

Metabolic alterations in the rat cerebellum following acute middle cerebral artery occlusion, as determined by ¹H NMR spectroscopy

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Abstract. Supratentorial focal ischemia may reduce cerebral blood volume and cerebellar glucose metabolic rate contralateral to the region of ischemia. The present study investigated the effects of middle cerebral artery occlusion (MCAO) on cerebral metabolism in the ischemic cerebral hemisphere and the non-ischemic cerebellum in rats 1, 3, 9 and 24 h following ischemia using *ex vivo* proton nuclear magnetic resonance (¹H NMR) spectroscopy. The results demonstrated that focal ischemia induced increases in the levels of lactate and alanine, and a decrease in succinate, as early as 1 h following ischemia in the left cerebral hemisphere and the right cerebellum. A continuous increase in lactate levels and

decrease in creatine levels were detected in both cerebral areas 3 and 24 h post-MCAO. The most obvious difference between the two cerebral areas was that there was no statistically significant difference in N-acetyl aspartate (NAA) levels in the right cerebellum at all time points; however, the amino acid levels of NAA in the left cerebral hemisphere were markedly decreased 3, 9 and 24 h post-MCAO. In addition, an obvious increase in glutamine was observed in the right and left cerebellum at 3, 9 and 24 h post-MCAO. Furthermore, the present study demonstrated that γ-aminobutyric acid levels were decreased at 1 h in the left and right cerebellum and were evidently increased at 24 h in the right cerebellum post-MCAO. In conclusion, supratentorial ischemia has been indicated to affect the activities of the non-ischemic contralateral cerebellum. Therefore, these results suggested that an NMR-based metabonomic approach may be used as a potential means to elucidate cerebral and cerebellar metabolism following MCAO, which may help improve understanding regarding cerebral infarction at a molecular level. *Ex vivo* ¹H NMR analysis may be useful for the assessment of clinical biopsies.

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Abbreviations: MCAO, middle cerebral artery occlusion; ¹H NMR, proton nuclear magnetic resonance; CCD, crossed cerebellar diaschisis; TCA, tricarboxylic acid; CCA, common carotid artery; ECA, external carotid artery; ICA, internal carotid artery; PLS-DA, partial least squares discriminate analysis; NAA, N-acetyl aspartate; Lac, lactate; Ala, alanine; GABA, γ-aminobutyric acid; Glu, glutamate; Gln, glutamine; Suc, succinate; Asp, aspartate; Cre, creatine; m-Ins, myo-inositol; Cho, choline; Gly, glycine; ¹H MRS, ¹H magnetic resonance spectroscopy

Key words: middle cerebral artery occlusion, ischemia, rat, cerebellum, metabolites, ¹H NMR

Introduction

Acute ischemic stroke is a type of focal brain injury, which causes functional depression due to a disruption in normal signal propagation between the ischemic area and regions that are connected to it by nerve fiber bundles (1). Crossed cerebellar diaschisis (CCD), which was first mentioned by Baron *et al* (2), is a condition in which blood flow and metabolism on the side contralateral to a damaged cerebral area are decreased (3). It has previously been reported that an interruption in corticopontocerebellar pathways is the most likely mechanism underlying CCD (3-5). Following a supratentorial stroke, cortical excitability cannot be transmitted to the contralateral cerebellum due to pathway disruption, which leads to functional inhibition and a decrease in metabolism in the contralateral cerebellar hemisphere (6). To date, numerous techniques, including single photon emission computed tomography, positron emission tomography, dynamic susceptibility contrast magnetic resonance (MR) perfusion imaging,

arterial spin-labeling MR imaging and computed tomography perfusion, have been used to estimate CCD within stroke patients (7-14). These techniques attempt to diagnose CCD based on the rate of regional cerebral blood flow and oxygen metabolism in the brain; however, the mechanisms underlying CCD remain unclear.

Regional metabolic differences in the mammalian brain, including glucose and glycogen stores, have been detected in *ex vivo* analyses (15), and have also been determined from non-invasive measurements in humans and mice (16-18). Håberg *et al* (1) reported that glucose metabolism and the metabolic activity of intermediates from the astrocytic tricarboxylic acid (TCA) cycle were markedly decreased in the whole rat cerebellum in the superacute stage of middle cerebral artery occlusion (MCAO), as determined using ^{13}C MR spectra. In addition, it was demonstrated that the cerebellum could control hemispheric activity. Therefore, it may be suggested that, to enhance the recovery of cerebral hemispheric function, it would be beneficial to maintain the cerebellum in a low-activity state. However, this previous study did not explore metabolism in the bilateral cerebellum or observe variations between left and right sides. Previous studies have reported that CCD may not be just a concomitant phenomenon of stroke, but may be regarded as a crucial prognostic indicator, which may benefit the treatment and rehabilitation of brain ischemia (10,19).

The present study aimed to identify the effects of MCAO on alterations in cerebral metabolism in the ischemic brain regions and in the contralateral cerebellum in rats 1, 3, 9 and 24 h following ischemia using proton nuclear MR (^1H NMR) spectroscopy. In addition, the study aimed to: i) Evaluate the regional metabolic differences induced by CCD between the ischemic cerebral hemisphere and the contralateral cerebellum; and ii) to identify the mechanisms underlying CCD.

Materials and methods

Animal preparation and treatment. A total of 38 male Sprague-Dawley rats (8-10 weeks old; 250-320 g; Shanghai SLAC Laboratory Animal Co., Ltd., Shanghai, China) were maintained in the Specific-Pathogen-Free Animal Experimental Center of Wenzhou Medical University (Wenzhou, China). All rats were kept under a temperature of $23\pm 20^\circ\text{C}$ and a relative humidity of $55\pm 10\%$, and were maintained under a 12-h light/dark cycle with free access to food and drink. The present study was approved by the Animal Ethics Committee of Wenzhou Medical University and was strictly conducted according to the National Institutes of Health Guide for the Care and Use of Laboratory Animals (20).

Development of the MCAO model. An MCAO model was developed using the intraluminal filament technique, as previously described (21). After 12 h of fasting, the rats were anesthetized with 10% chloral hydrate (300 mg/kg; intraperitoneal). Initially, an incision was made into the middle cervical fascia and the left common carotid artery (CCA) was exposed. The external carotid artery (ECA) and internal carotid artery (ICA) were then separated. Subsequently, the bifurcation close to the ECA was ligated with the filament. A ready-made suture (Beijing Sunbio Biotech Co., Ltd., Beijing, China) was inserted

via the left CCA into the ICA, in order to occlude the MCA. The depth of the suture within the vessel was 16-18 mm and the redundant part was cut off with a ligature. In this procedure, the suture was maintained around the vessel. Finally, the incision was stitched and the rats were fed separately to improve survival rate. The temperature was maintained at $25\text{-}26^\circ\text{C}$ during the surgery. In the sham operation group ($n=10$), the neck was incised to expose the left CCA; however, the MCA was not occluded. MCAO rats ($n=28$) were randomly sacrificed by prompt decapitation at the following time points: 1, 3, 9 and 24 h after MCAO ($n=7/\text{group}$). The rats in the sham operation group were decapitated 24 h after surgery and were compared with the MCAO rats at all other time points. Tissue specimens were obtained from the left cerebral hemisphere, and the left and right cerebellum, within 15 sec; tissue specimens were frozen at -80°C .

