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The water-absorption region of ventral skin of several semiterrestrial and aquatic anuran amphibians identified by aquaporins

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1Integrated Bioscience Section, Graduate School of Science and Technology, Shizuoka University, Shizuoka, Japan; 2Department of Biology, Faculty of Science, Shizuoka University, Shizuoka, Japan; 3Protein Purify Company, Gunma, Japan; and 4School of Dental Medicine, University of Nevada, Las Vegas, Nevada

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Ogushi Y, Tsuzuki A, Sato M, Mochida H, Okada R, Suzuki M, Hillyard SD, Tanaka S. The water-absorption region of ventral skin of several semiterrestrial and aquatic anuran amphibians identified by aquaporins. Am J Physiol Regul Integr Comp Physiol 299: R1150–R1162, 2010. First published September 1, 2010; doi:10.1152/ajpregu.00320.2010.—Regions of specialization for water absorption across the skin of Bufonid and Ranid anurans were identified by immunohistochemistry and Western blot analysis, using antibodies raised against arginine vasotocin (AVT)-stimulated aquaporins (AQPs) that are specific to absorbing regions of Hyla japonica. In Bufo marinus, labeling for Hyla urinary bladder-type AQP (AQP-h2), which is also localized in the urinary bladder, occurred in the ventral surface of the hindlimb, pelvic, and pectoral regions. AQP-h2 was not detected in any skin regions of Rana catesbeiana, Rana japonica, or Rana nigromaculata. Hyla ventral skin-type AQP (AQP-h3), which is found in the ventral skin but not the bladder of H. japonica, was localized in the hindlimb, pelvic, and pectoral skins of B. marinus, in addition to AQP-h2. AQP-h3 was also localized in ventral skin of the hindlimb of all three Rana species and also in the pelvic region of R. catesbeiana. Messenger RNA for AQP-x3, a homolog of AQP-h3, could be identified by RT-PCR from the hindlimb, pectoral, and pelvic regions of the ventral skin of Xenopus laevis, although AVT had no effect on water permeability. In contrast, 10−8 M AVT-stimulated water permeability and translocation of AQP-h2 and AQP-h3 into the apical membrane of epithelial cells in regions of the skin of species where they had been localized by immunohistochemistry and Western blot analysis. Finally, water permeability of the hindlimb skin of B. marinus and all the Rana species was stimulated by hydrons 1 and 2 to a similar level as seen for AVT. The present data demonstrate species differences in the occurrence, distribution, and regulation of AQPs in regions of skin specialized for rapid water absorption that can be associated with habitat and also phylogeny.

hindlimb skin; arginine vasotocin; immunohistochemistry; water permeability; frogs

Many adult anuran amphibians do not drink through their mouth. Rather, they absorb water across their skin and form dilute urine that is stored in their urinary bladder and can be reabsorbed when foraging away from a hydration source (4, 5). Hillman et al. (14) refer to this water balance strategy as semiterrestrial to distinguish it from terrestrial species that are completely independent of water. The semiterrestrial classification applies to tree frogs in the family Hylidae and toads in the family Bufonidae that have been traditionally classified as arboreal and terrestrial, respectively, in that both have large urinary bladder capacity and specializations for rapidly rehydrating when water is available. Specifically, they utilize an area of skin in the posteroventral region of the body that is specialized for rapid water absorption from shallow water sources or moist substrates. This region, termed the pelvic patch or seat patch, extends laterally to the ventral surface of the hindlimbs and shows a pattern of elevations and grooves termed verrucae hydrophilicae (14). The seat patch is also an area where capillaries form intimate contact with the basement membrane that underlies the epithelium. These structures are less well developed or absent in anuran species such as frogs in the family Ranidae that Hillman et al. (14) refer to as semiaquatic, because they are more dependent on a permanent source of water but may forage considerable distances in moist (mesic) habitats (10).

Literature values in vivo and with isolated skin indicate the semiaquatic Ranid species show a smaller response to arginine vasotocin (AVT) relative to the more xeric Hylid and Bufonid species (4) that have specialized regions for cutaneous water absorption. The mechanism for baseline and AVT-stimulated water permeability is controlled by aquaporins (AQPs), a class of integral membrane proteins that form selective water channels (pores) in the plasma membranes of various cells in virtually all organisms (1, 17, 23, 30). AVT stimulation results in the fusion of subapical vesicles containing the AQPs, with the apical membrane of epithelial tissues specialized for water absorption and reabsorption.

