colocalization between uroplakin and the raft marker GM1 (Figure 5R) indicates that the uroplakin-containing TEMs are distinct from the “rafts” (Hemler, 2003; Espenet et al., 2008). The fact that uroplakin/CD9 also colocalize with Fyn (Figure 5, P and Q) further suggests that CD9 may modulate/enhance UPIIIa/Fyn signaling. This idea may seem at odds with an earlier suggestion that xCD9 does not interact with xUPIIIa based on the absence of the former in the uroplakin-containing Triton X-100 resistant raft fraction (Mahbub Hasan et al., 2007). However, this discrepancy can be explained by the fact that tetraspanin association usually does not survive Triton X-100 (Indig et al., 1997; Miao et al., 2001). Indeed, their data showed that centrifugation in Brij 98, but not in Triton, allowed the cofloating of xCD9 with xUPIIIa and xUPIIb (Figure 1E in Mahbub Hasan et al., 2007). Finally, even though our data indicate CD9/uroplakin association in mouse egg TEMs (Figure 3A), it remains to be established whether they interact directly (Kovalenko et al., 2004;
Unlike most

FIGURE 9: Schematic diagrams showing the distinct patterns of uroplakin trafficking in (A) mouse oocyte/egg and (B) a terminally differentiated umbrella cells of mammalian bladder urothelium. (A) In mouse egg, uroplakins (red triangle luminal domain with a cytoplasmic tail; existing possibly as heterotetramers; see Discussion) are delivered to the cell surface via exocytosis or to exosomes via MVBs. (B) In urothelial umbrella cells, uroplakins are assembled in trans-Golgi network into 16-nm particles (red circles; 16-nm particles), which form growing two-dimensional crystals delivered via DV and FV to the urothelial apical surface, where they form the characteristic urothelial plaques (Wankel et al., 2016). Some of the apical surface-associated uroplakins can be endocytosed into multivesicular vesicles for lysosomal degradation (Vieira et al., 2014). Other abbreviations: EE (early endosome) and TGN (trans-Golgi network). Thickness of the arrows approximates the relative abundance of the pathways in the two cell types.

Dahmene et al., 2014) or form separate nanoclusters coexisting in TEMs (Zuidscherwoude et al., 2015).

Remarkable consistency of the mouse and frog data. The remarkable overall consistency between the mouse and Xenopus data suggests that uroplakins play a highly conserved role in egg fertilization and that Xenopus egg provides a relevant model for studying the biochemical and certain other aspects of uroplakin involvement in this process. In this regard, it should be noted that Sato, Fukami, and coworkers proposed a model, based on their Xenopus studies, suggesting that the cleavage of the extracellular domain of xUPIIIa at the juxta-membrane Gly-Arg-Arg (GRR) site by a sperm-derived protease(s) releases the xUPIb/IIIa-inhibition of Src, which then Tyr phosphorylates some of the remaining intact UPIIIa (Mahbub Hasan et al., 2005, 2007, 2014; Hasan et al., 2011). Additional studies are needed to identify the sperm protease(s) hypothesized to be involved in cleaving the juxta-membrane GRR site of the extracellular domain of xUPIIIa, and the functional consequences of UPIIIa phosphorylation, which may include the modulation of its interaction with a downstream target(s) and/or of the physical properties of the egg and/or exosome membranes (see below).

Modulation of membrane stability

Possible structural roles of uroplakins in sperms. Unlike most mammalian sperms that have an oval or paddlelike head, sperms of the myomorph rodents have an apical rostral hook, which contains part of the acrosome. The evolutionary and functional significance of such a unique hook structure is unclear, although it has been suggested to play a role in sperm velocity and postcopulatory competition (Moore et al., 2002; Varea-Sanchez et al., 2016). Since the uroplakin-enriched domain covers the concave (inside) surface of the hook (corresponding possibly to the “hook rim” structure) (Lin et al., 2013), which is far from the acrosome-abutting convex (outside) surface involved in egg binding (Avella and Dean, 2011), this uroplakin-enriched subdomain is unlikely to play a role in sperm–egg interaction. Given the possible roles of uroplakin proteins in enhancing membrane rigidity as discussed above, their association with the inside curvature of the hook including the tip (Figure 6, H–K) suggests a structural role, perhaps in conjunction with the microtubule manchette, in generating and/or maintaining the hook structure. Although UP knockout of males did not affect the sperm morphology (Figure 6, D, E, L, and M) or their ability to fertilize the eggs in our IVF assays and in vivo (Figure 7, C and E), the overwhelmingly high sperm–egg ratio in both of these assays could have masked the suboptimal performance of the UP-deficient sperms.