Preparation of cerebral samples. The frozen brain tissues were weighed and homogenized in centrifuge tubes using an electric homogenizer. The samples were then vortexed with 4 ml/g ice-cold methanol and 0.85 ml/g distilled water. Subsequently, 2 ml/g chloroform and 2 ml/g distilled water were added to the tubes and mixed again. The specimen tubes were placed on ice for 15 min and were then centrifuged at $12,000 \times g$ for 15 min at 4°C . The supernatant was separated from the tubes and placed into a freeze-dryer to lyophilize for 24 h. Finally, the obtained extracts were dissolved in 500 μl 99.5% D_2O for NMR spectroscopy.

Acquisition of ^1H NMR spectra. All ^1H NMR spectra of the extracts were obtained at 25°C with a 90° flip angle on a spectrometer (Bruker Avance III 600-MHz; Bruker Corporation, Billerica, MA, USA). The spectral width was set at 12,000 Hz and 32 K data points. The collection time was 2.66 sec per scan and the number of scans was 128. In order to assure full relaxation, an extra 8 sec relaxation delay was set. Exponential line-broadening of 0.3 Hz was used in the free induction decay ahead of Fourier transformation and the spectra were zero-filled to 64 K. All spectra were carefully corrected by hand for phase as well as baseline. In addition, the methyl peak of lactate (Lac) (CH_3 ; 1.33 ppm) was used as a reference point for the spectra. Peak area integration was conducted using the Bruker Topspin software package (version 2.1; Bruker Corporation) with standard routines.

Data and statistical analysis. NMR spectra ($\delta 0.5\text{-}10.0$) were segmented into integral intervals with each width of 0.01 ppm (2.4 Hz) through AMIX package (Bruker Topspin 2.1; Bruker Corporation), so that all metabolic information embedded in the spectra could be exploited; the sum of each spectrum was then standardized. The normalized integral values were mean-centered for multivariate data analysis using software (Umetrics SIMCA-P+12.0; Sartorius Stedim Data Analytics AB, Umeå, Sweden). Partial least squares discriminate analysis (PLS-DA) was conducted to identify metabolites according to the separation of different groups (22). Data were visualized using a principal component scores plot of the first two principal components to provide the most efficient 2D representation of the information (20). Data were confirmed in 2D $^1\text{H}\text{-}^1\text{H}$ correlation spectroscopy and total correlation

spectroscopy spectra. The position of each point represents an individual spectrum of a sample. Differences in the sample compositions between the different groups were determined by PLS-DA, and differences in the metabolites between the groups were revealed as coefficient of variation plots (23). Model quality was assessed with the fitness of model (R^2) and the predictive ability of model (Q^2).

All values are presented as the mean \pm standard deviation. SPSS software (version 13.0; SPSS, Inc., Chicago, IL, USA) was used to determine the statistical differences between groups using one-way analysis of variance followed by a least significant difference post hoc test for multiple comparisons. $P < 0.05$ was considered to indicate a statistically significant difference.

Results

^1H NMR spectral analysis of samples. ^1H NMR was used to investigate metabolism within the contralateral side to the damaged area; the typical ^1H NMR spectra of the right rat cerebellum samples in the sham operation, and 1, 3, 9 and 24 h MCAO model groups are presented in Fig. 1. The allocation of metabolites on the spectrograms was based on our previous work (24). 2D ^1H - ^1H correlation spectroscopy and total correlation spectroscopy of the representative samples were performed to confirm the allocations on the ^1H NMR spectra. Numerous endogenous metabolites were simultaneously observed on the ^1H NMR spectra of cerebral samples.

Pattern recognition of cerebral extracts. The ^1H NMR data of the left cerebral hemisphere and the right cerebellum were used to determine differences between metabolic profiling of rats in the MCAO and sham groups after 1, 3, 9 and 24 h by multivariate data analysis. PLS-DA score plots displayed a prominent separation between the sham and MCAO groups along the direction of $t(1)$ in the left hemisphere (Fig. 2) and the right cerebellum (Fig. 3A), revealing a significant metabolic disturbance. However, with the increase in ischemic duration, the alteration in metabolic patterns moved gradually away from the $t(1)$ direction in the left cerebral hemisphere (Fig. 2), which was not detected in the right cerebellum at 9 h (Fig. 3B). Such alterations indicated that the two brain regions have differences in metabolic pattern.

The comparisons between rats in the sham and 1 h model groups, and the sham and 24 h model groups are presented in Figs. 4 and 5, respectively. The PLS-DA score plots of the sham operation group and the 1 h model group, and the sham operation group and the 24 h model group are presented in Figs. 4A and 5A, respectively. The model groups may be separated from the sham operation group along the first principal components horizontal direction. The results demonstrated that there was an obvious difference between the two groups with regards to the spectral features in the right cerebellum. In addition, the validation graph of permutation tests indicated that the PLS-DA models were robust and credible (Figs. 4B and 5B).

Figs. 4C and 5C illustrate the corresponding loading plots of metabolites between the MCAO and sham groups using color-coded correlation coefficients at 1 and 24 h in the right cerebellum. The findings indicated that the separation of the

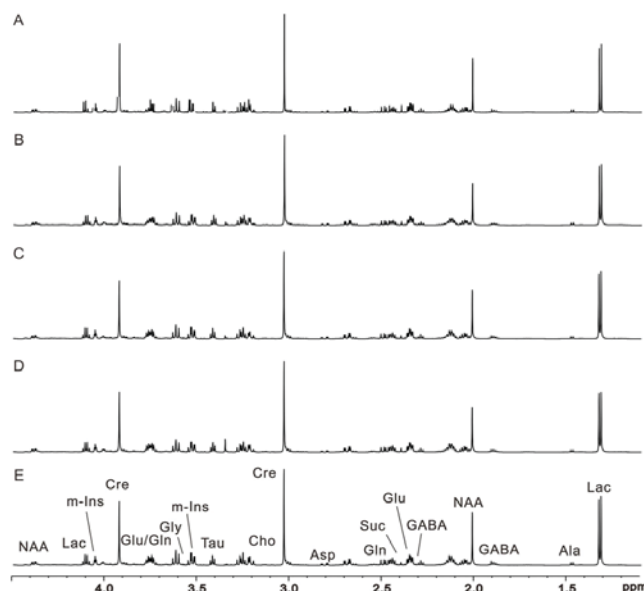


Figure 1. Representative 600-MHz proton nuclear magnetic resonance spectra of right cerebellum extracts obtained from rats in (A-D) the model groups (1, 3, 9 and 24 h after middle cerebral artery occlusion, respectively) and (E) the sham operation group. Ala, alanine; Asp, aspartate; Cre, creatine; Cho, choline; GABA, γ -aminobutyric acid; Gln, glutamine; Glu, glutamate; Lac, lactate; m-Ins, myo-inositol; NAA, N-acetyl aspartate; Suc, succinate.

