

REVIEW

Open Access



Function of aquaporins in sepsis: a systematic review

Katharina Rump*  and Michael Adamzik

Abstract

Background: Sepsis is a common cause of death in intensive care units worldwide. Due to the high complexity of this immunological syndrome development of novel therapeutic strategies is urgent. Promising drug targets or biomarkers may depict aquaporins (AQPs) as they regulate crucial key mechanisms of sepsis.

Main body: Here we report on base of the current literature that several AQPs are involved in different physiological processes of sepsis. In immune system mainly AQPs 3, 5 and 9 seem to be important, as they regulate the migration of different immune cells. Several studies showed that AQP3 is essential for T cell function and macrophage migration and that AQP5 and AQP9 regulate neutrophil cell migration and impact sepsis survival. Additionally, to the function in immune system AQPs 1 and 5 play a role in sepsis induced lung injury and their downregulation after inflammatory stimuli impair lung injury. By contrast, AQP4 expression is up-regulated during brain inflammation and aggravates brain edema in sepsis. In kidney AQP2 expression is downregulated during sepsis and can cause renal failure. Some studies also suggest a role of AQP1 in cardiac function.

Conclusion: In conclusion, AQPs are involved in many physiological dysfunctions in sepsis and their expressions are differently regulated. Additional research on the regulatory mechanisms of aquaporins may identify potential therapeutic targets.

Keywords: Aquaporin, AQP, Expression, Immune cells, Migration, Brain, Kidney, Liver, Lung, Heart, LPS, sepsis

Background

Sepsis is one of the most common complications in Intensive Care Units in Germany and the United States [1, 2], and mortality remains unrestrainable high due to the extreme complexity of this immunological syndrome. Predictive biomarkers which characterize this immunological syndrome properly are still missing; hence no individual therapy adapted on the immune status of the unique patient can be conducted. Aquaporins might be convenient biomarkers because they play an important role in inflammation and especially in sepsis as revealed by experimental and association studies [3–6].

Aquaporins (AQPs) are a group of to date 13 identified membrane proteins, which are essential for the regulation

of water and salt in- and out flux of the cell. In addition, some AQPs facilitate the passive transport of glycerol and other small solutes such as urea and carbon dioxide through the cell membrane [7]. The water-selective AQPs are involved in many biological functions, including trans-epithelial fluid transport, cell migration, brain edema and neuroexcitation [7], whereas the aquaglyceroporins participate in cell proliferation, adipocyte metabolism and epidermal water retention. With this study we want to elucidate the possible role of AQPs in pathomechanisms of sepsis on base of the current literature.

Approach of literature research and methodology

A literature search was undertaken using various online sources of English journal articles including Science-Direct, PubMed and Web of Science. The keywords “aquaporin AND sepsis”, “aquaporins AND sepsis” and “AQP(xy) AND sepsis” were used to search all relevant articles dealing with the role of aquaporins in sepsis. In

*Correspondence: katharina.k.rump@rub.de

Klinik für Anästhesiologie, Intensivmedizin und Schmerztherapie, Universitätsklinikum Knappschaftskrankenhaus Bochum-Langendreer, Ruhr-Universität Bochum, In der Schornau 23-25, 45882 Bochum, Germany

total 51 studies were found. 10 articles were excluded because they either did not deal with sepsis or with aquaporins. One article was excluded because it was in Russian. The workflow of literature research can be found in Fig. 1. Due to the relative low number of articles dealing with real bacterial sepsis models, endotoxemia models using LPS injection were included in the analysis.

Aquaporin expression during inflammation

To completely understand the role of AQP in sepsis, it is important to know how their expression is altered during inflammation. It was demonstrated that in leucocytes of septic patients *AQP3* expression is reduced 2.5 [8] fold and that simultaneously *AQP1* expression is increased twofold [8]. In line with this our group showed that *AQP1* expression is increased in the monocytic cell line THP-1 after lipopolysaccharide (LPS) administration, but *AQP5* mRNA expression is reduced [9]. *AQP6* expression in contrast might play a role in viral infections as it is decreased after viral infection and in turn can reduce the infectivity of Hazara virus [10]. Furthermore, *AQP8* is reduced in hepatocytes after LPS administration [11]. In addition, patients with systemic inflammatory response syndrome (SIRS) show increased *AQP9* expression in neutrophils compared to healthy controls [12]. Moreover, Gram-negative bacteria as *P. aeruginosa* induce increased expression, distribution and re-organization of *AQP9* in macrophages with is accompanied by changes in macrophage size and morphology. This in turn affects motility, migration and phagocytosis [13].

Aquaporins in cell migration of immune cells

The importance of aquaporins in cell migration has been demonstrated several times before [7, 14–17].

