



Protein expression levels in the medullary visceral zone of rats with subarachnoid hemorrhage

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ABSTRACT. We investigated protein expression in the medullary visceral zone (MVZ) of rats with multiple-organ dysfunction syndrome (MODS) caused by subarachnoid hemorrhage (SAH) to discuss the possible regulatory mechanism of the MVZ in the course of SAH-induced MODS. A SAH-induced MODS model was established in rats by injecting arterial blood into the Willis' circle. Protein expression in the MVZ was analyzed by immunohistochemistry assay. Protein expression in the MVZ peaked 24-36 h after SAH, and was significantly higher than in the control and sham operation groups. Organs at each time point exhibited inflammatory injuries to varying degrees after SAH, which reached a maximum at 24-36 h. Incidences of systemic inflammatory response syndrome and MODS were 100 and 71.67%, respectively, after SAH. There is a consistency between MVZ protein expression and inflammatory changes in each organ after SAH. This

prompts the suggestion that the MVZ may be one of the direct regulative centers in SAH-induced MODS, and may be involved in the functional regulation of the surrounding organs after SAH.

Key words: Subarachnoid hemorrhage; Medullary visceral zone; Rat model; Protein expression; Multiple-organ dysfunction syndrome

INTRODUCTION

The mechanisms of systemic inflammatory response syndrome (SIRS) and multiple-organ dysfunction syndrome (MODS) caused by acute cerebrovascular disease (ACVD) have not been fully elucidated either in China or internationally (Whiteley et al., 2011; Koton et al., 2012). In the past, research has focused on the hypothalamus. In particular, a series of studies have been carried out on changes of the neuroendocrine function of the hypothalamic-pituitary-target gland axis in the course of ACVD-induced MODS (Arauz et al., 2010; Yavuz et al., 2012). Recent studies have found that a structural function region called the medullary visceral zone (MVZ), located in the middle and tail of the medulla oblongata, participates in the regulation of visceral activities under various stress states. However, whether the MVZ is also involved in the process of MODS caused by ACVD has rarely been reported (Rösler et al., 2010; Abdo et al., 2012).

According to the diagnostic criteria of SIRS and MODS for laboratory animals, we successfully established an animal model for MODS induced by subarachnoid hemorrhage (SAH) in the early stage. We used the model to explore the central regulative role of the MVZ in the functions of the surrounding organs after SAH by observing protein expression in the MVZ of the model rats. Our objective was to provide a new theoretical and experimental basis for the clinical prevention of ACVD-induced MODS.

MATERIAL AND METHODS

Animal grouping

Eighty healthy adult male Wistar rats (clean-grade) weighing 200-220 g were provided by the SLRC Laboratory Animal Co., Ltd., Shanghai, China. The rats were divided into a normal control group, a sham operation group, and six experimental subgroups of 10 rats each for observations at 6, 12, 24, 36, 48, and 72 h following SAH using a random number table.

Modeling

The SAH-induced MODS model was established in the rats by injecting arterial blood into the Willis' circle (Ozel et al., 2009). Life signs of the rats at each time point were observed postoperatively, and blood routine, hepatic and renal function, and cardiac enzymes were detected. The tissues of the lung, liver, kidney, and small intestine were taken for pathological examination by hematoxylin and eosin staining. Protein expression in the MVZ was determined using the immunohistochemical ABC method after freezing sections of cerebral tissue. A set of medullary slices was randomly selected from each rat to calculate the sum of positive MVZ cells in the set of slices at a magnification of 20 (ocular lens) x 20 (objective lens).

Statistical analysis

All data are reported as means \pm standard error and were analyzed by SPSS 13.0. The data were subjected to the *t*-test and analysis of variance. $P < 0.05$ was considered to be statistically significant.

RESULTS

Vital signs and blood biochemical indices

The differences in breathing, heart rate, body temperature, peripheral white blood cell count, alanine aminotransferase, aspartate aminotransferase, blood urea nitrogen, creatinine, and creatine kinase were not statistically significant between the sham operation group and the normal control group ($P > 0.05$). The above indicators of the SAH group were all higher than those of the normal control group and the sham operation group ($P < 0.01$; Tables 1 and 2).