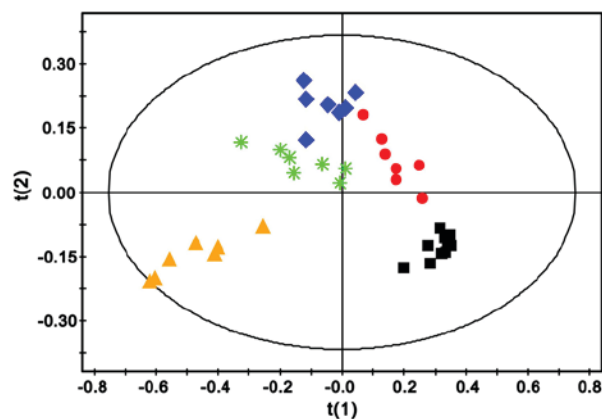


Figure 2. Partial least squares discriminate analysis score plots of the first and second principal components based on the proton nuclear magnetic resonance spectra of the left cerebral hemisphere samples obtained from rats in the sham operation (\blacksquare), MCAO 1 h (\bullet), MCAO 3 h (\blacklozenge), MCAO 9 h (\ast) and MCAO 24 h (\blacktriangle) groups. MCAO, middle cerebral artery occlusion.

different groups may be due to variation in metabolite levels. The square of the correlation coefficient was used as the weight of a variable, and color-coding was used to indicate low and high values (low, blue; high, red). An increase in the corresponding metabolites in MCAO rats was displayed in the negative area, whereas a decrease in the corresponding metabolites was displayed in the positive area. The results demonstrated that MCAO rats had lower levels of N-acetyl aspartate (NAA), creatine (Cre), glutamate (Glu) and succinate (Suc), and higher levels of lactate (Lac), γ -aminobutyric acid (GABA) and glutamine (Gln) compared with the control groups. The results at the other time points are presented in Figs. 6 and 7, and were in accordance with those presented in Figs. 4 and 5.

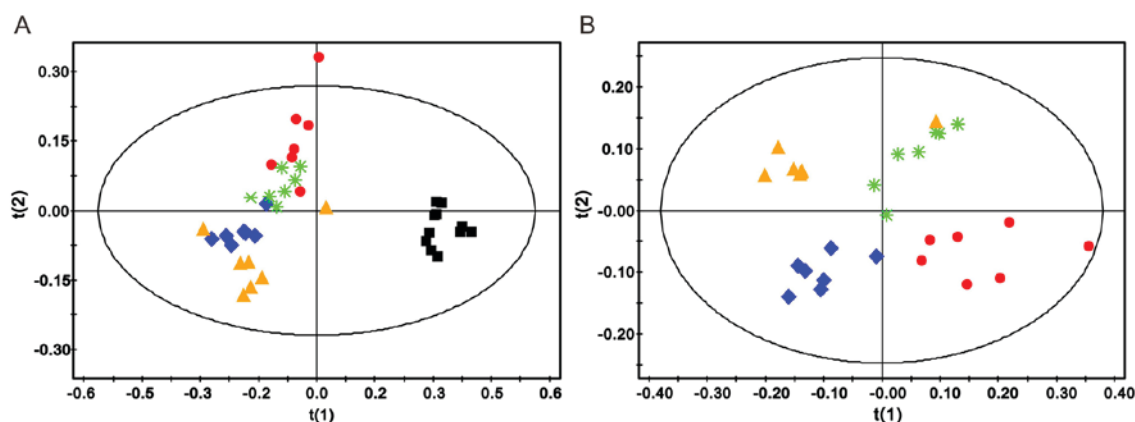


Figure 3. (A) Partial least squares discriminate analysis score plots of the first and second principal components based on the proton nuclear magnetic resonance spectra of the right cerebellum samples obtained from rats in the sham operation (■), MCAO 1 h (●), MCAO 3 h (◆), MCAO 9 h (◊) and MCAO 24 h (▲) groups. (B) Partial least squares discriminate analysis score plots of the first and second principal components based on the proton nuclear magnetic resonance spectra of the right cerebellum samples obtained from rats in the MCAO 1 h (●), MCAO 3 h (◆), MCAO 9 h (◊) and MCAO 24 h (▲) groups. MCAO, middle cerebral artery occlusion.