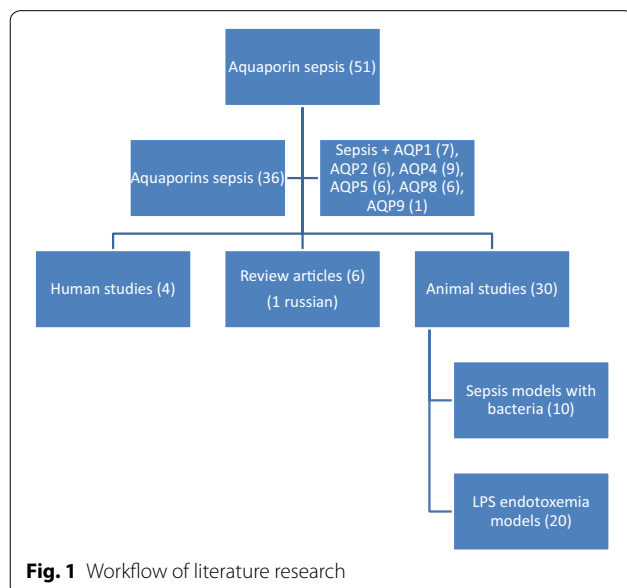
The proposed mechanism by which AQP enhance cell migration is that they facilitate water influx at the cell's leading edge. This causes membrane expansion and formation of a concentration gradient of actin polymers which is followed by actin repolymerization to stabilize the membrane protrusion and lamellipodia formation [17]. As immune cell migration is an essential mechanism in sepsis, AQP might depict key players in this process. Considerable AQP for immune cell migration are AQP1, AQP3, AQP5, AQP7 and AQP9 as they are expressed in activated B and T lymphocytes (AQP1, 3, 5) [15] as well as immature dendritic cells (AQP3, 5, 7) [15] and neutrophils (AQP9) [15, 18, 19] (Fig. 2f).

AQP5 seems to be of special interest, because in the past our group demonstrated that the C-allele of the functional *AQP5* A(-1364)C promoter polymorphism (rs3759129) is associated with increased survival in severe sepsis [3] but decreased *AQP5* expression [20]. Recently we showed that *Aqp5*-knockout (KO) mice show increased survival compared to wildtype mice after LPS induced endotoxemia. Furthermore, *AQP5* overexpression caused increased migration of the T-lymphocytic cell line Jurkat. In addition, neutrophil granulocytes from C-allele carriers showed decreased migration compared to A-allele carriers. Therefore we concluded that the *AQP5* genotype and *AQP5* protein expression seem to alter neutrophil cell migration and may influence survival in sepsis by altering neutrophil cell migration. Hence *AQP5* might be a key protein in inflammation and depict a novel target for developing sepsis therapeutics [21].

Similar to our study Zhu et al. analyzed the effects of *Aqp3* expression in a sepsis mouse model. They found that mouse resident peritoneal macrophages (mRPMs) express the aquaglyceroporin *Aqp3* and to a low extent *Aqp7* and *Aqp9* in a plasma membrane pattern [22]. In contrast to our study, *Aqp3*-KO mice show significantly greater mortality than wildtype mice in a model of bacterial peritonitis. In addition, *Aqp3*-KO is accompanied by reduced migration of macrophages [22]. Besides to macrophage function, *AQP3* seems also to be crucial for T-cell migration. It is suggested that *AQP3*-mediated H₂O₂ uptake is required for chemokine-dependent T-cell migration and a sufficient immune response [23].

AQP4 plays a role in the development of regulatory T-cells in the thymus. *Aqp4*-KO mice show decreased levels of CD4+ CD25+ regulatory T-cells. The decreased amount of regulatory T-cells causes increased microglial inflammatory response in a mouse model of Parkinson with *Aqp4*-KO mice [24].

Similar to the role of *AQP5* and *AQP3*, *AQP9* seems to be responsible for neutrophil migration, as *Aqp9*-KO mice show reduced neutrophil migration to fMLP [25]. In addition, *Aqp7*-KO mice have reduced migration of



cutaneous dendritic cells. Beside its role in cell migration AQP7 seems to be responsible for antigen uptake as it could be demonstrated that *Aqp7*-deficient DCs showed a decreased cellular uptake of low-molecular-mass compounds and high-molecular-mass substances [19].