Recently, our group used RT-PCR to identify two forms of AVT-stimulated AQP in epithelial tissues of the skin of the tree frog, Hyla japonica: AQP-h2 was found in the urinary bladder, as well as the pelvic skin region (13, 20, 31). AQP-h2-like AQPs are also localized in the urinary bladder of Rana and Bufo species (20, 27) and have been referred to as the “urinary bladder-type” amphibian AQP (20). The second isoform, AQP-h3, was characterized from the ventral skin but not the urinary bladder of H. japonica (31). AQP-h3-like AQPs have similarly been characterized in ventral skin but not the bladder of Hyla, Bufo, and Rana species and have been termed the “ventral skin-type” AQP (20). This tissue distribution of AQP-h2 and AQP-h3 isoforms is provided in Table 1.

A subfamily of AQPs, termed aquaglyceroporins occurs in the basolateral membranes of water-transporting epithelia (1, 17) and is permeated by glycerol and urea in addition to water. In amphibians, these include AQP-h3BL, homologue of mammalian AQP3, in H. japonica (2), and AQP HC-3 in H. chrysoscelis (37). Such channels facilitate the movement of
water out of the cells into the vasculature, but the rate-limiting step is apical entry that is regulated by AVT and possibly other factors. The first objective of this study was to critically examine the relationship between AQP distribution in the apical membranes and AVT stimulation of water permeability in three zones (hindlimb, pelvic, and pectoral) of the ventral skin of toads (Bufo marinus) and frogs (Rana catesbeiana, Rana japonica, and Rana nigromaculata). This allowed a more detailed examination of the hypothesis put forth by Bentley and Main (4) that water permeability and its regulation by AVT differ among anuran species depending upon their habitat, and we suggest that phylogenetic factors should also be considered.

More recently, we have found AQP-x3 mRNA, that is homologous to AQP-h3, is expressed in the pelvic skin of the aquatic species, Xenopus laevis, but the mRNA is not translated to AQP-x3 protein (20). Given the different patterns of regional specialization seen in terrestrial, arboreal, and semiaquatic species, the second objective of this study was to examine the expression of AQP-x3 mRNA in the skin of the hindlimb, pelvic, pectoral, and dorsal regions.

In addition to AVT, hydrins are intermediate peptides derived from a provasotocin-neurophysin precursor. Like AVT, hydrins stimulate osmotic water movement across the skin and bladder but are devoid of antidiuretic activity in the kidney (19).

**MATERIALS AND METHODS**

**Animals.** Sexually mature Rana japonica, Rana nigromaculata, Rana catesbeiana, and X. laevis of both sexes were purchased from a commercial dealer (Ouchi, Saitama, Japan). Sexually mature Bufo marinus were collected in the field on Ishigaki Island, Okinawa. While X. laevis was kept in water, the other frogs and toads were in containers with a small amount of water for about 2 wk at 25–27°C before use. They were fed European crickets or trout pellets twice a week. All the experiments were performed from June to September. Skin for experimental treatments was removed under anesthesia (MS222; Nacalai Tesque, Kyoto, Japan). All animal experiments were carried out in compliance with the Guide for Care and Use of Laboratory Animals of Shizuoka University and approved by its ethics committee.

**Immunohistochemistry.** Ventral skin from the pectoral, pelvic, and hindlimb regions of B. marinus and the three Rana species was fixed overnight in periodate-lysine-paraformaldehyde (PLP) fixative, dehydrated, and embedded in Paraplast. Four-micrometer sections were cut and mounted on gelatin-coated slides. The deparaffinized sections were rinsed with PBS. For Rana species, the sections were incubated with rabbit anti-AQP-h3 (ST-141; 1:10,000, 31) and then reacted with Alexa Fluor 488-labeled donkey anti-rabbit IgG (1:200; Molecular Probes, Eugene, OR) containing 4’,6-diamidino-2-phenylindole (DAPI) for nuclear counterstaining. To investigate the specificity of the immunoreaction, we performed an absorption test by preincubating anti-AQP-h3 antibody with the antigen peptide (10 μg/ml). For B. marinus skin, double-immunofluorescence labeling was carried out according to Hasegawa et al. (13). In brief, specimens were also incubated with guinea pig anti-Hyla AQP-h2 serum (ST-140; 1:5,000, 13) followed by indocarbocyanine (Cy3)-labeled goat anti-guinea pig IgG (1:200; Jackson Immunoresearch, West Grove, PA). As with AQP-h3, specificity was performed by preincubating anti-AQP-h3 antibody with the antigen peptide (10 μg/ml). Specimens were examined with an Olympus BX61 microscope equipped with a BX-epifluorescence attachment (Olympus Optical, Tokyo, Japan).