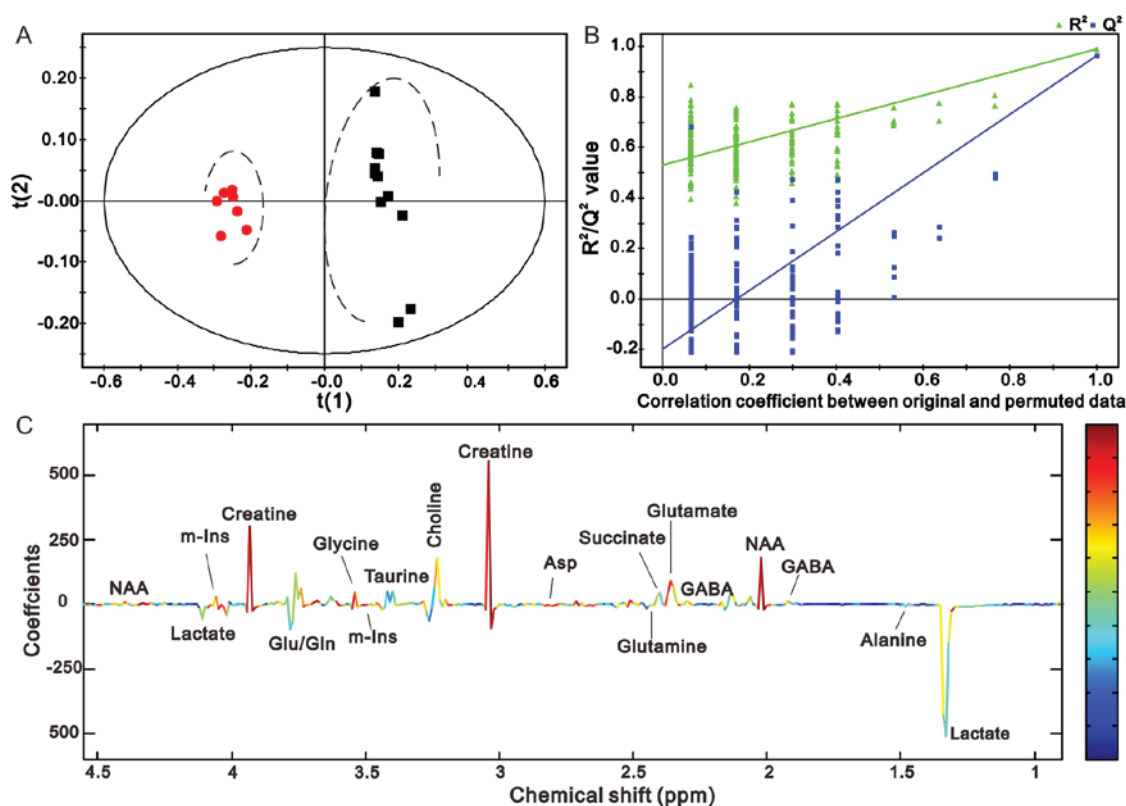


Figure 4. (A) PLS-DA score plot ($R^2=0.099$, $Q^2=0.97$) and (B) validation plot based on the proton nuclear magnetic resonance spectra of right cerebellum samples obtained from rats in the middle cerebral artery occlusion 1 h (●) and sham operation (■) groups. (C) Coefficient-coded loading plot corresponding to the PLS-DA plot revealed that metabolites with varying intensities between the groups were responsible for the separation between the groups in the corresponding score plot. Asp, aspartate; GABA, γ -aminobutyric acid; m-Ins, myo-inositol; Gln, glutamine; Glu, glutamate; NAA, N-acetyl aspartate; PLS-DA, partial least squares discriminate analysis.

Alterations in the levels of cerebral metabolites. Metabolite levels were quantified, in order to investigate the metabolic alterations in left cerebral tissue (Fig. 8). The results demonstrated that the levels of GABA, glycine (Gly), choline (Cho), Lac and alanine (Ala) were markedly increased in the ischemic cerebral hemisphere of rats compared with in the control group. The levels of Gln were not markedly increased until 9 h post-MCAO. The levels of Glu, aspartate (Asp),

NAA and Cre were markedly decreased in the left cerebral hemisphere 3, 9 and 24 h post-MCAO. In addition, the levels of Suc were decreased 24 h post-MCAO, and an obvious decrease in myo-inositol (m-Ins) levels were also detected 24 h after ischemic insult in the left cerebral hemisphere.

Conversely, ischemia induced marked increases in the levels of Gln, Asp and Lac, and concomitant decreases in the levels of Suc and Cre in the right cerebellum at all studied

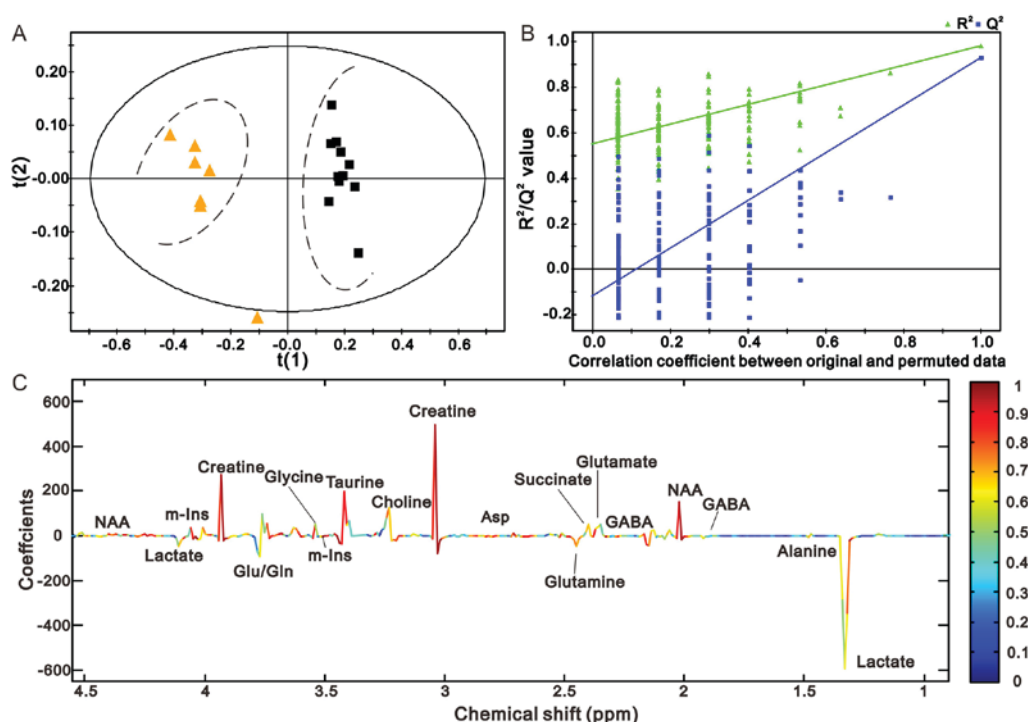


Figure 5. (A) PLS-DA score plot ($R^2=0.98$, $Q^2=0.93$) and (B) validation plot based on the proton nuclear magnetic resonance spectra of right cerebellum samples obtained from rats in the middle cerebral artery occlusion 24 h (▲) and sham operation (■) groups. (C) Coefficient-coded loading plot corresponding to the PLS-DA plot revealed that metabolites with varying intensities between the groups were responsible for the separation between the groups in the corresponding score plot. Asp, aspartate; GABA, γ -aminobutyric acid; m-Ins, myo-inositol; NAA, N-acetyl aspartate; PLS-DA, partial least squares discriminate analysis.