Table 1. Basic data for animals.

| Group | Respiratory rate (bpm) | Heart rate (bpm) | Body temperature ($^{\circ}$ C) | WBC ($\times 10^9$ /L) |
|----------------|------------------------|----------------------|----------------------------------|-------------------------|
| Normal | 59.42 \pm 2.16 | 146.52 \pm 26.17 | 35.92 \pm 0.64 | 7.06 \pm 0.28 |
| Sham operation | 60.18 \pm 2.32 | 149.29 \pm 24.91 | 36.04 \pm 0.59 | 7.13 \pm 0.29 |
| SAH | 107.81 \pm 9.16** | 294.65 \pm 39.15** | 39.48 \pm 0.81** | 18.46 \pm 2.61** |

Data are reported as means \pm SE. bpm = breaths (or beats) per minute; WBC = white blood cell count; SAH = subarachnoid hemorrhage. Compared with the normal group, # $P < 0.01$; compared with the sham operation group, * $P < 0.01$.

Table 2. Blood biochemical indices at each time point.

| Group | ALT (U/L) | AST (U/L) | BUN (μ M) | Cr (μ M) | CK (U/L) |
|----------------|--------------------------------|---------------------------------|------------------------------|-------------------------------|----------------------------------|
| Normal | 64.58 \pm 9.12 | 143.52 \pm 16.18 | 5.89 \pm 0.87 | 26.84 \pm 3.29 | 692.25 \pm 39.46 |
| Sham operation | 67.29 \pm 9.45 | 146.82 \pm 15.64 | 5.93 \pm 0.91 | 28.56 \pm 3.81 | 703.15 \pm 38.42 |
| SAH | | | | | |
| 6 h | 76.49 \pm 10.61 | 159.53 \pm 24.89 | 7.92 \pm 1.06 | 36.54 \pm 4.91** | 953.55 \pm 61.98 ^{ab} |
| 12 h | 159.29 \pm 26.14** | 194.27 \pm 29.18** | 15.57 \pm 2.64** | 49.82 \pm 5.66** | 1652.94 \pm 95.56** |
| 24 h | 216.18 \pm 29.49** | 299.54 \pm 39.15** | 25.26 \pm 3.27** | 69.92 \pm 6.57** | 1968.28 \pm 96.25** |
| 36 h | 204.56 \pm 22.17** | 261.59 \pm 29.68** | 19.67 \pm 2.51** | 60.68 \pm 5.91** | 1539.67 \pm 89.26** |
| 48 h | 142.38 \pm 16.28** | 205.92 \pm 18.16** | 14.65 \pm 2.09** | 52.94 \pm 4.68** | 1256.26 \pm 76.16** |
| 72 h | 97.46 \pm 12.57 ^a | 177.84 \pm 13.58 ^a | 8.99 \pm 1.67 ^a | 39.19 \pm 4.46 ^a | 991.58 \pm 71.65 ^a |

Data are reported as means \pm SE. ALT = alanine aminotransferase; AST = aspartate aminotransferase; BUN = blood urea nitrogen; Cr = creatinine; CK = creatine kinase; SAH = subarachnoid hemorrhage. Compared with the normal group, * $P < 0.01$, ^a $P < 0.05$; compared with the sham operation group, # $P < 0.01$, ^b $P < 0.05$.

Incidence rates of SIRS and MODS in the SAH group

The incidences of SIRS and MODS in the SAH group were 100 and 71.67%, respectively. There were seven deaths before the time points, accounting for 11.67% of the rats in the SAH group.

