

The Roles of Aquaporins in Inflammatory and Ischemic Diseases

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Abstract : Aquaporins (AQPs) are membrane proteins that function as a water channel and, in some cases, also transport small solutes such as glycerol and CO₂. AQPs are expressed in various tissues such as the kidney, lung, brain, skin, and glandular epithelia. Abnormalities of water metabolism are one of characteristics of inflammatory and ischemic diseases. However, their molecular mechanisms are poorly understood. Pathophysiological stimuli modulate the AQP expression level in inflammatory and ischemic tissues. This short review summarizes the pathophysiological roles of AQPs in abnormal water metabolism of lung, skin, and kidney, based on studies with AQPs-deficient mice.

Key words : Aquaporin, Gene Expression, Inflammation, Ischemia/Reperfusion, Water Metabolism

Introduction

Water constitutes about 70% of our body mass and the appropriate regulation of water transport is therefore necessary to maintain a sufficient water balance in various organs. Aquaporins (AQPs) are a family of small (about 30 kDa), hydrophobic, integral membrane proteins that dominantly function as water channels¹⁾. AQPs facilitate trans-epithelial water transport in response to osmotic gradients such as fluid absorption by kidney proximal tubule and fluid secretion in the salivary gland. Some AQPs also transport small solutes such as glycerol and CO₂ across the plasma membrane. Phenotype studies of mice lacking each AQP suggest that the physiological functions of AQPs are involved in urinary concentration, glandular fluid secretion, and maintenance of water balance in the brain and skin. Therefore, AQPs may be therapeutic targets for abnormalities of water metabolism.

The dysregulation of the water metabolism is observed in inflammatory and ischemic diseases. For example, inflammation in the lung induces lung edema which is an end-stage feature in alveolus, and mucus retention (low water content in mucus) in the airway. In addition, brain edema, which is most frequently associated with the destruction of water homeostasis, is caused by a brain

stroke. Therefore, inflammation and ischemia are the main causes of an abnormal water metabolism, although their molecular mechanisms are still poorly understood. This review focuses on the pathophysiological roles of AQPs in lung, skin, and kidney, based on studies with AQPs-deficient mice.

Physiological functions of AQPs

AQPs are constitutively expressed in the plasma membranes of various organs (Table 1). Thirteen AQPs (AQP0-12) have been identified in mammals and divided into two groups on the basis of their permeability¹⁾⁻⁴⁾.

Table 1. Tissue distribution and permeability of human aquaporins.

AQP	Tissue expression	Permeability
AQP0	Lens	
AQP1	Kidney, endothelium, erythrocyte and lung	Water
AQP2	Kidney	Water
AQP3	Skin, kidney and lung	Water and glycerol
AQP4	Brain, kidney and muscle	Water
AQP5	Lung, salivary, lacrimal and sweat glands	Water
AQP6	Kidney	
AQP7	Adipose tissue, testis and kidney	Water and glycerol
AQP8	Liver, pancreas, intestine and testis	Water
AQP9	Liver, leukocytes, brain and testis	Water
AQP10	Small intestine	Water and glycerol
AQP11	Kidney and liver	
AQP12	Pancreas	

AQP1, 2, 4, 5 and 8 are selectively permeable to water. Computer simulations based on the AQP1 crystal structure suggest the single-file passage of water through a narrow pore within the molecule⁵. On the other hand, AQP3, 7, 9 and 10 transport not only water but also glycerol, therefore, they are called ‘aquaglyceroporins’.

AQPs have no gating system in water permeable pores. Therefore, functions of AQPs depend on their expression level in the plasma membrane. For example, AQP2, which normally exists in intracellular storage vesicles of the collecting duct cells, translocates to the apical plasma membrane in response to vasopressin stimulation, leading to water reabsorption for urine concentration⁶. On the other hand, the expression level of other AQPs is regulated by changing the transcriptional-level. NF-κB, a pro-inflammatory transcription factor, decreases AQP5 gene transcription in lung epithelial cells⁷.