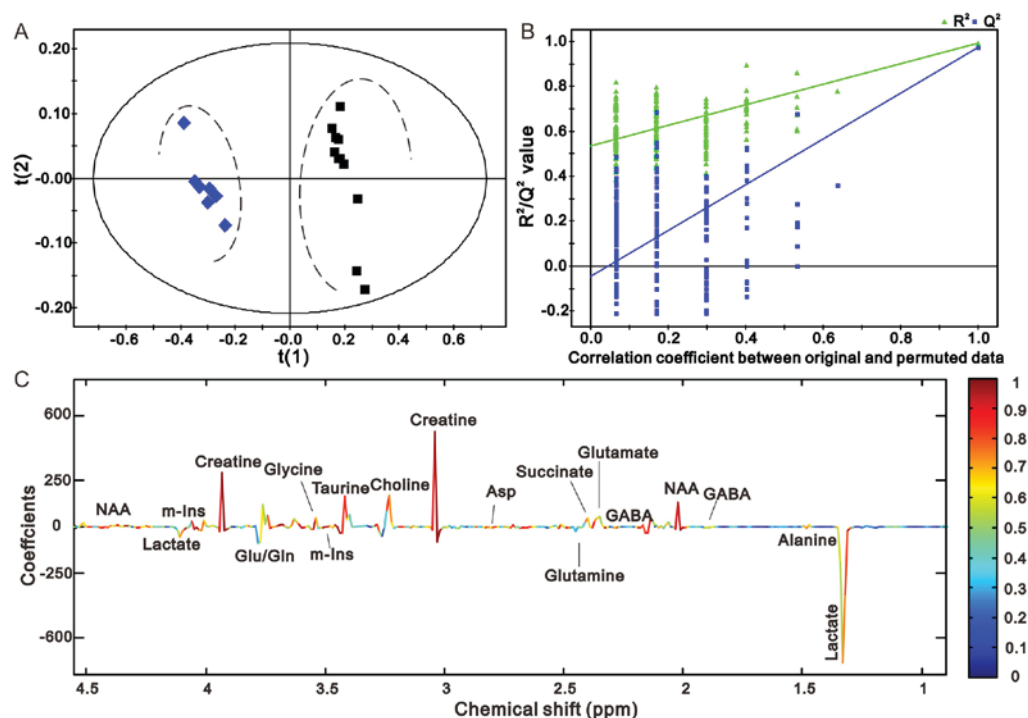


Figure 6. (A) PLS-DA score plot ($R^2=0.99$, $Q^2=0.97$) and (B) validation plot based on the proton nuclear magnetic resonance spectra of right cerebellum samples obtained from rats in the middle cerebral artery occlusion 3 h (◆) and sham operation (■) groups. (C) Coefficient-coded loading plot corresponding to the PLS-DA plot revealed that metabolites with varying intensities between the groups were responsible for the separation between the groups in the corresponding score plot. Asp, aspartate; GABA, γ -aminobutyric acid; m-Ins, myo-inositol; NAA, N-acetyl aspartate; PLS-DA, partial least squares discriminate analysis.

time points post-MCAO (Fig. 9); however, there were no significant differences in NAA levels between the groups

(data not shown). Cho levels were not markedly increased until 24 h post-MCAO. In addition, Ala levels were increased

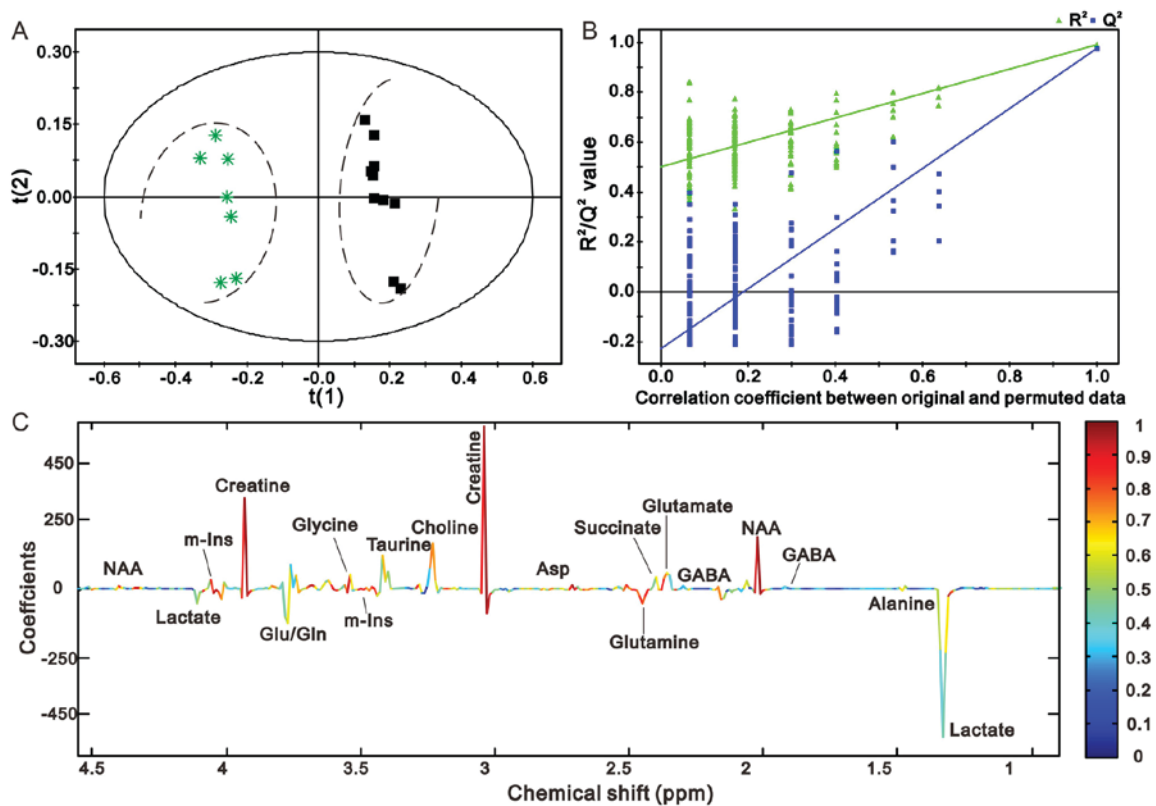


Figure 7. (A) PLS-DA score plot ($R^2=0.99$, $Q^2=0.97$) and (B) validation plot based on the proton nuclear magnetic resonance spectra of right cerebellum samples obtained from rats in the middle cerebral artery occlusion 9 h (*) and sham operation (■) groups. (C) Coefficient-coded loading plot corresponding to the PLS-DA plot revealed that metabolites with varying intensities between the groups were responsible for the separation between the groups in the corresponding score plot. Asp, aspartate; GABA, γ -aminobutyric acid; m-Ins, myo-inositol; NAA, N-acetyl aspartate; PLS-DA, partial least squares discriminate analysis.

at 1 h post-MCAO, and GABA levels were elevated at 3 and 24 h following ischemic insult. Conversely, Glu levels were significantly decreased at 1 and 3 h post-MCAO.

As presented in Fig. 10, there were variations in Glu, Gln, GABA and total levels (Gln + Glu + GABA), and the Glu/GABA and Glu/Gln ratios in the non-ischemic left cerebellum and right cerebellum 1, 3, 9, and 24 h post-MCAO and sham operation. The alterations in Glu/Gln ratio and Gln + Glu + GABA had just the same trend between the right and left cerebellum; however, for Glu and Glu/GABA ratio, a similar trend was observed between the right and left cerebellums. A marked increase in Gln levels and a concomitant decrease in the Glu/Gln ratio was detected in both regions at all studied time points post-MCAO. In addition, an obvious increase in Gln was observed in the right and left cerebellum at 3, 9 and 24 h post-MCAO. Furthermore, the present study demonstrated that GABA levels were decreased at 1 h in the left and right cerebellum and were evidently increased at 24 h in the right cerebellum post-MCAO.