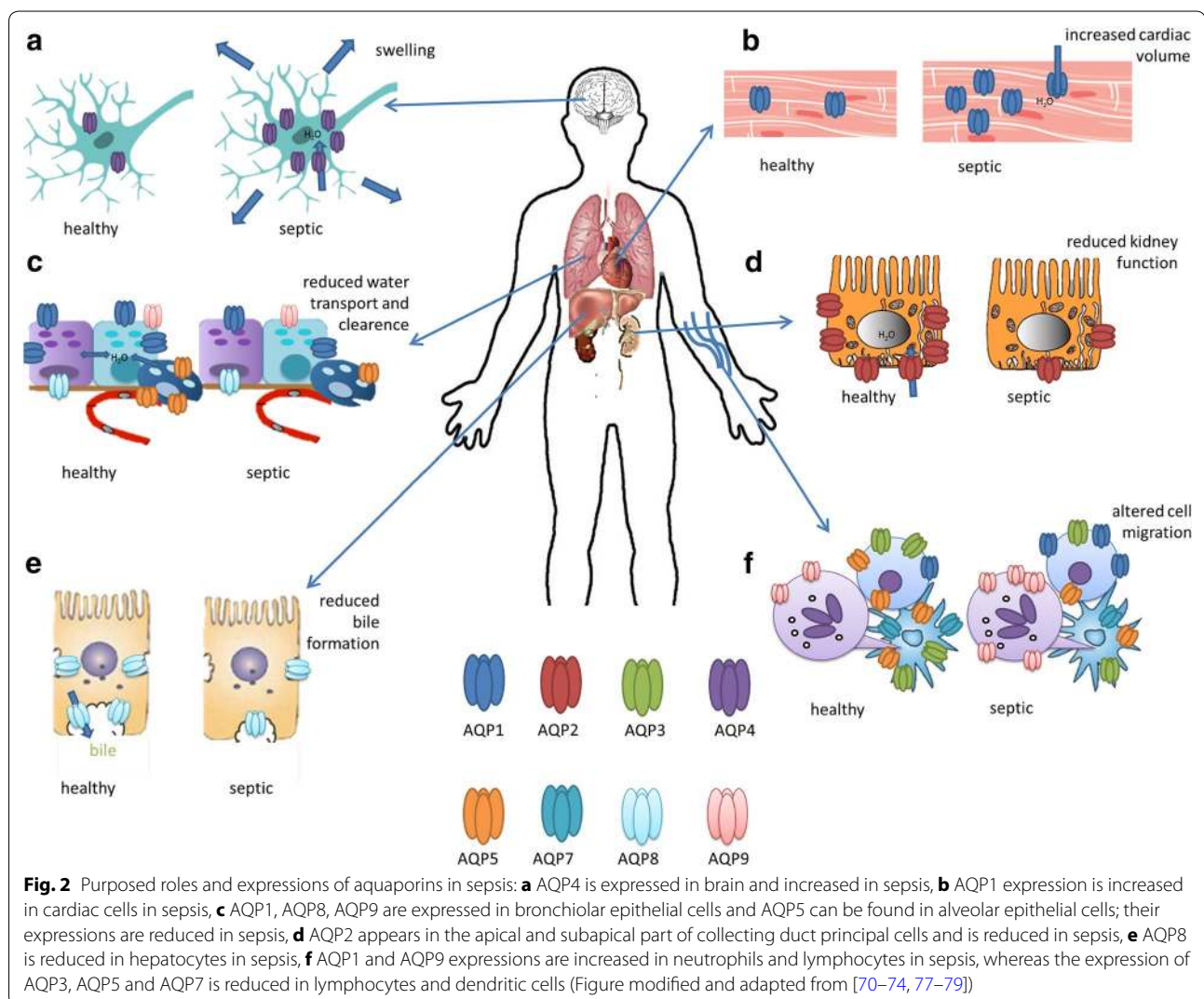
Role of aquaporins in the inflammasome

The inflammasome is an important key modulator of the immune response and affects the immune response by the release of proinflammatory cytokines. It can be found in macrophages and neutrophil granulocytes and can recognize pathogens like bacteria. The inflammasome inter alia consists of NLR family pyrin domain containing 3 (NLRP3) which is up-regulated in sepsis [26]. Activation of NLRP3 inflammasome causes interleukin 1 beta (IL-1 β) release. The IL-1 β release depends on the pH of the cell and its regulation is caused by water influx mediated

by aquaporins. AQP-mediated water movement in macrophages therefore appears as the common element unifying the variety of NLRP3 inflammasome activators [27].

Aquaporins in sepsis induced brain inflammation

One devastating complication of sepsis is septic encephalopathy (SE) [28]. In this context, aquaporins might play an important role, as SE is associated with vasogenic brain edema [29, 30]. The inflammation of the brain occurring in SE is mediated by neutrophil infiltration and causes *Aqp4* upregulation which aggravates brain edema [31, 32] (Fig. 2a). Upregulation of *Aqp4* in brain after LPS exposure can be attenuated by dexamethasone and this mechanism is mainly regulated by tumor necrosis factor alpha (TNF- α) [33]. However the use of corticosteroids like dexamethasone in sepsis is still discussed and its usage is only recommended under certain conditions [34].



In addition, AQP4 expression is upregulated in astrocytes during sepsis induced delirium (SID) and exosomes carrying AQP4 proteins from astrocytes to the peripheral blood may be utilized as biomarker for SID [35].

Aquaporins in kidney injury

Another common complication in sepsis is acute kidney injury (AKI), former called acute renal failure (ARF), which is frequently associated with polyuria and urine concentration defects and it increases the mortality rate in sepsis [36]. A cecal ligation and puncture (CLP) mouse model for sepsis showed that *Aqp2* expression is downregulated through NF- κ B pathway and may therefore cause acute renal failure during sepsis [37] (Fig. 2d). Pretreatment of rats with continuous erythropoietin receptor activator (CERA) preserves *Aqp2* expression in rat kidneys and protects against sepsis induced AKI [38].

The downregulation of *Aqp2* in sepsis models is confirmed by animal models using LPS induced endotoxemia after short time exposure (6 h) [39–42], whereas after long time exposure (18 h) *Aqp2* expression is increased in kidney [43]. Another study shows that *Aqp2* is downregulated after LPS administration in an LPS sepsis model in rats [44] and that pretreatment but not post-treatment with propofol prevents *Aqp2* downregulation and protects renal function during endotoxemia and that this effect may be mediated by regulation of Intercellular Adhesion Molecule 1 (ICAM-1), TNF- α and mediators of apoptosis [44]. Another possibility for *Aqp2* preservation after LPS exposure is treatment with α -lipoic acid [45].

Aquaporins in liver dysfunctions during sepsis

Liver has numerous functions in sepsis and is itself a target for sepsis induced injury [46]. For example septic shock and its toxins can cause hypoxic hepatitis, cholestasis due to altered bile metabolism or hepatocellular and acute liver injury [46]. In cholestasis AQP8 might play a role as it is downregulated after LPS stimulation in hepatocytes via TNF- α [11, 47]. The reduced AQP8 expression in turn causes reduced water permeability of hepatocytes, which can result in reduced bile formation and aggravates cholestasis [48, 49] (Fig. 2e). Beside, AQP8 can modulate hepatocellular mitochondria function by modifying water transport [50]. A loss of mitochondria function in turn can cause kidney injury due to loss of cellular energy [51]. In an endotoxemia rat model hepatic mitochondrial *Aqp8* expression is reduced [52]. Regulation of *Aqp8* in endotoxemia and septic models by substances like tetramethylpyrazine or ethyl pyruvate could stabilize the mitochondria membrane potential, protect hepatocellular mitochondria from damage and might therefore be a therapeutic option in sepsis [51, 53].