Pathological changes of main organs

The histological structures of each organ in the normal control group and the sham

operation group were basically normal. The lungs of the SAH group exhibited telangiectasia congestion, and exudation in alveolar space and perivascular area at 6 h. Alveolar space was obviously expanded, and fibrin effusion was observed in the bronchial and tracheal cavities at 12 h. Mild chronic lymphocytic bronchial pneumonia was found after 24 and 36 h (Figure 1A). The inflammatory change mentioned above was alleviated at 48 h. Edema and thickening appeared in the mucosa and submucosa of the small intestine at 24 h. Edema, thickening, congestion and mild inflammatory cell infiltration were observed between the muscular layer and the mucosal layer after 36 and 48 h (Figure 1B). Mucosal and submucosal congestion and edema occurred occasionally after 72 h. Cloudy swelling was found in a small number of hepatocytes after 8 h. Spotty necrosis of hepatocytes and hepatic sinusoid expansion were observed after 24 h (Figure 1C). A small amount of interstitial chronic inflammatory cell infiltration and acidophilic change of hepatocytes appeared after 36 h, and was alleviated after 48 h. Renal interstitial congestion began to occur and cloudy swelling of a few epithelial cells of the proximal renal tubule was found after 12 h. Glomerular atrophy and renal tubular dilatation of varying degrees appeared after 24 and 36 h (Figure 1D), which were alleviated after 48 h.

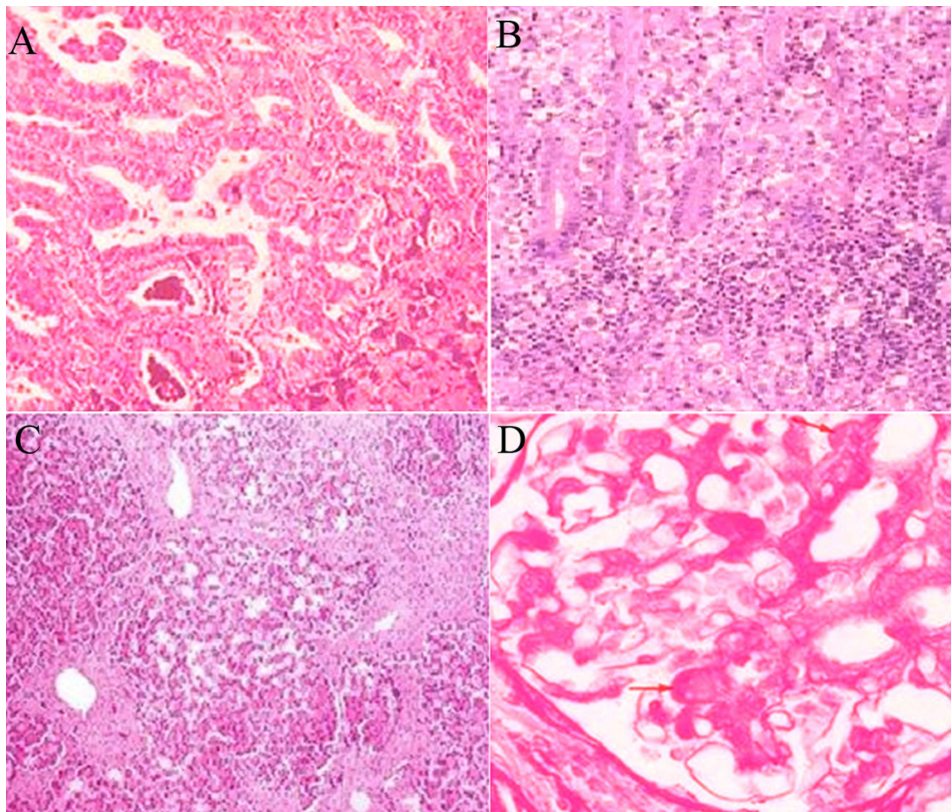


Figure 1. Pathological changes of main organs in the subarachnoid hemorrhage (SAH) group. **A.** Pathological changes of lung tissues 24 h after hemorrhage (SP 400X). **B.** Pathological changes of small intestine tissues 36 h after hemorrhage (SP 400X). **C.** Pathological changes of liver tissues 24 h after hemorrhage (SP 400X). **D.** Pathological changes of kidney tissues 36 h after hemorrhage (arrows; SP 400X).