**Western blot analysis.** Pieces of ventral skin from pectoral, pelvic, and hindlimb regions of B. marinus and the Rana species were homogenized in cell lysis buffer (50 mM Tris-HCl, pH 8.0, 0.15 M NaCl, 1% Triton X-100, 0.1 mg/ml PMSF, 1 mg/ml aprotinin) and homogenized in cell lysis buffer (50 mM Tris-HCl, pH 8.0, 0.15 M NaCl, 1% Triton X-100, 0.1 mg/ml PMSF, 1 mg/ml aprotinin) and centrifuged at 16,000 g in a microcentrifuge for 10 min to remove insoluble materials. The protein (10 μg) was denatured at 70°C for 10 min in denaturation buffer comprising 3% SDS, 70 mM Tris-HCl, pH 6.8, 11.2% glycerol, 5% 2-mercaptoethanol, and 0.01% bromphenol blue, subjected to electrophoresis on a 12% polyacrylamide gel, and then transferred to an Immobilon-P membrane (Millipore, Tokyo, Japan). The proteins on the membrane were reacted sequentially with rabbit anti-AQP-h3 serum (ST-141) or guinea pig anti-AQP-h2 serum (ST-140) followed by biotinylated goat anti-rabbit IgG (Jackson Immunoresearch, West Grove, PA) or biotinylated goat anti-guinea pig IgG (Jackson Immunoresearch). Immunopositive bands were visualized as reaction products of streptavidin-conjugated horseradish peroxidase (DAKO Japan, Kyoto, Japan) using an ECL Western Blot Detection kit (Amersham Pharmacia Biotech, Buckinghamshire, UK).

To check the specificity of the immunoreaction, we performed an absorption test by preincubating the antibodies with the antigen peptide (10 μg/ml).

**RT-PCR of Xenopus ventral skin.** RT-PCR for AQP-x3 was conducted for ventral pectoral, pelvic, hindlimb, and dorsal skin. TRIzol reagent (Invitrogen, Tokyo, Japan) was used to prepare total RNA from parts of ventral skin. Total RNA (20 μg) was first treated with DNase I (4 U; Takara, Kyoto, Japan), following which a 10-μl aliquot of the total RNA product was reverse transcribed with M-MLV-reverse transcriptase (Invitrogen) at 37°C for 1 h and then inactivated at 70°C for 30 min. The RT-PCR analysis was performed according to the method of Ogushi et al. (22), using the following primers: primer 1 (sense) GTGACACTAGCATTCCCTT (224–244 b) and primer 2 (antisense) GCAGCGGATGAATCATTC (537–557 b). *Xenopus* β-actin (accession no.: BC082343) primers were used as a control for the RT-PCR. β-actin sense primer was 5′-ACGTGACCTGACAGACTACC-3′ (579–598 bp) and the antisense, was CAGATTGCGATAGGTCCC-3′ (906–917 bp). The RT-PCR products were amplified in a 25-μl reaction mixture containing 5 μl of cDNA, 12.5 μl of 2× PCR master mix (Evrogen, Russia), 0.5 μl of each primer (10 μM), and 3 μl of water. All reactions were performed in a thermal cycler (Biometra, Germany), with the following program: 5 min at 94°C, 35 cycles of 30 s at 94°C, 30 s at 55°C, and 1 min at 72°C, followed by 7 min at 72°C.

**Table 1. Phylogenetic distribution of aquaporins in the ventral pelvic skins of anuran species living in different habitats**

<table>
<thead>
<tr>
<th>Habitat</th>
<th>Species</th>
<th>Pelvic Skin AQP-x3-Like Protein (Ventral Pelvic-Type)</th>
<th>Urinary Bladder AQP-h2-Like Protein (Urinary Bladder-Type)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Arboral</td>
<td><em>Hyla japonica</em></td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>Terrestrial</td>
<td><em>Bufo japonica</em></td>
<td>-</td>
<td>+</td>
</tr>
<tr>
<td>Semiaquatic</td>
<td><em>Rana catesbeiana</em></td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>Semiaquatic</td>
<td><em>Rana nigromaculata</em></td>
<td>-</td>
<td>+</td>
</tr>
<tr>
<td>Semiaquatic</td>
<td><em>Rana japonica</em></td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>Aquatic</td>
<td><em>Xenopus laevis</em></td>
<td>-</td>
<td>+</td>
</tr>
</tbody>
</table>

+, presence; –, absence. *Unpublished data
were analyzed on a 2% agarose gel containing etidium bromide (0.5 μg/ml).

The PCR products were purified following gel electrophoresis and sequenced using DNA sequencer [model Lic-4200L(S); Aloka, Tokyo, Japan].