Aquaporins in cardiac dysfunction

40–50% of patients with prolonged septic shock develop cardiac dysfunction [54] and newer studies indicate that cardiac dysfunction can occur in all stages of sepsis [55]. The underlying molecular mechanisms are not fully understood yet, but a notable cause is mitochondrial dysfunction which contributes to cardiac dysfunction by causing myocardial energy depletion [56]. Here AQP1 might be important because *Aqp1* knockout causes cardiac hypertrophy in mice [57] (Fig. 2b). Another animal study tested the hypothesis if *Aqp1* may play a role in cardiac dysfunction during sepsis. They found that *Aqp1* expression is increased after LPS exposure in cardiac tissue and that this might influence cardiac function [58].

Aquaporins in acute lung injury

Another common complication in sepsis is acute lung injury that can cause acute respiratory distress syndrome (ARDS), which is associated with increased risk of in-hospital mortality [59]. In lung mainly the aquaporins AQP1 and 5, 8 and to a lower extent AQP9 are expressed [60]. Here, *Aqp1* is expressed in all vascular endothelial cells, *Aqp5* in the alveolar type I cells and *Aqp8* and *Aqp9* can be found in the bronchial epithelial cells in lung [61] (Fig. 2c). In 2016 in a small group of septic patients suffering from diffuse alveolar damage is was demonstrated

Table 1 Overview of AQP regulation during inflammation (\uparrow upregulation, \downarrow downregulation, ? unknown regulation, = unaffected)

Aquaporin	Tissue	Regulation during inflammation	References
AQP1	Immune cells	\uparrow In leukocytes and cell lines (THP-1)	[8, 9]
	Heart	\uparrow In cardiac cells	[58]
	Lung	\downarrow In lung tissue after LPS	[5, 6]
AQP2	Kidney	\downarrow In renal tissue after LPS	[37]
AQP3	Immune cells	\downarrow In leukocytes of septic patients	[8]
AQP4	Brain	\uparrow In brain and anterior pituitary gland	[31, 75]
AQP5	Lung	\downarrow In lung tissue after LPS	[65]
	Immune cells	\downarrow In THP-1 cells after LPS	[9]
AQP7	Immune cells	? Mouse resident peritoneal macrophages	[76]
AQP8	Liver	\downarrow In hepatic cells	[11]
	Lung	= In bronchial epithelial cells	[61]
AQP9	Immune cells	\uparrow In neutrophils of SIRS patients	[12]
	Immune cells	? Mouse resident peritoneal macrophages	[76]
	Lung	= In bronchial epithelial cells	[61]

that they have increased expression of AQP3 and AQP5 in the alveolar septum compared to healthy controls [62]. Recently it was demonstrated that *Aqp5* expression is decreased after sepsis induction with cecal ligation puncture (CLP) in the lung of rats [63, 64]. This effect can be attenuated by emodin [65] and is regulated by the microRNAs miR-96 and miR-330 [66]. In line with this *Aqp1* expression is decreased after LPS exposure in rat lungs [6, 67]. As a therapeutic option it was demonstrated that hydrogen rich saline and parenteral vitamin C can be protective in sepsis related lung injury and that it can attenuate the LPS induced reduction of *Aqp1* and *Aqp5* expression [5, 68]. In addition, *Aqp1* and *Aqp5* expression in lung is reduced in lung after an inflammatory pancreatitis models, whereas *Aqp8* and *Aqp9* expression remains unaffected [61]. Here the traditional Chinese prescription Dai-Huang-Fu-Zi-Tang can upregulate *Aqp1* and 5 and attenuate inflammation [61].

Conclusion

The regulatory mechanisms of aquaporins by LPS after endotoxemia and in sepsis seem to be tissue and aquaporin specific, as it can be seen in Table 1 and Fig. 2. As an example and it was demonstrated that AQP8 is downregulated in hepatic cells after LPS administration, though TNF- α pathway [11], while AQP9 expression remains unaffected [33, 69].

In summary, AQPs protein expressions seem to alter differential pathological mechanisms in sepsis and might be key proteins in inflammation. As a limitation of this review it has to be mentioned that several results were concluded from animal studies and that they potentially might to be fully adopted to human physiology. Elucidating the differential regulatory mechanisms of AQP expression in human studies might be helpful for developing novel sepsis therapeutics.

Abbreviations

AQP: aquaporin; CLP: cecal ligation and puncture; KO: knockout; LPS: lipopolysaccharide; SE: septic encephalopathy; TNF: tumor necrosis factor; WT: wildtype.

Authors' contributions

KR analyzed and interpreted the current literature and designed and wrote the manuscript. MA designed the workflow and discussed the topics of the manuscript. Both authors read and approved the final manuscript.

Acknowledgements

We acknowledge support by the DFG Open Access Publication Funds of the Ruhr-Universität Bochum.

Competing interests

The authors declare that they have no competing interests.

Availability of data and materials

Not applicable.

Consent for publication

All authors agree with the publication in Cell & Bioscience.

Ethics approval and consent to participate

Not applicable.