Phenotype studies of AQPs-deficient mice suggest multiple physiological roles of AQPs. Deficiency of AQP1 and AQP2, which are expressed in the kidney, caused polyuria in mice due to abnormal urinary concentration^{8,9}. In addition, AQP2 mutations in humans cause congenital nephrogenic diabetes insipidus (NDI). NDI is characterized by the inability to concentrate urine despite a normal level of vasopressin in plasma, resulting in a massive loss of water from the kidney¹⁰⁻¹². A study with AQP5-deficient mice revealed that AQP5, which is expressed in salivary glandular epithelial cells, is thought to play an important role in the secretion of saliva¹³.

Roles of AQP1 and AQP5 in the lung

AQP1 and AQP5 are main subtypes in lung. AQP1 is expressed in vascular endothelial cells and AQP5 is expressed in type 1 alveolar epithelial cells^{14,15}. Double knockout mice for AQP1 and AQP5 show decreased alveolar-capillary water permeability¹⁶. Therefore, AQP1 and AQP5 play critical roles in the regulation of water content in the alveolus. Lung inflammation, such as adenoviral infection and acute lung injury is a respiratory dysfunction, which finally induces pulmonary edema. Pulmonary edema is destruction of water homeostasis in the alveolus. Intratracheal infection with adenovirus, which causes airway inflammation and pulmonary edema, decreases the expression levels of AQP1 and AQP5 in the lung¹⁷. In addition, pro-inflammatory cytokines such as a tumor necrosis factor-α (TNF-α) also decrease AQP5 expression in lung epithelial cells¹³. These findings suggest that decreases in AQP1 and AQP5 expression in

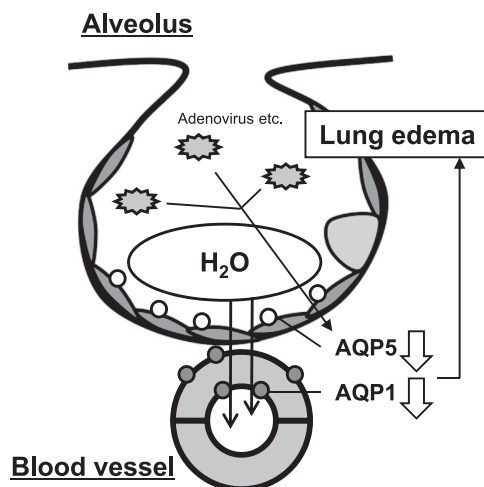


Figure. 1 Schematic model of down-regulation of AQP1 and AQP5 in lung inflammation.

Inflammatory stimuli (adenovirus infection, LPS, and inflammatory cytokines) decrease the AQP1 and AQP5 expression level in the alveolus. Inflammation-induced decreases in AQP1 and AQP5 reduce alveolar-capillary water permeability and subsequently cause pulmonary edema.

the lung induce low water excretion from the alveolus and finally cause pulmonary edema in lung inflammation (Fig. 1). Interestingly, AQP1 is also permeable to CO₂ gas^{18,19}. Therefore, the AQP1 and AQP5 expression levels may contribute not only to the dysregulation of water homeostasis in the alveolus, but also to an impairment of gas exchange in lung inflammation.

AQP5 is also expressed in airway epithelial cells. The airway has a lot of water, so-called ‘airway-surface liquid (ASL)’. ASL is a first defense system which removes pathogens and viruses and maintains airway humidity. However, a decrease in ASL induced by airway inflammation such as cystic fibrosis (CF) causes mucus retention (low water content in mucus). Decreases in AQP5 expression in the airway may reduce water secretion to the airway lumen and decrease ASL volume. Therefore, AQP5 may also play an important role in the pathophysiology of airway inflammation.

Role of AQP3 in the skin

Impaired hydration of the stratum corneum, so-called ‘dry skin’, is a characteristic feature of inflammatory skin diseases, such as atopic dermatitis, eczema, psoriasis, senile xerosis, and hereditary ichthyosis²⁰⁻²⁴. Low hydration of the stratum corneum is frequently correlated to disruption of the skin barrier function²⁵ and probably