Experimental protocol for water permeability. Water permeability was measured across isolated skin from the pectoral, pelvic, and hindlimb regions of B. marinus and the Rana species. Animals were immersed in water for 30 min before being killed to ensure a fully hydrated state for control observations (21). The isolated skin was washed with Ringer solution and then mounted between two chambers connected by an opening having a diameter of 1 cm. The chamber on the serosal side of the skin was filled with a Ringer solution (in millimoles per liter): 113 NaCl, 1.9 KCl, 1.1 CaCl2, 0.06 NaH2PO4, and 1.4 NaHCO3 having an osmolality of 220 mOsm/l, while the mucosal chamber was filled with water. Water movement from the mucosal to the serosal side of the skin was measured directly using a 0.1-ml pipette attached the serosal chamber of the apparatus (21). Water movement recorded over a 30-min period with the Ringer solution in the mucosal chamber followed by a 30-min period with Ringer solution containing 10−8 M AVT. After incubation, the skins were examined by immunofluorescence microscopy, as described above, to evaluate incorporation of AQP-h3 and AQP-h2 into the apical membrane of the first reacting cell (FRC) layer that is joined by tight junction to form a continuous barrier between the outside and inside of the body (6, 7, 11, 13, 35).

In a second set of experiments the effect of AVT on water permeability across the hindlimb skin only was compared with that to the same concentration (10−8 M) of hydrin 1 and hydrin 2. To verify that AVT-stimulated water flux was mediated by AQPs, the effect of HgCl2 on water transport across the hindlimb skins was measured. Skins were pretreated with 10−5 M AVT for 10 min to increase the number of AQPs inserted in the apical plasma membrane followed by treatment of the skins with 0.3 mM HgCl2 solution for 10 min. Water movement with continued AVT treatment was measured for an additional 30 min. Results from 5 or 6 individuals were expressed as means ± SE.

Statistical analysis. The data were compared by the Steel-Dwass’s test using PASW Statistics 18 software (SPSS, Chicago, IL). Statistical significance was considered as P < 0.05.

RESULTS

Characterization of aquaporins in three regions of ventral skin. Ventral skin from R. japonica and R. nigromaculata (Figs. 1A and 2A), showed immunoreactive staining for AQP-h3 in the principal cells of the hindlimb region (Figs. 1B and 2B). Nomarski images of the same sections (Figs. 1C and 2C) showed that intense labeling was predominantly found in the basolateral membrane of cells in the FRC. This positive reaction was abolished by preincubation with the antigen peptide (Figs. 1, D and E, and 2, D and E). Positive labeling was not visible in the pelvic (Figs. 1, F and G and 2, F and G) or pectoral (Figs. 1, H and I and 2, H and I) regions. Positive bands were seen in Western blots at ∼29 kDa with smearable bands of higher molecular mass in the hindlimb skin (Fig. 1, J, a, lane I and Fig. 2, J, a, lane I). A positive band was not detected in skin from the pelvic (Fig. 1, J, a, lane II and Fig. 2, J, a, lane II) or pectoral (Fig. 1, J, a, lane III and Fig. 2, J, a, lane III) regions. Positive bands were not seen when the antibody was preincubated with the antigen peptide (Figs. 1, J, b and 2, J, b).

In ventral skin of R. catesbeiana (Fig. 3A), the greatest number of labeled cells and the intensity of labeling with AQP-h3 antibody were similarly observed in principal cells of the hindlimb (Fig. 3, A–C), and the positive reaction was abolished by preincubation with the respective antigen peptide (Fig. 3, D and E). Unlike R. japonica or R. nigromaculata, a small number of principal cells in the pelvic skin were moderately labeled by the AQP-h3 antibody (Fig. 3, A, F, and G). For both hindlimb and pelvic skin, the label was localized in the basolateral plasma membrane. However, in the pectoral region, a positive reaction was observed as a dot spot only in the cytoplasm of a few principal cells in the FRC layer (Figs. 3, A, H, and I). When the extracts of the hindlimb, pelvic, and pectoral skins from R. catesbeiana were subjected to Western blot analysis, positive bands were found at ∼29 kDa (Fig. 3, J, a), although the intensity of the band decreased gradually from the hindlimb skin to the pectoral skin (Fig. 3, J, a, lanes I–III). These positive bands disappeared when antiserum was preincubated with the antigen (Fig. 3, J, b).