MVZ protein expression levels

Both the normal control group and the sham operation group exhibited a small number of lightly stained positive MVZ cells (Figure 2A), between which the difference was not statistically significant ($P > 0.05$). The number of positive MVZ cells in the SAH group was significantly higher than in the normal control and sham operation groups, in which the majority were deeply stained (Figure 2B). The positive cells were relatively numerous in the nucleus tractus solitarii (NTS), the area postrema, and the dorsal vagal nucleus (DMV) in the dorsomedial region and the lateral reticular nucleus (LRN) in the ventrolateral medulla; they peaked at 24 h and gradually decreased after 6-48 h, but there was still positive expression at 72 h (Table 3).

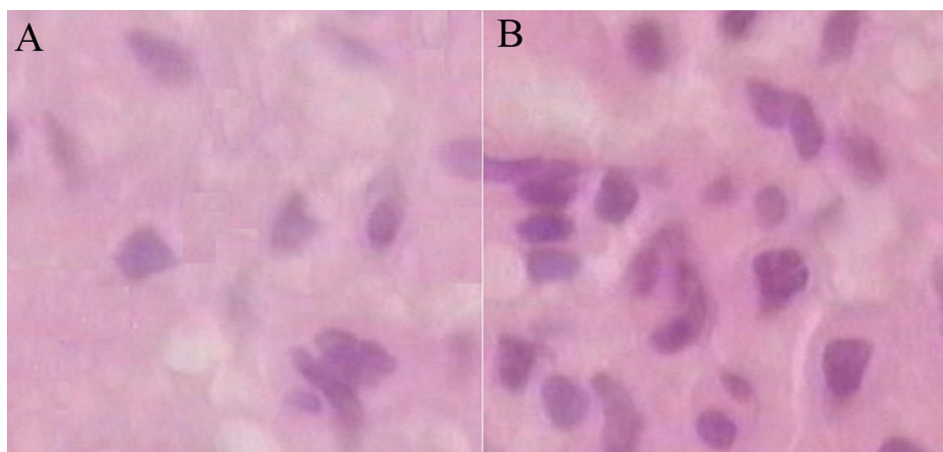


Figure 2. Medullary visceral zone (MVZ) protein expression levels. **A.** A small number of lightly stained positive MVZ cells in the ventrolateral medulla (VLM) of the sham operation group (SP 400X). **B.** Numerous darkly stained positive MVZ cells in the VLM of the subarachnoid hemorrhage group (SP 400X).

Table 3. Positive expressions of medullary visceral zone (MVZ) proteins in the ventrolateral medulla (VLM) at each time point.

| Group | N | Number of positively expressed MVZ proteins |
|----------------|----|---|
| Normal | 10 | 8.46 ± 2.07 |
| Sham operation | 10 | 9.38 ± 2.43 |
| SAH | | |
| 6 h | 10 | 34.51 ± 4.65** |
| 12 h | 10 | 59.16 ± 6.29** |
| 24 h | 10 | 98.32 ± 10.65** |
| 36 h | 10 | 88.49 ± 7.59** |
| 48 h | 10 | 67.56 ± 7.18** |
| 72 h | 10 | 42.65 ± 4.69** |

Data are reported as means ± SE. SAH = subarachnoid hemorrhage. Compared with the normal group, # $P < 0.01$; compared with the sham operation group, * $P < 0.01$.

DISCUSSION

Recent studies have found that there is a ribbon called the MVZ that runs from the dorsomedial to the ventrolateral region on the transverse section of the middle and tail of the