Funding

We acknowledge support by the DFG Open Access Publication Funds of the Ruhr-Universität Bochum.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Received: 7 August 2017 Accepted: 2 February 2018

Published online: 09 February 2018

References

1. Mayr FB, Yende S, Angus DC. Epidemiology of severe sepsis. *Virulence*. 2014;5(1):4–11.
2. Schorr CA, Zanotti S, Dellinger RP. Severe sepsis and septic shock: management and performance improvement. *Virulence*. 2014;5(1):190–9.
3. Adamzik M, Frey UH, Mohlenkamp S, Scherag A, Waydhas C, Marggraf G, Dammann M, Steinmann J, Siffert W, Peters J. Aquaporin 5 gene promoter—1364A/C polymorphism associated with 30-day survival in severe sepsis. *Anesthesiology*. 2011;114(1528–1175; 0003–3022; 4):912–17.
4. Liu L, Xie C. Effects of downregulation of aquaporin1 by peptidoglycan and lipopolysaccharide via MAPK pathways in MeT-5A cells. *Lung*. 2011;189(1432–1750; 0341–2040; 4):331–40.
5. Tao B, Liu L, Wang N, Wang W, Jiang J, Zhang J. Effects of hydrogen-rich saline on aquaporin 1, 5 in septic rat lungs. *J Surg Res*. 2016;202(2):291–8.
6. Liu LD, Wu XY, Tao BD, Wang N, Zhang J. Protective effect and mechanism of hydrogen treatment on lung epithelial barrier dysfunction in rats with sepsis. *Genet Mol Res*. 2016;15(1). <https://doi.org/10.4238/gmr.15016050>.
7. Verkman AS, Anderson MO, Papadopoulos MC. Aquaporins: important but elusive drug targets. *Nat Rev Drug Discov*. 2014;13:259–77.
8. Vassiliou AG, Maniatis NA, Orfanos SE, Mastora Z, Jahaj E, Pappourantas T, Armaganidis A, Roussos C, Aidinis V, Kotanidou A. Induced expression and functional effects of aquaporin-1 in human leukocytes in sepsis. *Crit Care*. 2013;17(5):R199.
9. Rump K, Brendt P, Frey UH, Schafer ST, Siffert W, Peters J, Adamzik M. Aquaporin 1 and 5 expression evoked by the ss-2 adrenoreceptor agonist terbutaline and LPS in mice and in the human monocytic cell line THP-1 is differentially regulated. *Shock*. 2013;40:430–6.
10. Molinas A, Mirazimi A, Holm A, Loitto VM, Magnusson K, Vikström E. Protective role of host aquaporin 6 against Hazara virus, a model for Crimean–Congo hemorrhagic fever virus infection. *FEMS Microbiol Lett*. 2016;363(8). <https://doi.org/10.1093/femsle/fnw058>.
11. Lehmann GL, Carreras FI, Soria LR, Gradilone SA, Marinelli RA. LPS induces the TNF-alpha-mediated downregulation of rat liver aquaporin-8: role in sepsis-associated cholestasis. *Am J Physiol Gastrointest Liver Physiol*. 2008;294(2):567.
12. Matsushima A, Ogura H, Koh T, Shimazu T, Sugimoto H. Enhanced expression of aquaporin 9 in activated polymorphonuclear leukocytes in patients with systemic inflammatory response syndrome. *Shock*. 2014;42(4):322–6.
13. Holm A, Karlsson T, Vikström E. *Pseudomonas aeruginosa* lasI/rhlI quorum sensing genes promote phagocytosis and aquaporin 9 redistribution to the leading and trailing regions in macrophages. *Front Microbiol*. 2015;6:915.
14. Huang YH, Zhou XY, Wang HM, Xu H, Chen J, Lv NH. Aquaporin 5 promotes the proliferation and migration of human gastric carcinoma cells. *Tumour Biol*. 2013;34(3):1743–51.
15. Karlsson T, Glogauer M, Ellen RP, Loitto VM, Magnusson KE, Magalhaes MA. Aquaporin 9 phosphorylation mediates membrane localization and neutrophil polarization. *J Leukoc Biol*. 2011;90(5):963–73.