Uroplakins enhance membrane rigidity and/or stability. Several lines of evidence suggest that uroplakins enhance the stability and rigidity of the membrane as indicated by enhanced detergent insolubility. As mentioned earlier, uroplakins form two-dimensional crystals of 16-nm particles, the urothelial plaques, which form rigid-looking concave plaques covering the mammalian urothelial apical surface (Hicks, 1965; Kachar et al., 1999). Unlike other tetraspan-in enriched microdomains that can survive only relatively weak detergents such as Brij 99 and CHAPS (Claas et al., 2001), urothelial plaques can largely survive more stringent detergents such as Triton X-100 or Sarkosyl (Mu et al., 1990; Liang et al., 1999). Interestingly, even the regular-looking plasma membranes that interconnect the plaques (the “hinge” areas), which contain no or largely collapsed uroplakin particles, are also insoluble in Triton and Sarkosyl (Liang et al., 1999). This suggests that uroplakins, even when they do not form 16-nm particles, can still enhance urothelial membrane stability as gauged by detergent insolubility. Taken together, these results suggest that uroplakins can increase the stability of membranes in urothelial hinges and possibly the uroplakin/CD9/CD81-containing egg TEM microdomains (Figure 3, A and B). Whether uroplakin mutations can cause fertilization problems in humans is unknown, although it should be noted that sequencing of human CD9 gene, which is critical for egg fertilization in mice, has not revealed any CD9 mutations causing human female fertility problems (Nishiyama et al., 2010), probably because defects in molecules critically important for fertility are subject to negative selection.

Evolution of the uroplakin gene family: a unifying model

Our results indicate that uroplakins function in oocytes where they play essential and more ancestral roles in fertilization without forming two-dimensional crystals (Figure 7). Coupled with the fact that mammalian uroplakins have abundant mammal-specific sequences (Figure 8, A and B, and Supplemental Figure S6), these data suggest that uroplakins play highly conserved roles such as egg

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fertilization in nonurothelial cells, and that they acquired an additional role in mammalian urothelium. This idea can explain why uroplakin genes have been preserved in most vertebrates, including nonmammals that do not even have a urinary bladder (Garcia-España et al., 2006, 2008; Desalle et al., 2014; Chicote et al., 2017), and it can also explain the expression of uropilakins in nonurothelial cells in which uropilakins may perform additional cell type-specific functions.