Ventral skin from B. marinus (Fig. 4A) showed immunopositive labeling for AQP-h3 and AQP-h2 in principal cells of the hindlimb (Fig. 4, B and D), pelvic (Fig. 4, F and H), and pectoral (Fig. 4, J and L) regions. The positive reactions in each region were abolished when the antibodies were preabsorbed with the respective peptides (Fig. 4, C, E, G, I, K, and M). Labeling was predominantly observed in the cytoplasm just beneath the apical membrane; however, the number of the principal cells labeled with anti-AQP-h3 and AQP-h2 varied among toads. In some toads, the number of cells immunopositive for AQP-h3 and AQP-h2 was high in all regions, while in other toads the greatest number of cells and intensity of labeling were observed in the hindlimb and pelvic skins, but less labeling was seen in the pectoral skin. Fig. 4 (B–M) illustrates a representative experiment in which the number of principal cells labeled for AQP-h3 and AQP-h2 decreased from the hindlimb skin to the pectoral skin. Western blot analysis revealed a major positive band for AQP-h3 at ∼29 kDa in the hindlimb, pelvic, and pectoral skin (Fig. 4, N, a, lanes I, II, and III). A smearable band at higher molecular mass was frequently observed as well. A major band for AQP-h2 protein with a smearable band at higher molecular mass was similarly observed at ∼29 kDa in the hindlimb, pelvic, and pectoral regions (Fig. 4, O, a, lanes I–III). The intensity of both the bands decreased from the hindlimb skin to the pectoral skin. These positive bands disappeared when the antiserum were preabsorbed with the respective antigen peptides (Fig. 4, N, b and O, b).

In X. laevis, we examined AQP-x3 mRNA levels in the same three regions of the ventral skin (Fig. 5A). Using RT-PCR, we were able to detect AQP-x3 mRNA expression in skin from the pectoral, pelvic, and hindlimb regions but not in dorsal skin (Fig. 5B). The cDNA fragments of AQP-x3 amplified by the RT-PCR showed a range of 223 to 555 bp in the full-length sequence (data not shown), indicating that the RNA amplified by the RT-PCR is a product of the AQP-x3.

Water permeability and dynamic movement of AQP-h3 and AQP-h2 immunopositive substances after stimulation of AVT and hydrins. Water permeability across the skin from the hindlimb, but not the pelvic or pectoral regions, of R. japonica and R. nigromaculata was significantly stimulated by AVT (Fig. 6, A and B). AVT stimulation of water permeability of skin from R. catesbeiana increased in order of the pectoral, pelvic, and hindlimb regions, but it was significantly greater in the hindlimb region, reaching levels similar to those of R. japonica and R. nigromaculata (Fig. 6C). AVT stimulation of
water permeability across the ventral skin of *B. marinus* was variable depending upon individuals and regions of ventral skin but was consistently above the control values (Fig. 6D). In three of the six toads, the response to AVT was greatest in the hindlimb skin and declined in the pelvic and pectoral skin, while in the other three, the response was greatest in the pelvic skin.

Following AVT treatment, hindlimb skin of *R. japonica* and *R. nigromaculata*, showed positive immunostaining for anti-AQP-h3 in the apical plasma membrane in the principal cells of the hindlimb skin.
FRC layer (Fig. 7, A and C). As before, Nomarski images were shown for each slide to identify the site of immunostaining (Fig. 7, B and D). For *R. catesbeiana*, the translocation of immunoreactive AQP-h3 protein to the apical plasma membrane of principal cells in the FRC layer was also greater in the hindlimb region and decreased in pelvic and pectoral regions that showed a lower stimulation of water permeability by AVT (Figs. 6C and 8). In *B. marinus*, the translocation of AQP-h3- and AQP-h2-positive proteins to the apical plasma membrane of principal cells in the FRC layer of the hindlimb, pelvic, and pectoral regions occurred in proportion to the stimulation of water permeability following AVT treatment (Fig. 9). Water permeability across *X. laevis* skin was not stimulated by AVT in any of the three regions (data not shown).
AVT, hydrid 1, and hydrid 2, significantly increased the water permeability of hindlimb skin (P < 0.05) in *R. japonica* > *R. nigromaculata* > *R. catesbeiana* > *B. marinus* (Fig. 10). There were no significant differences among the hormone responses within a given species (Fig. 10). The increased rates of water flux by AVT and its related peptides relative to the control values were 30–38 times in *R. japonica* (Fig. 10A), ~15 times in *R. nigromaculata* (Fig. 10B), 8–12 times in *R. catesbeiana* (Fig. 10C).
and 3 or 4 times in *B. marinus* (Fig. 10D). When hindlimb skin from each species was stimulated with AVT following the HgCl₂ treatment, the ratio of water flux significantly decreased compared with the AVT stimulation groups (Fig. 10).