16. Jung HJ, Park JY, Jeon HS, Kwon TH. Aquaporin-5: a marker protein for proliferation and migration of human breast cancer cells. *PLoS ONE*. 2011;6(12):e28492.
17. Papadopoulos MC, Saadoun S, Verkman AS. Aquaporins and cell migration. *Pflugers Arch*. 2008;456(0031–6768; 0031–6768; 4):693–700.
18. Moon C, Rousseau R, Soria JC, Hoque MO, Lee J, Jang SJ, Trink B, Sidransky D, Mao L. Aquaporin expression in human lymphocytes and dendritic cells. *Am J Hematol*. 2004;75(0361–8609; 0361–8609; 3):128–33.
19. Hara-Chikuma M, Sugiyama Y, Kabashima K, Sohara E, Uchida S, Sasaki S, Inoue S, Miyachi Y. Involvement of aquaporin-7 in the cutaneous primary immune response through modulation of antigen uptake and migration in dendritic cells. *FASEB J*. 2012;26(1):211–8.
20. Adamzik M, Frey UH, Bitzer K, Jakob H, Baba HA, Schmieder RE, Schneider MP, Heusch G, Peters J, Siffert W. A novel-1364A/C aquaporin 5 gene promoter polymorphism influences the responses to salt loading of the renin-angiotensin-aldosterone system and of blood pressure in young healthy men. *Basic Res Cardiol*. 2008;103(1435–1803; 0300–8428; 6):598–610.
21. Rump K, Unterberg M, Bergmann L, Bankfalvi A, Menon A, Schäfer S, Scherag A, Bazzi Z, Siffert W, Peters J, Adamzik M. AQP5-1364A/C polymorphism and the AQP5 expression influence sepsis survival and immune cell migration: a prospective laboratory and patient study. *J Transl Med*. 2016;14(1):321.
22. Zhu N, Feng X, He C, Gao H, Yang L, Ma Q, Guo L, Qiao Y, Yang H, Ma T. Defective macrophage function in aquaporin-3 deficiency. *FASEB J*. 2011;25(12):4233–9.
23. Hara-Chikuma M, Chikuma S, Sugiyama Y, Kabashima K, Verkman AS, Inoue S, Miyachi Y. Chemokine-dependent T cell migration requires aquaporin-3-mediated hydrogen peroxide uptake. *J Exp Med*. 2012;209(10):1743–52.
24. Chi Y, Fan Y, He L, Liu W, Wen X, Zhou S, Wang X, Zhang C, Kong H, Sonoda L, Tripathi P, Li CJ, Yu MS, Su C, Hu G. Novel role of aquaporin-4 in CD4⁺ CD25⁺ T regulatory cell development and severity of Parkinson's disease. *Aging Cell*. 2011;10(3):368–82.
25. Moniaga CS, Watanabe S, Honda T, Nielsen S, Hara-Chikuma M. Aquaporin-9-expressing neurophilis are required for the establishment of contact hypersensitivity. *Sci Rep*. 2015;5:15319.
26. Esquerdo KF, Sharma NK, Brunialti MKC, Baggio-Zappia GL, Assunção M, Azevedo LCP, Bafi AT, Salomao R. Inflammasome gene profile is modulated in septic patients, with a greater magnitude in non-survivors. *Clin Exp Immunol*. 2017;189(2):232–40.
27. Rabolli V, Wallemme L, Lo Re S, Uwambayinema F, Palmi-Pallag M, Thomassen L, Tyteca D, Octave JN, Marbaix E, Lison D, Devuyt O, Huaux F. Critical role of aquaporins in IL-1beta-mediated inflammation. *J Biol Chem*. 2014;289:13937–47.
28. Tauber SC, Eiffert H, Brück W, Nau R. Septic encephalopathy and septic encephalitis. *Expert Rev Anti Infect Ther*. 2017;15(2):121–32.
29. Tong D, Zhou Y, Wang G, Chen X, Yang T. Early prediction and outcome of septic encephalopathy in acute stroke patients with nosocomial coma. *J Clin Med Res*. 2015;7(7):534–9.
30. Davies DC. Blood-brain barrier breakdown in septic encephalopathy and brain tumours. *J Anat*. 2002;200(6):639–46.
31. Alexander JJ, Jacob A, Cunningham P, Hensley L, Quigg RJ. TNF is a key mediator of septic encephalopathy acting through its receptor, TNF receptor-1. *Neurochem Int*. 2008;52(3):447–56.
32. Rama Rao KV, Jayakumar AR, Norenberg MD. Brain edema in acute liver failure: mechanisms and concepts. *Metab Brain Dis*. 2014;29(4):927–36.
33. Du Y, Meng Y, Lv X, Guo L, Wang X, Su Z, Li L, Li N, Zhao S, Zhao L, Zhao X. Dexamethasone attenuates LPS-induced changes in expression of urea transporter and aquaporin proteins, ameliorating brain endotoxemia in mice. *Int J Clin Exp Pathol*. 2014;7(12):8443–52.
34. Rhodes A, Evans LE, Alhazzani W, Levy MM, Antonelli M, Ferrer R, Kumar A, Sevransky JE, Sprung CL, Nunnally ME, Rochweg B, Rubenfeld GD, Angus DC, Annane D, Beale RJ, Bellinhan GJ, Bernard GR, Chiche J, Coopersmith C, De Backer DP, French CJ, Fujishima S, Gerlach H, Hidalgo JL, Hollenberg SM, Jones AE, Karnad DR, Kleinpell RM, Koh Y, Lisboa TC, Machado FR, Marini JJ, Marshall JC, Mazuski JE, McIntyre LA, McLean AS, Mehta S, Moreno RP, Myburgh J, Navalesi P, Nishida O, Osborn TM, Perner A, Plunkett CM, Ranieri M, Schorr CA, Seckel MA, Seymour CW, Shieh L, Shukri KA, Simpson SQ, Singer M, Thompson BT, Townsend SR, Van der Poll T, Vincent J, Wiersinga WJ, Zimmerman JL, Dellinger RP. Surviving sepsis campaign: international guidelines for management of sepsis and septic shock: 2016. *Crit Care Med*. 2017;45(3):486–552.