Although our phylogenetic studies of uropilakin genes shed light on how uropilakin genes are evolved (Garcia-España et al., 2006, 2008; Desalle et al., 2014), there are several features of this process remain paradoxical. Here we propose a unifying model of uropilakin gene evolution (Figure 8C), which integrates all our phylogenetic data, including our current findings (Figure 8, A and B, and Supplemental Figure S6), and provides possible explanations for these puzzling features. 1) At the beginning of vertebrates, a tetraspanin-originated proto-UPI gene gave rise to proto-UPIb and proto-UPIa (Garcia-España et al., 2006, 2008; Desalle et al., 2014). 2) The proto-UPIb recruited a precursor of the phosphotyrosine phosphatase receptor (PTPR) gene that evolved to become proto-UPIII, which then gave rise to proto-UPII by gene duplication (Chicote et al., 2017); this idea can explain why the UP1I sequences are much more distant than UP1II, from PTPR. 3) The proto-UPII gene gave rise to the current UP1IIs (referred to so far as UPII); present in all vertebrates, including mammals) and an early branch of UP1IIs that exists only in fish and reptiles; and proto-UP1II gave rise to UPIIIa, UPIIIb, and UPIIIc (present in all vertebrates, including mammals), as well as UPIIIId (fish only; corresponds to the UP3a-like protein described in Mitra et al. (2012) and Desalle et al. (2014). 4) Throughout uropilakin evolution, members within the UP1I/II and UP1I/III pairs strongly coevolved (Garcia-España et al., 2006, 2008; Desalle et al., 2014), consistent with their known specific protein/protein interactions (Wu et al., 1995; Tu et al., 2002; Hu et al., 2005). 5) Coinciding with the divergence of mammals, most of the mammalian uropilkins acquired a significant amount of mammal-specific changes (Figure 8, A and B, and Supplemental Figure S6, A–E; asterisks in Figure 8C). Thus, we have identified the evolutionary branches of uropilkins that have experienced statistically significant departure from neutrality in their dN/dS ratios in the uropilakin geneologies, with the single exception of the UP1b gene (Figure 8C) (Desalle et al., 2014). It is likely that such an acquisition of mammal-specific sequences forms the basis for neofunctionalization, including two-dimensional crystal formation. 6) There are some data indicating that UP1b protein has a broader tissue distribution than UP1a (Human-Protein-Atlas, [2018]; also see the mRNA data from Lobban et al. [1998] and Olsburgh et al. [2003]), suggesting that UP1b may interact with additional nonuroplakin partners which may impose extra constraints to UP1b evolution. This can explain why UP1b has undergone the least amount of mammal-specific sequence changes (Figure 8, B and C, and Supplemental Figure S6, A–E) (Desalle et al., 2014). Overall, uropilakin evolution provides a good example of genetic piracy. The ancestral and essential function of uropilkins in, for example, egg fertilization may have locked these genes into the genome. These genes are then pirated during mammal divergence for an additional cellular function, that is, uroepithelial structure, via gene duplication and neofunctionalization.

In conclusion, our data blurred the artificial boundary between the "general" and "specialized" tetraspanins, as uropilkins and their associated proteins, which were thought to be urothelium-specific, are also expressed in several nonurothelial cells, including oocytes of mouse and Xenopus laevis, in which they may interact with CD tetraspanins to play conserved and essential roles in fertilization. Only later during mammalian divergence uropilkins acquired ability to form two-dimensional crystalline urothelial plaques to support umbrella cell enlargement, as well as the formation of effective permeability and mechanical barriers (Hu et al., 2000, 2002; Kong et al., 2004), features that are essential for the formation of a functional modern-day mammalian urothelium. Collectively, our data indicate that uropilkins have a broader tissue distribution and play more diverse functions than hitherto appreciated. Additional work is needed to further decipher the functions and disease implications of uropilkins in not just urothelial cells but also nonurothelial cells.

MATERIALS AND METHODS

Reagents and antibodies

The reagents and their sources (in parentheses) are listed below. Hormones gonadotropin from pregnant mare serum (PMSG), chorionic gonadotropin human (hCG); hyaluronidase, acidic Tyrode’s solution, reduced L-glutathione (GSH), Src kinase inhibitor SU6656 and Ficol 400 (Sigma, St. Louis, MO); HEPES-buffered modified human tubal fluid (HTF; Irvine Scientific, Irvine, CA); Radioactive Fertilization Medium (RVF) medium (Cook Medical, Bloomington, IN); M2 and KSOM media (EMD Millipore Corp., Billerica MA); Ca/Mg-free Quinn’s Advantage medium with HEPES (Cooper Surgical, Trumbull, CT); Src kinase inhibitor SKI606 (Selleck Chemicals, Houston, TX); RNasey Mini Kit (Qiagen, Valencia, CA); Capacity Reverse Transcription Kit (Thermo Fisher Scientific, Waltham, MA); Lowicryl K4M (Polysciences, Warrington, PA); glutaraldehyde, paraformaldehyde, sodium cacodylate, osmium tetroxide, and EMbed 812 (Electron Microscopy Sciences, Hatfield, PA); and HQ Silver enhancement kit (Nanoprobes). Table 1 for the properties of 10 newly developed mouse monoclonal antibodies to uropilkins (plus the previously described AU1 antibody to UP1llla) (Liang et al., 2001). See Supplemental Table S1 for a complete list of our anti-uropilakin antibodies, including rabbit polyclonal antibodies against mouse and Xenopus uropilakin peptides and subdomains. All our mouse antibodies were purified using Protein G–coupled Sepharose beads, and rabbit antibodies were immunofinity-purified using gel-purified uropilkins or synthetic peptides. See Supplemental Table S2 for a list of other antibodies.