**DISCUSSION**

Importance of the AQP-rich hindlimbs for water absorption by frogs and toads. Because skin from the hindlimbs was studied separately from the pelvic skin immediately anterior to the hindlimbs, the present study provides additional details regarding the specialization of ventral skin regions for regulated water absorption by anuran amphibians. Of particular interest, the area-specific rate of AVT-stimulated water flow across isolated skin of the hindlimbs was similarly large for the more mesic and xeric adapted frog and toad species studied. If anything, the response of toad skin was smaller than that of the frog species and tended to be highly variable. This is in contrast with the
study of Bentley and Main (4) that used isolated skin from the ventral posterior skin “near the hindlimbs” and found the magnitude of the AVT response to be greatest in *B. marinus*/*H. moori*/*R. pipiens*/*Neobatrachus pelobatoides*. These authors also showed that AVT stimulated water flow across the pectoral skin of *B. marinus* but not the other species. This is consistent with our observations that correlated AVT-stimulated water flow with the presence of an AQP-h2-like water channel in the pectoral region in addition to the pelvic and hindlimb regions of *B. marinus*. For *R. catesbeiana*, AQP-h3-like AQP was observed in the hindlimb and the more anterior pelvic and pectoral regions, while for *R. japonica* and *R. nigromaculata*, it was only observed in the hindlimb. The smaller response to AVT that was observed by Bentley and Main (4) for *Rana* vs. *Bufo* species could arise from skin obtained near the hindlimb that included a mixture of AQP-rich and AQP-poor regions in contrast with the more extensive distribution of AQPs in the toad skin. Thus, the greater response of *Bufo* vs. *Rana* species, in vivo, could be the result of the relative area of skin that contains AQPs rather than an area-specific response to AVT. As predicted, HgCl2 inhibited water flux across the hindlimb skin under AVT-stimulation, supporting our assumption that the measured water flux is mediated by mercury-sensitive AQP proteins.

**Possible expression of new-type AQPs in the ventral pelvic skin of Ranid species.** Western blot analysis of the extract of hindlimb skin from *R. japonica*, *R. nigromaculata*, and *R. catesbeiana* revealed a clear band detected at ~29 kDa, which is consistent with the molecular mass of AQP-x3 mRNA, the *Xenopus* ventral pelvic type-AQP, was expressed in the hindlimb, pelvic, and pectoral skins.

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**Fig. 5.** mRNA expression of the AQP-x3 in the hindlimb, pelvic, and pectoral skins in *Xenopus laevis*. A: ventral skin was divided into three regions: hindlimb (I), pelvic (II), and pectoral skins (III). B: RT-PCR products obtained using primers as described in Materials and Methods were separated on a 2% agarose gel and stained with ethidium bromide. – RT and water represent the negative control using the hindlimb skin sample without a RT reaction and with water instead of cDNA, respectively. β-actin was used as a positive control. A positive band of AQP-x3 mRNA, the *Xenopus* ventral pelvic type-AQP, was expressed in the hindlimb, pelvic, and pectoral skins.

**Fig. 6.** In vitro water permeability experiment using skin from the hindlimb, pelvic, and pectoral regions evaluated as water flux during 30-min periods before (open bars) and following (solid bars) treatment with $10^{-8}$ M AVT in the serosal solution. Bars with a different letter (a–c) are significantly different at $P < 0.05$. For *R. japonica* (A) and *R. nigromaculata* (B), AVT treatment significantly increased water permeability in the hindlimb skin, but not in the pelvic or pectoral regions. For *R. catesbeiana* (C), the greatest response to arginine vasotocin (AVT) was similarly observed in the hindlimb skin. Smaller responses were detected in the pelvic and pectoral skins. For *B. marinus* (D), the response to AVT was different from individual to individual, and even further from tissue to tissue. The same symbol is used to indicate data for each skin region from the same animal.
deduced from the amino acid sequences of AQP-h3 isoforms previously cloned for these species (20). It should be noted that AQP-h3 isoforms obtained from extracts of previously cloned for these species (20). It should be noted that the antibody raised against AQP-h3 from *H. japonica* is not immunoreactive with these oocyte extracts (unpublished observation) but does react in the present Western blot analysis of tissue homogenates and immunohistochemically labels hindlimb skins of the three frog species. We have recently cloned a new AQP from the ventral pelvic skin of *R. japonica* that has a higher homology (~85%) with AQP-h3 from *H. japonica*. Extracts of *Xenopus* oocytes injected with cRNA for this AQP-h3-like protein are immunoreactive with the anti-AQP-h3 antibody (unpublished observation). Consequently, the ventral pelvic skin-type AQPs detected by anti-AQP-h3 may be different from those reported previously by Ogushi et al. (20). However, this antibody-specificity was also proven by the absorption test in the present immunohistochemistry and Western blot analyses.