35. Sfera A, Price AI, Gradini R, Cummings M, Osorio C. Proteomic and epigenomic markers of sepsis-induced delirium (SID). *Front Mol Biosci*. 2015;2:59.
36. Bellomo R, Kellum JA, Ronco C, Wald R, Martensson J, Maiden M, Bagshaw SM, Glassford NJ, Lankadeva Y, Vaara ST, Schneider A. Acute kidney injury in sepsis. *Intensive Care Med*. 2017;43(6):816–28.
37. Höcherl K, Schmidt C, Kurt B, Bucher M. Inhibition of NF-kappaB ameliorates sepsis-induced downregulation of aquaporin-2/V2 receptor expression and acute renal failure in vivo. *Am J Physiol Renal Physiol*. 2010;298(1):196.
38. Rodrigues CE, Sanches TR, Volpini RA, Shimizu MHM, Kuriki PS, Camara NOS, Seguro AC, Andrade L. Effects of continuous erythropoietin receptor activator in sepsis-induced acute kidney injury and multi-organ dysfunction. *PLoS ONE*. 2012;7(1):e29893.
39. Olesen ETB, de Seigneux S, Wang G, Lütken SC, Frøkiaer J, Kwon T, Nielsen S. Rapid and segmental specific dysregulation of AQP2, S256-pAQP2 and renal sodium transporters in rats with LPS-induced endotoxaemia. *Nephrol Dial Transplant*. 2009;24(8):2338–49.
40. Wang W, Li C, Summer SN, Falk S, Wang W, Ljubanovic D, Schrier RW. Role of AQP1 in endotoxemia-induced acute kidney injury. *Am J Physiol Renal Physiol*. 2008;294(6):1473.
41. Versteilen AMG, Heemskerk AEJ, Groeneveld ABJ, van Wijhe M, van Lambalgen AA, Tangelder G. Mechanisms of the urinary concentration defect and effect of desmopressin during endotoxemia in rats. *Shock*. 2008;29(2):217–22.
42. Grinevich V, Knepper MA, Verbalis J, Reyes I, Aguilera G. Acute endotoxemia in rats induces down-regulation of V2 vasopressin receptors and aquaporin-2 content in the kidney medulla. *Kidney Int*. 2004;65(1):54–62.
43. Chagnon F, Vaidya VS, Plante GE, Bonventre JV, Bernard A, Guindi C, Lesur O. Modulation of aquaporin-2/vasopressin2 receptor kidney expression and tubular injury after endotoxin (lipopolysaccharide) challenge. *Crit Care Med*. 2008;36(11):3054–61.
44. Cui W, Tian A, Bai T. Protective effects of propofol on endotoxemia-induced acute kidney injury in rats. *Clin Exp Pharmacol Physiol*. 2011;38(11):747–54.
45. Suh SH, Lee KE, Kim IJ, Kim O, Kim CS, Choi JS, Choi H, Bae EH, Ma SK, Lee JU, Kim SW. Alpha-lipoic acid attenuates lipopolysaccharide-induced kidney injury. *Clin Exp Nephrol*. 2015;19(1):82–91.
46. Strnad P, Tacke F, Koch A, Trautwein C. Liver—guardian, modifier and target of sepsis. *Nat Rev Gastroenterol Hepatol*. 2017;14(1):55–66.
47. Bhogal HK, Sanyal AJ. The molecular pathogenesis of cholestasis in sepsis. *Front Biosci (Elite Ed)*. 2013;5:87–96.
48. Marinelli RA, Lehmann GL, Soria LR, Marchisio MJ. Hepatocyte aquaporins in bile formation and cholestasis. *Front Biosci (Landmark Ed)*. 2011;16:2642–52.
49. Lehmann G, Larocca M, Soria L, Marinelli R. Aquaporins: their role in cholestatic liver disease. *World J Gastroenterol*. 2008;14(46):7059–67.
50. Xu X, Shi Z, Hu J, Yuan B, Huang H, Fang H, Yin X, Nie N, Sheng X. Identification of differentially expressed genes associated with burn sepsis using microarray. *Int J Mol Med*. 2015;36(6):1623–9.
51. Wang J, Zhang L, Tao X, Wei L, Liu B, Huang L, Chen Y. Tetramethylpyrazine upregulates the aquaporin 8 expression of hepatocellular mitochondria in septic rats. *J Surg Res*. 2013;185(1):286–93.
52. Soria LR, Marrone J, Molinas SM, Lehmann GL, Calamita G, Marinelli RA. Lipopolysaccharide impairs hepatocyte ureagenesis from ammonia: involvement of mitochondrial aquaporin-8. *FEBS Lett*. 2014;588(9):1686–91.
53. Jiang Z, Li X, Lin Z, Chen J, Guan X, Chen M. Ethyl pyruvate reduces hepatic mitochondrial swelling and dysfunction in a rat model of sepsis. *Int J Clin Exp Pathol*. 2015;8(7):7774–85.
54. Rudiger A, Singer M. Mechanisms of sepsis-induced cardiac dysfunction. *Crit Care Med*. 2007;35(6):1599–608.
55. Flynn A, Chokalingam Mani B, Mather PJ. Sepsis-induced cardiomyopathy: a review of pathophysiologic mechanisms. *Heart Fail Rev*. 2010;15(6):605–11.
56. Cimolai MC, Alvarez S, Bode C, Bugger H. Mitochondrial mechanisms in septic cardiomyopathy. *Int J Mol Sci*. 2015;16(8):17763–78.
57. Montiel V, Leon Gomez E, Bouzin C, Esfahani H, Romero Perez M, Loby-sheva I, Devuyt O, Dessy C, Balligand JL. Genetic deletion of aquaporin-1