Discussion

Since the brain has high sensitivity to ischemic hypoxia, focal ischemia can result in a reduced supply of glucose and oxygen to the corresponding brain areas, resulting in disruption to the TCA cycle (25). Consequently, the levels of Lac, as the main product of anaerobic glycolysis, were markedly increased in the present study in response to ischemia, which is in agreement with the results of previous studies (26-29).

^1H MR spectroscopy (MRS) has been widely used to research the pathological mechanism underlying neuronal and cerebral metabolic alterations in response to cerebral ischemia in humans and animals (26,30,31). Alterations in the spectral peaks of patients with acute ischemic stroke may have prognostic value in clinical practice (32,33).

The present study aimed to use ^1H NMR spectroscopy to analyze the metabolic alterations of CCD between the left cerebral hemisphere and the contralateral cerebellum in rats after permanent MCAO. The main finding of the present study was that metabolic alterations were detected in the contralateral cerebellum, which is a region involved in CCD, as determined using PLS-DA. The results indicated that: i) Focal ischemia induced marked increases in the levels of Lac, Ala, Gln and GABA, and decreases in the levels of Cre, Suc and Glu in the left cerebral hemisphere and the right cerebellum; ii) supratentorial ischemia induced metabolic alterations between left and right cerebellum, particularly in the contralateral cerebellum, iii) alterations in Glu metabolism may be associated with CCD; however, further studies are required.

Glu and Asp are the major excitatory amino acids, which have significant roles in the central nervous system. In previous studies, dynamic equilibrium of excitatory and inhibitory amino acids (Glu and GABA) was elevated to the highest level 1-2 h after ischemia; however, it was decreased 3 h after ischemic injury in rats (34,35). The present study detected alterations in the levels of Glu, Asp, NAA and Cre at 3, 9 and 24 h in the brain of ischemic rats. In addition, GABA and Gly levels were

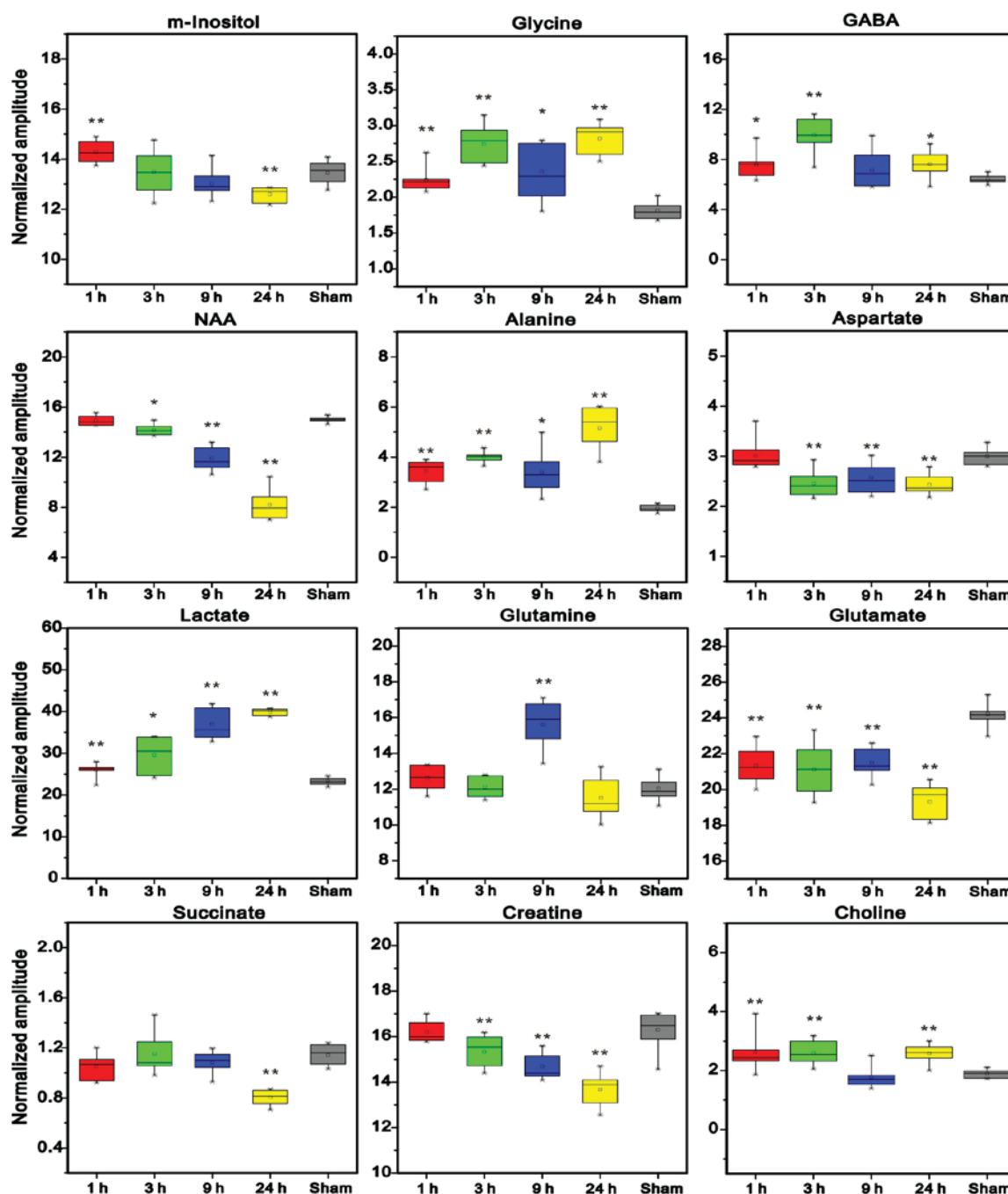


Figure 8. Relative normalized concentrations of the significantly altered metabolites in the left ischemic cerebral hemisphere. Red, green, blue, yellow and grey represent relative normalized concentrations in the middle cerebral artery occlusion 1, 3, 9 and 24 h, compared with sham operation groups, respectively. * $P < 0.05$; ** $P < 0.01$ vs. the sham group. GABA, γ -aminobutyric acid.

significantly increased at 1, 3 and 24 h post-MCAO, which indicated that GABA and Gly had protective effects on ischemic insult in the rat brain against excitatory amino acid toxicity.