**Physiological and behavioral variables that can affect water absorption.** Because our experiments were conducted during the normal summer activity period on freshly collected animals that were kept with water available ad libitum, it seems unlikely that seasonal variability was a factor in the responses of the different species to AVT. It should be noted that Bentley and Main (4) did not report on the seasons during which their study was conducted. Seasonality in the stimulation of water absorption by AVT has been reported in spadefoot toads, *Scaphiopus couchi*, which remain dormant in burrows during much of the year and only respond to AVT during the brief period of emergence during summer rainfall in the southwestern United States (15).

The highly variable values for area-specific water flux across isolated *B. marinus* skin could result from a greater dependence on vascular perfusion relative to the thinner frog skin. Christensen (9) found that AVT stimulation of water flux across isolated skin from *B. bufo* required the cutaneous vasculature to be perfused with a Ringer solution. The tissue electrolyte content was diluted in nonperfused skin, indicating the circulation removed water as it was absorbed to prevent diminution of the osmotic gradient across the epidermis. Water absorption by anurans occurs in conjunction with a behavior, termed the water absorption response (16, 26), during which the skin is pressed to a moist surface and there is a large increase in blood flow to the absorbing area of the seat patch (33, 34), in conjunction with the insertion of AQPs into the apical membranes of the FRC layer of the skin (27). A better understanding of adaptations for semiterrestrial vs. semiaquatic species will require an understanding of all three parameters from a physiological and phylogenetic perspective.

**Phylogenetic significance on the presence of AQPs of ventral pelvic skin.** Both AQP-h2 and AQP-h3 isoforms are anuran specific AQPs that are structurally distinct from mammalian AQP2 but similarly become inserted into the apical membranes of epithelial cells by antidiuretic hormone stimulation (27). The species differences in the expression and localization of AQP-h2- and AQP-h3-reactive AQPs in the skin raise interesting hypotheses in the context of recent phylogenetic analyses of anuran taxa. On the basis of analysis of morphological and molecular data, Frost et al. (12) suggested that the radiation of anuran families giving rise to modern species occurred when Pangea was breaking up into what would become the current continental land masses, during the Jurassic and Cretaceous periods. The largest superfamilies of extant anurans are the Hylidae that includes modern Hylid and Bufonid species, among others, and the Ranidae that includes Ranid species, among others (12, 14, 25). Both *Hyla* and *Bufo* species examined to date have more specialized “seat patch” regions of skin and express AQP-h2-like proteins in the skin and bladder,
indicating this to be an apomorphic character. Further experiments are needed to compare the distribution of this trait and determine its synapomorphic distribution in other families of the Hyloidea.

AQP-h3 has been found in the ventral skin of Bufonid, Hylid, and Ranid species (20, 27), suggesting this pattern of expression is a more pleisiomorphic characteristic. As noted above, we observed AQP-h3 immunoreactivity to be present in the pelvic, as well as hindlimb, skin of *R. catesbeiana*, while it was only present in the hindlimbs of *R. japonica* and *R. nigromaculata*. Recent phylogenetic studies (12) have also noted differences between Ranid species to the point that many taxonomists have reclassified *R. catesbeiana* as *Lithobates catesbeiana*. The North American leopard frog, *Rana pipiens* has similarly been reclassified in the genus *Lithobates* and, like *R. catesbeiana*, the water permeability of its pelvic skin was observed to be modestly stimulated by AVT (4). So-called “old-world” Ranid species, including *R. japonica* and *R. nigromaculata* remain in the genus *Rana*. As with the tissue distribution of AQP-h2-like AQPs, future studies are needed to better determine whether the spatial localization of AQP-h3-like AQPs corresponds with recent advances in phylogenetic analysis of species that have historically been classified as terrestrial, arboreal, and semiaquatic.

The expression of two AVT-stimulated AQPs in the skin of *Bufo* and *Hyla* species. An AQP-h2 homolog has been detected in the urinary bladder of all species examined but only in the skin of *Hyla* and *Bufo* species (Table 1). In contrast, mRNA encoding an AQP-h3 homolog has been identified in the skin but not the urinary bladder of all species examined (20, 28). Molecular diversity of vasotocin-dependent AQPs closely associated with water adaptation strategy in anuran amphibians (29). Thus, anurans possess both AQP-h2- and AQP-h3-like AQPs, which respond to AVT, like mammalian AQP2, but are classified into an anuran-specific type, AQPα2 by phylogenetic analyses (27, 28). For *Xenopus tropicalis*, whose genome data are available in the Ensembl genome browser (http://www.ensembl.org/index.html), two types of AQPα2 genes, i.e., AQP-x2 (ENSXETG00000024581), and AQP-x3 (ENSXETG00000013389), are sited with AQP5 between *FAIM2* (Fas apoptotic inhibitory molecules 2) and *RACGAP1* (Rac GTPase-activating protein 1), as are the human genes for AQP2, AQP5, and AQP6 on chromosome 12. On the basis of these findings and the close examination of *AQP* genes in the genome of fishes (medaka and pufferfish), it is likely that h2- and h3-like AQPα2 genes had been generated by local gene duplication of AQP2 in the amphibian lineage to anurans (28). For contemporary anurans, the upstream regions of these AQPα2 genes seem to be differentiated so that h2-like AQPα2
basically occurs in the urinary bladder, whereas h3-like AQPa2 is expressed in the ventral skin. In *Bufo* and *Hyla* species, the upstream region of h2-like AQPa2 gene is considered to have undergone a further change to express this gene in the ventral skin, as well as in the urinary bladder, which might give the terrestrial species an advantage with respect to cutaneous water absorption and thereby in adapting to drier environments.