- results in microcardia and low blood pressure in mouse with intact nitric oxide-dependent relaxation, but enhanced prostanoids-dependent relaxation. *Pflugers Arch*. 2014;466(2):237–51.
58. Madonna R, Jiang J, Geng YJ. Attenuated expression of gelsolin in association with induction of Aquaporin-1 and nitric oxide synthase in dysfunctional hearts of aging mice exposed to endotoxin. *Int J Immunopathol Pharmacol*. 2012;25(0394–6320; 0394–6320; 4):911–22.
 59. Kim W, Hong S. Sepsis and acute respiratory distress syndrome: recent update. *Tuberc Respir Dis (Seoul)*. 2016;79(2):53–7.
 60. Hong-Min F, Chun-Rong H, Rui Z, Li-Na S, Ya-Jun W, Li L. CGRP 8-37 enhances lipopolysaccharide-induced acute lung injury and regulating aquaporin 1 and 5 expressions in rats. *J Physiol Biochem*. 2016;73(3):381–6.
 61. Kang X, Lu X, Zhan L, Liang Z, Guo W, Ma Q, Wang Y, Song J, Feng J, Wang C, Bai L, Song Y, Liu G. Dai-Huang-Fu-Zi-Tang alleviates pulmonary and intestinal injury with severe acute pancreatitis via regulating aquaporins in rats. *BMC Complement Altern Med*. 2017;17(1):288.
 62. Pires-Neto RC, Del Carlo Bernardi F, Alves de Araujo P, Mauad T, Dolhnikoff M. The expression of water and ion channels in diffuse alveolar damage is not dependent on DAD etiology. *PLoS ONE*. 2016;11(11):e0166184.
 63. Chinnaiyan AM, Huber-Lang M, Kumar-Sinha C, Barrette TR, Shankar-Sinha S, Sarma VJ, Padgaonkar VA, Ward PA. Molecular signatures of sepsis: multiorgan gene expression profiles of systemic inflammation. *Am J Pathol*. 2001;159(4):1199–209.
 64. Bromberg Z, Raj N, Goloubinoff P, Deutschman CS, Weiss YG. Enhanced expression of 70-kilodalton heat shock protein limits cell division in a sepsis-induced model of acute respiratory distress syndrome. *Crit Care Med*. 2008;36(1):246–55.
 65. Sun Y, Sun L, Liu S, Song J, Cheng J, Liu J. Effect of emodin on aquaporin 5 expression in rats with sepsis-induced acute lung injury. *J Tradit Chin Med*. 2015;35(6):679–84.
 66. Zhang Y, Chen M, Zhang Y, Peng P, Li J, Xin X. miR-96 and miR-330 overexpressed and targeted AQP5 in lipopolysaccharide-induced rat lung damage of disseminated intravascular coagulation. *Blood Coagul Fibrinolysis*. 2014;25(7):731–7.
 67. Ma T, Liu Z. Functions of aquaporin 1 and α -epithelial Na⁺ channel in rat acute lung injury induced by acute ischemic kidney injury. *Int Urol Nephrol*. 2013;45(4):1187–96.
 68. Fisher BJ, Kraskauskas D, Martin EJ, Farkas D, Wegelin JA, Brophy D, Ward KR, Voelkel NF, Fowler AA, Natarajan R. Mechanisms of attenuation of abdominal sepsis induced acute lung injury by ascorbic acid. *Am J Physiol Lung Cell Mol Physiol*. 2012;303(1):20.
 69. Lehmann GL, Marinelli RA. Peritoneal sepsis downregulates liver expression of aquaporin-8: a water channel involved in bile secretion. *Liver Int*. 2009;29(2):317–8.
 70. Wang H, Thorling CA, Liang X, Bridle KR, Grice JE, Zhu Y, Crawford DHG, Xu ZP, Liu X, Roberts MS. Diagnostic imaging and therapeutic application of nanoparticles targeting the liver. *J Mater Chem B*. 2015;3(6):939–58.
 71. Eymael J, Smeets B. Origin and fate of the regenerating cells of the kidney. *Eur J Pharmacol*. 2016;790:62–73.
 72. Nielsen S, King LS, Christensen BM, Agre P. Aquaporins in complex tissues. II. Subcellular distribution in respiratory and glandular tissues of rat. *Am J Physiol*. 1997;273(5 Pt 1):1549.
 73. Nielsen S, Frøkiaer J, Marples D, Kwon T, Agre P, Knepper MA. Aquaporins in the kidney: from molecules to medicine. *Physiol Rev*. 2002;82(1):205–44.
 74. Kovach TK, Dighe AS, Lobo PI, Cui Q. Interactions between MSCs and immune cells: implications for bone healing. *J Immunol Res*. 2015;2015:752510.
 75. Kuwahara-Otani S, Maeda S, Tanaka K, Hayakawa T, Seki M. Systemic administration of lipopolysaccharide increases the expression of aquaporin-4 in the rat anterior pituitary gland. *J Vet Med Sci*. 2013;75(8):1081–4.
 76. Zhu N, Feng X, He C, Gao H, Yang L, Ma Q, Guo L, Qiao Y, Yang H, Ma T. Defective macrophage function in aquaporin-3 deficiency. *FASEB J*. 2011;25(1530–6860; 0892–6638; 12):4233–39.
 77. Anonymous: animal primary tissues—OpenStax CNX. <http://cnx.org/contents/f8b7e159-1112-46eabe19-5f492747a7b5@4/Animal-Primary-Tissues>.
 78. Anonymous: Brain cells|The Brain Tumour Charity. <https://www.thebraintumourcharity.org/understanding-brain-tumours/symptoms-and-information/braincells/>.
 79. Fliesler N. Reversing lung disease in mice by coaxing production of healthy cells. <https://vector.childrenshospital.org/2014/01/reversing-lung-disease-in-mice-by-coaxing-production-of-healthy-cells/>.

Submit your next manuscript to BioMed Central and we will help you at every step:

- We accept pre-submission inquiries
- Our selector tool helps you to find the most relevant journal
- We provide round the clock customer support
- Convenient online submission
- Thorough peer review
- Inclusion in PubMed and all major indexing services
- Maximum visibility for your research

Submit your manuscript at
www.biomedcentral.com/submit