Animal strains

The generation and characterization of the UP1II-knockout, UP1llla-knockout, and UP1II/UP1IIa double-knockout mice were described earlier (Hu et al., 2000; Kong et al., 2004; Zocher et al., 2012). Oocytes, eggs, and sperms were collected from CB6F1 mice, 6–8 wk of age, from Charles River Laboratories, Wilmington, MA. Mouse tissues were collected from Swiss Webster mice (Taconic Biosciences, Hudson, NY). The Xenopus ovaries, eggs, and testis were purchased from Xenopus1.com, Dexter, MI. Our animal work was performed with the approval from the NYU School of Medicine’s Institutional Animal Care and Use Committee.

Immunofluorescence staining

Deparaffinized tissue sections were incubated with 5% bovine serum albumin (BSA) in phosphate-buffered saline (PBS) buffer and incubated with primary antibodies at 4°C overnight and later with secondary antibodies for 1 h at room temperature. ZP-free oocytes and eggs were fixed and treated with 0.1% Triton X-100 for 10 min at room temperature, blocked with 5% BSA in PBS buffer, incubated with primary antibodies at 4°C overnight, followed by incubation with secondary antibodies. ZP-intact eggs were incubated with...
primary antibodies for 30 min followed by 30 min of fixation at room temperature and then incubated with secondary antibodies for 1 h at room temperature. The oocyte and egg nuclei were stained with Hoechst 33258 (nonpermeabilized). Mouse sperms were collected from caput and cauda epididymis as described in the IVF method, fixed with 4% formaldehyde and permeabilized with 0.1% Triton X-100, blocked with 5% BSA, and incubated with primary antibodies and stained with DAPI.

Electron microscopy
Mouse oocytes and eggs were fixed in the fixative containing 2.5% glutaraldehyde and 2% paraformaldehyde in 0.1 M sodium cacodylate buffer (pH 7.2) for 2 h, postfixed with 1% osmium tetroxide for 1.5 h at room temperature, and then processed in a standard manner and embedded in EMbed 812 for transmission electron microscopy. For immunoelectron microscopy, mouse eggs were fixed in freshly made 3% paraformaldehyde in 0.1 M sodium cacodylate buffer containing 0.1% glutaraldehyde and 4% sucrose (pH 7.4). After washing and dehydration, the tissues were embedded in Lowicryl K4M and polymerized under UV (360 nm) at 35°C. Ultrathin sections were mounted on formvar-/carbon-coated nickel grids, incubated with primary antibodies at 4°C overnight and then with nanogold- or 18-nm colloidal gold-conjugated secondary antibodies followed by silver enhancement in the dark for 8 min. Grids were stained with uranyl acetate and lead citrate by standard methods, examined with a Philips CM12 electron microscope (FEI, Eindhoven, The Netherlands) and photographed with a Gatan (4 k × 2.7 k) digital camera (Gatan, Pleasanton, CA).

RT-PCR
Total RNA was prepared using the RNeasy Mini Kit; RNA concentration was determined using Nanodrop (Thermo Scientific, Waltham, MA); RT-PCR was performed on a GeneAmp PCR System (Applied Biosystems, Foster City, CA); and PCR products were analyzed by electrophoresis on 2% agarose gels stained with ethidium bromide.