Glu is a neurotransmitter that is released by neurons and can be absorbed by astrocytes, where it is converted into Gln. Subsequently, Gln can be transferred to neurons and once again converted to Glu. Furthermore, Gln is a major precursor of neuronal Glu and GABA. This important circulatory pathway between astrocytes and neurons is known as the Gln-Glu-GABA cycle (36). Elevated levels of GABA and Gln, and decreased levels of Glu and Asp, were observed in the ischemic cerebral hemisphere of MCAO rats in the present study,

which was consistent with the results of a previous study (37). A recent study indicated that the potential mechanisms underlying a decrease in Glu levels may be associated with increased utilization and decreased synthesis (1). Furthermore, Glu may undergo retrograde transport in axons (38). Therefore, it may be hypothesized that excitotoxic action in the ischemic region is increased if Glu is transported from the non-ischemic cerebellum to the ischemic cerebral hemisphere. In the present study, Glu concentrations were significantly decreased in the left cerebral hemisphere 1, 3, 9 and 24 h post-MCAO; however, in the right cerebellum Glu was markedly decreased at 1 and 3 h post-MCAO, but was increased at 9 h, without reaching

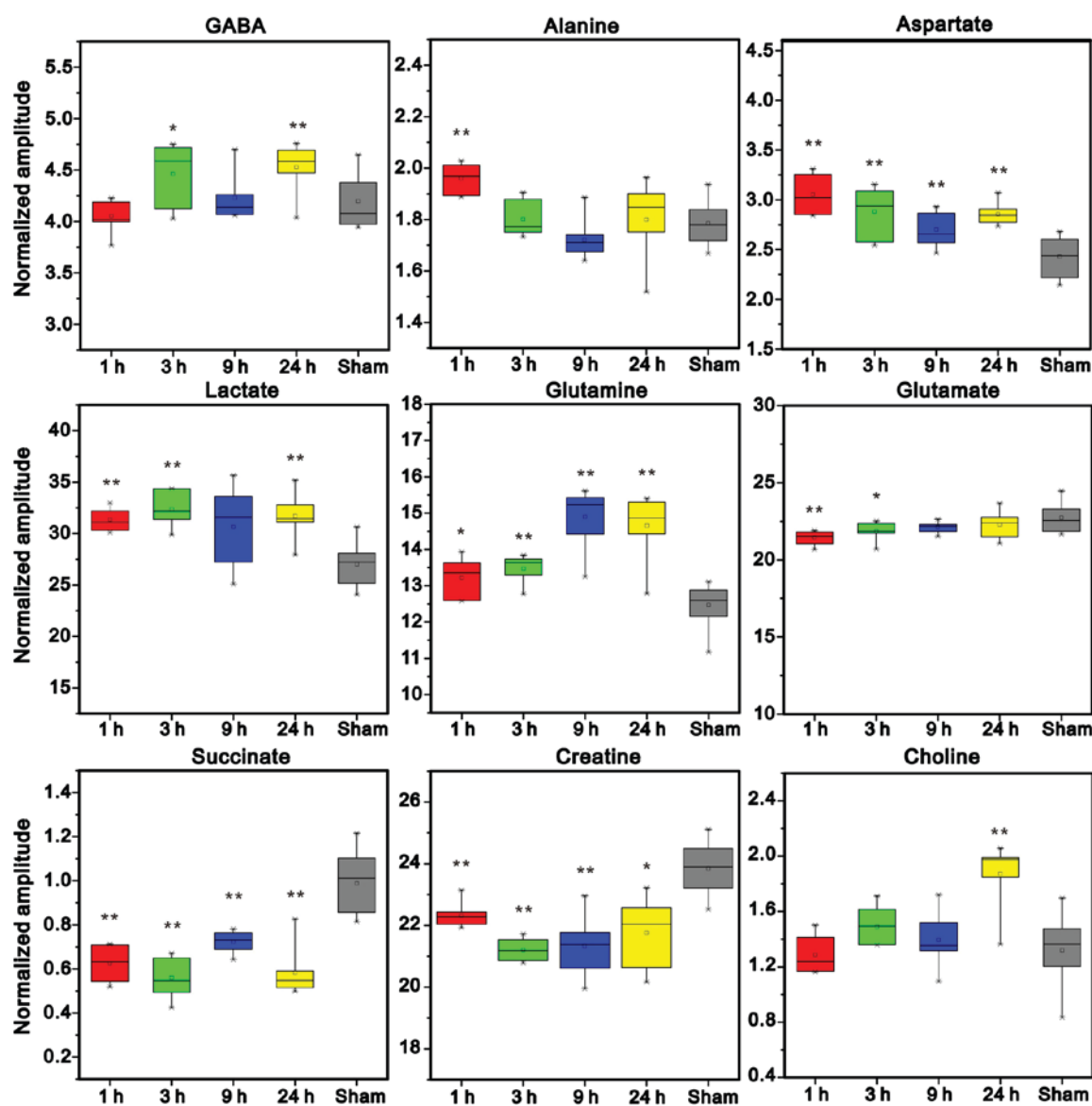


Figure 9. Relative normalized concentrations of the significantly altered metabolites in the right cerebellum. Red, green, blue, yellow and grey bar charts represent relative normalized concentrations in the middle cerebral artery occlusion 1, 3, 9 and 24 h, compared with sham operation groups, respectively. * $P < 0.05$; ** $P < 0.01$ vs. the sham group. GABA, γ -aminobutyric acid.

statistical significance. In addition, according to the PLS-DA score plots, as the duration of ischemia increased, the metabolic pattern in the left cerebral hemisphere moved gradually away from the sham group along the $t(1)$ direction; however, this was not the case for the metabolic pattern in the right cerebellum at 9 h. It may be hypothesized that Glu was not transferred from the non-ischemic right cerebellum to the left ischemic brain at 9 h post-MCAO. As a result, the excitotoxic burden in the ischemic region is decreased. This finding also suggested that CCD is reversible 9 h after ischemic injury. The metabolic alterations of Glu may be associated with CCD. It has previously been reported that following permanent ischemia, metabolic alterations can be longitudinally followed using *in vivo* localized ^1H MRS (39,40). Berthet *et al* (40) detected marked metabolic alterations in the ischemic core following permanent focal ischemia, including increases in GABA, Gly and Cre, and decreases in Gln, Glu and NAA. However, the results of the present study, including the increases in Gln levels in the ischemic cerebral hemisphere,

were not consistent with the previous *in vivo* results. Further studies are required to assess these inconsistencies.

NAA is regarded as a sensitive marker of neuronal function (41). The levels of NAA were significantly decreased in the ischemic cerebral hemisphere of rats. This indicated that the ischemic brain damage may lead to obvious neuronal dysfunction. Cre is a biological marker for the energy metabolism of neurons (42,43). In the present study, the levels of Cre were markedly decreased in the left cerebral hemisphere and the right cerebellum, which revealed that energy metabolism of the brain was disordered in the ischemic rats. Cho is involved in lipid metabolism and membrane function (44). An increase in the levels of Cho in the brain suggests a significant alteration in membrane metabolism caused by ischemia. In addition, the reduction in the levels of m-Ins detected in the present study was consistent with previous findings (45) and may be a suggestive of alterations in the local osmotic pressure of cells induced by the ischemia.