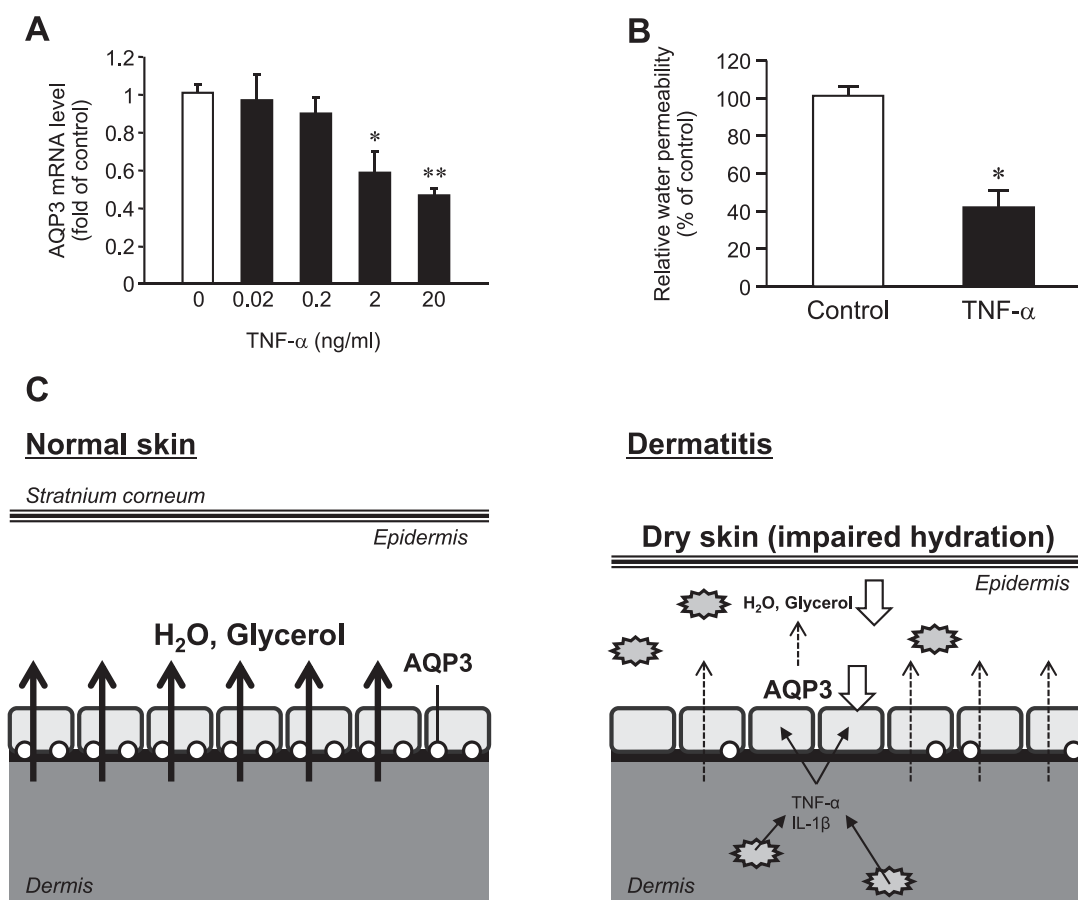


Figure 2 Decrease of AQP3 expression induced by inflammatory cytokines in the skin.
 A) TNF- α decreased AQP3 mRNA expression in human keratinocytes cell lines.
 B) TNF- α decreased water permeability of plasma membrane in human keratinocytes cell lines.
 C) A schematic model of down-regulation of AQP3 in skin inflammation.

influences epidermal differentiation and wound repair. AQP3 is the major subtype in skin and is specifically expressed in the basal layer of keratinocytes²⁶⁾. Impaired stratum corneum hydration, reduced skin elasticity, and delayed recovery of barrier function after removal of the stratum corneum are observed AQP3-deficient mice^{27), 28)}. However, the molecular mechanisms that control the AQP3 expression level in inflammatory skin are poorly understood. TNF- α decreases AQP3 expression in keratinocytes and this reduction affects water permeability in the plasma membrane (Fig. 2)²⁹⁾. AQP3 down-regulation may also affect glycerol transport. Keratinocytes-mediated water and glycerol transport are important for maintaining hydration of the stratum corneum. Therefore, impaired stratum corneum hydration due to skin inflammation may be caused by a decrease in AQP3 expression in keratinocytes.

Skin inflammation causes a delay of wound-healing.