A unique AQP in aquatic Xenopus. The purely aquatic species in the family, Pipidae, are believed to have diverged from other anurans as far back as the Permian period (12). *X. laevis* is a commonly used species for a variety of molecular and developmental studies but lacks a hydroosmotic response to AVT (3, 4, 36). Nonetheless, we have identified mRNA of AQPa-x3 in the pelvic skin of *X. laevis* that is homologous to AQPa-h3, but contains an extra C-terminal tail, consisting of 33 nucleotides within the coding region, which appears to attenuate translation (22). Despite the lack of functional expression of the protein, we have shown mRNA for AQPa-x3 to be present in all three regions of the ventral skin. In their native habitat, ponds may dry up and leave the animals dependent on substrate moisture (32). Under these conditions, complete processing of the mRNA and expression of AQPa-x3 could be advantageous, but we are unaware of data to support this possibility.

Regulation of AQP expression by AVT and its related peptides. In the present study, we found that hydrin 1 and hydrin 2 stimulated the water permeability of the hindlimb skin of *Bufo* and *Rana* species at a level equivalent to that of AVT. The $K_m$ for cAMP production by tree frog V2-type AVT receptor expressed by transfected CHO-KI cells was 3.6, 8.6, and 13.5 nM for AVT, hydrin 1, and hydrin 2, respectively (18), suggesting they may share a common receptor. Both peptides are generated from a down-regulation in posttranslational processing, i.e., hydrin 1 by a decrease in carboxypeptidase E activity and hydrin 2 by a reduction in the activity of the $\alpha$-amidating enzymatic system. Of interest, *X. laevis* secretes hydrin 1, as well as AVT but shows no hydroosmotic response of skin to either. However, AVT and hydrin 1 both stimulate water reabsorption from the urinary bladder and may be involved in the water balance in *X. laevis* during aestivation (24).

Fig. 9. Immunofluorescence images of the ventral skin of *B. marinus* following in vitro challenge with $10^{-6}$ M AVT. The label for AQP-h3 (green) and AQP-h2 (red) is visible in the apical membrane (arrows) of principal cells in the first-reacting cell layers of the hindlimb skin (A, B). Similar results were obtained in the pelvic (D, E) and pectoral (G, H) skin, to those in the hindlimb skin, although fewer positive principle cells were labeled. The corresponding Nomarski images (C, F, and I) are to A–B, D–E, and G–H, respectively. The box in each figure is the enlarged image. Nuclei are counterstained with DAPI (blue). Scale bar = 10 $\mu$m.
Perspectives and Significance

Anuran amphibians have two AQP isoforms that are stimulated by AVT to increase water absorption across the skin and reabsorption from the urinary bladder. Suzuki and Tanaka (28) suggested the two isoforms are derived from gene duplication of AQP2 in an ancestral amphibian and are homologs of AQP-h2 and AQP-h3 that were first characterized in *H. japonica*. Amniotes, including mammals, are thought to have evolved from a primitive amphibian ancestor in the carboniferous (8) and retain AQP2 as a single isoform regulated by antidiuretic hormone, while amphibian evolution has proceeded independently. Given the importance of regulated water permeability in amphibian skin and bladder, the two isoforms have retained the same function in the course of evolution.

Within the limited number of species examined, anurans in the superfamily Hyloidea express both isoforms in the ventral skin and have a specialized seat patch region that is also highly vascularized. Anurans in the superfamily Ranidae lack specialization in the pelvic region and express only AQP-h3, primarily in the ventral surface of the hindlimbs. The aquatic frog, *X. laevis* transcribes mRNA for homologs of both isoforms but a C-terminal sequence prevents translation. However, all species examined to date express AQP-h2-like AQPs in the urinary bladder. Future studies are needed to examine species differences in the functional expression of AQP-h2 and AQP-h3 to characterize phylogenetic relationships associated with water balance adaptations among anuran families.

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DISCLOSURES

No conflicts of interest, financial or otherwise, are declared by the authors.
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