Hormonal priming, antibody blocking, and in vitro fertilization
Female wild-type (WT), UPII KO, UPIIIa KO, and UPII- and UPIIIa double-KO mice were injected intraperitoneally with 10 IU of PMSG for 48 h and then 10 IU of hCG for 15 h before killing. IVF and in vitro culture IVF procedures were modified from (Hogan, 1994; Chang et al., 2005). Male mice were killed by cervical dislocation under anesthesia. The vas deferens was cut close to the cauda epididymis, and the sperms were extruded into a dish containing HEPES-HTF medium. The dish was then placed in a 37°C incubator for 0.5 h for sperm swimming out and capacitation. Freshly mature eggs were placed into plates containing 250-µl drops of high calcium (5.14 mM) RVF medium with 1 mM GSH covered by mineral oil. Sperm mixture (5–10 µl) was pipetted to each microdrop to a final concentration of (1–2) × 10⁶ sperm/ml. After 56 h of coincubation at 37°C, the two-cell stage embryos were collected and washed with M2 medium and then transferred to microdrop of KSOM and placed in a 5% CO₂ incubator at 37°C. The embryos were monitored daily and cultured to blastocyst stage. The mature eggs were collected from female CB6F1 mice injected with PMSG and HCG and treated with hyaluronidase (300 IU/ml) in HEPES-HTF medium to remove cumulus cells. After three washes, mature eggs were transferred into five drops of RVF medium, which contained the antibodies against UPIIa, UPIIb, UPIIIa, CD9, and normal rabbit immunoglobulin G (IgG) (200 µg/ml each), respectively, and incubated in the 5% CO₂ for 30 min at 37°C. The eggs then were washed three times in fresh drops of media, and regular IVF was performed as described above.

Collection of mouse tissues, oocytes, and eggs
Urinary bladder, eye, kidney, prostate, epididymis, stomach, ovary, testis, and oviduct with ovulated eggs were collected from Swiss Webster, CB6F1, UPII KO, and UPIIIa KO mice, fixed in 4% formaldehyde and embedded in paraffin, and sectioned (4 µm thick). Immature oocytes (Prophase I arrested) were obtained from the ovaries of female CB6F1 mice in HTF-BSA medium (modified from Kryzak et al. [2013]). The mature egg collection procedure was modified from those of Hogan (1994). The cumulus egg complexes were obtained from the oviduct of CB6F1 mice. Then cumulus cells were removed mechanically by pipetting vigorously in HEPES-HTF medium + hyaluronidase (300 IU/ml). The oocytes and eggs were transferred to acidic Tyrode’s solution, pH 2.5, for 1–2 min to remove the zona pellicula. The eggs were washed three times with HEPES-HTF medium and then fixed with 4% formaldehyde for 1 h at room temperature.

Parthenogenetic activation and inhibitor treatment of mouse and Xenopus eggs
The mouse experiments were done according to Cuthbertson (1983) with modification. Briefly, mouse eggs were treated to remove the zona pellucida and split into two groups. One group was kept in Quinn’s medium with 1% BSA (Quinn’s BSA), plus 5 µM SU6656, and the other group was kept in Quinn’s BSA plus dimethyl sulfoxide (DMSO). After incubation for 30 min at 37°C, half of the eggs from each group were transferred to fresh Quinn’s medium at 37°C, and the other half was exposed to 8% ethanol in Quinn’s medium at 37°C, 5% CO₂ for 6.5 min. The eggs were washed three times with Quinn’s medium and then fixed with 4% formaldehyde. The Xenopus fertilization experiments were done according to Sakakibara et al. (2005) with minor modifications. The eggs were divided into three groups. In Group A, the eggs were incubated with normal rabbit IgG (100 µg/ml) or affinity-purified xIIIa IgG in fertilization buffer plus 1% BSA for 30 min at room temperature, then washed with fertilization buffer three times. The eggs were then incubated with sperms for 10 min. In Group B, eggs were incubated with 500 mM H₂O₂ in fertilization buffer for 2 min at room temperature. In Group C, eggs were incubated with 2-µM SKI606 in fertilization buffer for 30 min at room temperature. Finally, some eggs were kept in fertilization buffer with DMSO as controls. After three washes, the eggs of Groups A and C were also treated with H₂O₂. All three groups of eggs were washed with fertilization buffer three times after the treatments and fixed with 4% formaldehyde.

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