Interestingly, delayed wound-healing is observed in AQP3-deficient mice²⁸⁾ and keratinocytes derived from AQP3-deficient mice have low migration activity, which causes delayed wound-healing^{30), 31)}. Inflammation-induced decrease in AQP3 expression in keratinocytes may affect the wound-healing activity in skin. Therefore, the AQP3 expression level is important for the skin barrier function via water homeostasis and epidermis regeneration and AQP3 plays a role in the pathophysiology of skin inflammation.

Role of AQPs in the kidney

AQP1, AQP2, and AQP3 are the main subtypes in the kidney, although other AQPs (AQP4, 6, 7, and 8)¹²⁾ are also expressed (Table 2). AQP1 is highly abundant in the proximal and descending thin limb and plays a critical role in the constitutive re-absorption of water in these segments and their role in urinary concentration³²⁾. In

Table 2. Subcellular localization and regulation of renal aquaporins.

AQP	Distribution	Subcellular localization	Regulation	Functions
AQP1	Proximal tubules	Apical and basolateral membrane	Not regulated by vasopressin	Reabsorption of 80% of water in filtrate secondary to solute reabsorption
	Descending thin limb of loop of Henle Vascular endothelial cells			
AQP2	Collecting duct (principle cells)	Apical plasma membrane and apical vesicles	Regulated by vasopressin	Water reabsorption from distal collecting duct
AQP3	Collecting duct (principle cells)	Basolateral membrane	Regulated by vasopressin	Water reabsorption from distal collecting duct
AQP4	Collecting duct (principle cells)	Basal membrane	Regulated by vasopressin	Water reabsorption from distal collecting duct
AQP6	Collecting duct (intercalated cells)	Intracellular vesicles	?	(Intracellular water and ion channels)
AQP7	Proximal tubules in segment 3	Apical brush border	?	?
AQP8	Proximal tubules collecting duct cells	Intracellular vesicles	?	?

addition, AQP2, AQP3, and AQP4 are expressed in the collecting duct. AQP2 is found in the apical membrane in the principal cells of the collecting duct. On the other hand, AQP3 is found in the in basolateral membrane. Therefore, AQP2 and AQP3 may be involved in water reabsorption from the collecting ducts.

Acute kidney injury (AKI) caused by ischemia/reperfusion (I/R) injury is a common clinical problem associated with a high morbidity and mortality. I/R injury may occur in hypo-perfusion following kidney transplantation, partial nephrectomy, aortic cross-clamping and shock³³. Renal expression levels of AQP1, AQP2, and AQP3 are markedly reduced after renal I/R injury^{6, 34, 35}. Mice deficient for AQP1, AQP2, and AQP3 showed an abnormal urine concentration and subsequently polyuria. Therefore, renal I/R-induced decrease in the expression of AQPs may be related to the pathophysiology in AKI.

Urinary concentration and tubular sodium reabsorption are markedly impaired in post-ischemic kidneys. Intriguingly, treatment with α -melanocyte stimulating hormone (α -MSH), a potent anti-inflammatory agent, inhibits the reduction in renal AQPs (AQP1, AQP2, and AQP3) and sodium transporter after renal I/R³⁶. α -Lipoic acid, which is a potent antioxidant, and erythropoietin (EPO), which is primarily released from renal cortical fibroblasts in response to hypoxia, represses the down-regulation of renal AQP2 and AQP3 expression^{37, 38}. In fact, α -MSH, α -lipoic acid, and EPO improve urinary concentration ability after I/R injury. These results

suggest that the pharmacological modulators of AQPs expression in the kidney are therefore novel drug candidates for renal diseases, such as AKI.

Conclusion

Phenotype studies with AQP-deficient mice suggest that AQPs may be involved in abnormalities of water metabolism in various tissues with inflammation and ischemia. Inflammatory stimuli (adenovirus infection, LPS, and inflammatory cytokines) decreased AQP1, AQP3, and AQP5 expression in mammalian tissues. Although it is necessary to obtain more information about the AQP expression level under specific pathophysiological conditions, inflammation-induced down-regulation of AQPs expression may cause dysfunction of water homeostasis in various organs. Therefore, the modulators of AQPs expression may have a therapeutic potential in inflammatory and ischemic diseases.

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