rat medulla oblongata. The MVZ, an important center for the regulation of stress response, is closely related to the transmission of somatic and visceral sensitivity and the regulation of visceral activities, exerting a strong influence on the surrounding organs during ACVD (Jiang et al., 2009; Yang et al., 2010a). Protein is expressed in the MVZ in the early stages of ACVD. As a cell transcription regulatory factor, the MVZ protein is involved in the transcription process of all enzymes effected by the signal transduction system. Many studies have confirmed that the MVZ protein, which fulfills a “tracer” role, can be used as a positioning and functional indicator of neuronal activity (Treger et al., 2008; He et al., 2013). Zhao et al. (2011) replicated MODS caused by cerebral hemorrhage in a rat model using different doses (0.5 and 1.0 U) of collagenase and found that the positive expression of MVZ protein increased after cerebral hemorrhage compared with the control group, and the number of positively expressing MVZ cells was significantly higher in the 1.0 U group than in the 0.5 U group. Thus, it can be seen that severe ACVD may cause changes in MVZ protein expression in the central nervous system, and to a certain extent the quantity of protein induced is related to the stimulation intensity suffered. Studies have shown that after rats were given 1% formaldehyde by intragastric administration, bulbar MVZ immunoreactive neurons were mainly located in the NTS ventrolateral region of the middle and tail and the reticular structure between the two, and the number of MVZ/tyrosine hydroxylase double-positive neurons accounted for more than 50% of the total number of tyrosine hydroxylase single-positive neurons, suggesting that more than half of the catecholaminergic neurons in the MVZ are involved in the stress response to noxious stimuli in the digestive tract (Gallas et al., 2011). In this study, the rats in the SAH group exhibited inflammatory damage in the small intestine 24 h after SAH, which reached a peak between 36 and 48 h, and was significantly alleviated after 72 h. This is consistent with the MVZ expression pattern after SAH, especially the intensive expression of MVZ protein in the DMV. This prompts the suggestion that the functions of NTS and DMV are affected after SAH, causing inflammatory damage of the intestinal mucosa and submucosa (Yang et al., 2010b).

Using the cardiovascular stress response model of hypertension and acute myocardial ischemia induced by intravenous injection of Pituitrin, some researchers found that MVZ protein expression was limited to the MVZ region, and was negative in the surrounding nuclei. Its expression was more intensive in the NTS and ventrolateral medulla, in which 50% of neurons were MVZ/tyrosine hydroxylase double-positive, indicating that the catecholaminergic neurons in MVZ are involved in the stress response to cardiovascular noxious stimuli (Rotoli et al., 2011; Daubert et al., 2012). In view of the limited experimental conditions, this study did not investigate the changes of blood pressure and electrocardiogram measurements in the rats after SAH, but the changes in heart rate and myocardial enzymes did reflect myocardial ischemia. In this study, the MVZ protein was markedly expressed 6 h after hemorrhage in the SAH group, and was mainly distributed in the NTS, LRN, and DMV, and to a lesser extent in the area postrema. The protein expression peaked after 24 h, but was still relatively intensive after 36 h and gradually subsided after 72 h. The positive expression of MVZ protein was consistent with the changes of heart rate and myocardial enzymes, prompting the suggestion that positive MVZ protein expression is associated with peripheral myocardial ischemia. Therefore, it is inferred that the MVZ participates in the regulation of heart function after SAH.

Electric damaged the NTS in the MVZ of rats can lead to neurogenic pulmonary edema. It is thought that the ascending fibers of the caudal NTS have a wide range of fiber links to catecholaminergic neurons in the brainstem reticular structure, so as to cause neurogenic pulmonary edema by regulating the release of catecholamines (Mukadam et al., 2005;

Schreiber et al., 2007). This study found that the breathing frequency at each time point was significantly faster in the SAH group compared with the control group, the difference being more pronounced at 24, 36, and 48 h. In addition, non-specific pulmonary inflammation also appeared 6 h after SAH, reached a peak between 24 and 36 h, and subsided after 48 h. At the same time, the positive expression of MVZ protein could also be found at the same regular pattern, and was most pronounced in the caudal NTS and LRN. In the SAH group, MVZ-positive expression in the sites mentioned above appeared at 6 h, peaked at 24 h, was still sustained at a relatively high level at 36 h, and began to subside gradually at 48 h, which prompts the suggestion that a series of cerebral pathophysiological changes after SAH can affect the NTS and LRN functions in MVZ, so that the regulation of respiratory function of the lung becomes unbalanced under normal circumstances, thereby causing neurogenic pulmonary edema.

The mechanism of ACVD-induced MODS is complicated. This study found that there is an obvious relationship over time between MVZ protein expression and pathological changes in myocardial enzymes and the surrounding vital organs following SAH. Therefore, there may be a necessary relationship between them, suggesting that the MVZ is one of the direct regulative centers in SAH-induced MODS, and is involved in the functional regulation of the surrounding organs following SAH.

Conflicts of interest

The authors declare no conflict of interest.

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