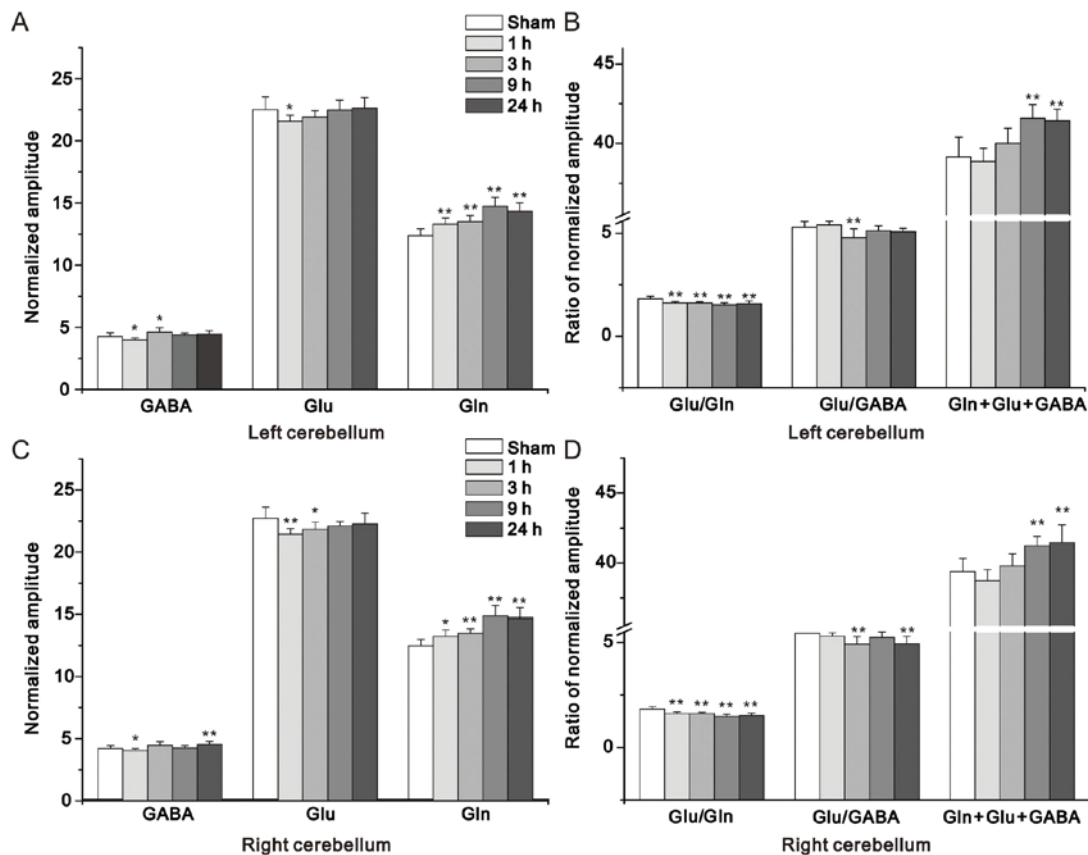


Figure 10. Summed concentration of Glu, Gln, GABA, Glu/Gln ratio, Glu/GABA ratio and Gln + Glu + GABA in (A and B) the non-ischemic left cerebellum and (C and D) right cerebellum in the 1, 3, 9 and 24 h middle cerebral artery occlusion, compared with sham groups. *P<0.05; **P<0.01 vs. the sham group. GABA, γ -aminobutyric acid; m-Ins, myo-inositol; Gln, glutamine; Glu, glutamate.

The effects of ischemia on metabolism in the ischemic cerebral hemisphere and in other brain regions have been reported in a previous *ex vivo* analysis (1) and in an *in vivo* study (46). However, some of the findings from the ischemic cerebral hemisphere in the present study were inconsistent with the findings of other studies regarding permanent MCAO (1,39,40,47), which may be due to numerous factors, as listed below.

Firstly, there are differences between the methods used to generate successful models of ischemia, including the control of cerebral blood circulation for filament insertion-induced focal ischemia (5). This may result in differences in the metabolic response to permanent ischemia in the cerebral cortex, consequently resulting in increased variability in metabolic results (21,39,40).

Secondly, fixation of cerebral tissue following decapitation has potential postmortem effects on Lac, Cre and GABA, which are key components used for PLS-DA in the present study. It is well known that fixation and extraction procedures intrinsically affect metabolic results. For example, decapitation is known to induce postmortem effects on metabolism (2,48,49). Typically, highly elevated Lac is expected following decapitation, due to the degradation of glucose and glycogen in the brain, and may also occur during extraction procedures. In addition, GABA is known to increase and the levels of phosphocreatine are known to immediately diminish after decapitation (48). The effects of fixation following decapitation have also been observed *in vivo* in the ischemic core following permanent ischemia using localized ^1H MRS in a horizontal 600MHz magnet (40). Therefore, the

majority of the significant metabolic alterations induced by permanent ischemia in the present study may have been affected by the fixation method used; for example, highly elevated Lac levels were detected in MCAO rats compared with in sham-operated rats and therefore the difference in Lac between sham-operated and ischemic rats was significantly reduced. In addition, a three-fold elevation of GABA (40) *in vivo* following permanent ischemia was reduced; however, the difference was not so marked *ex vivo* in the present study. Alternatively, microwave fixation (50) and funnel-freeze fixation (51), with careful extraction procedures, have been reported to reserve all carbohydrates, including glucose and glycogen, as illustrated by the lower Lac levels detected in the control animals.

Finally, the selection of specimens may have effects on the results. For example, the effects of focal ischemia were limited to only part of one side of the cerebral hemisphere in Igarashi *et al* (26), Berthet *et al* (40) and Håberg *et al* (1). Therefore, the results from the selected specimen (part of the left cerebral hemisphere) would intrinsically provide information regarding the metabolic alterations in the ischemic cerebral hemisphere and in some non-ischemic regions. This may explain the discrepancies between the metabolic alterations detected in the present study compared with other studies (1,39,40,47).

In conclusion, the present study used ^1H NMR-based metabolomics to evaluate cerebral metabolism in the ischemic cerebral hemisphere and the non-ischemic cerebellum post-MCAO in rats. The results indicated that focal ischemia affected non-ischemic cerebellum activities, neurotransmitter

synthesis and metabolic balance, particularly in the contralateral cerebellum. In addition, metabolic activity in the cerebellum, particularly with regards to Glu, may serve an important role in brain function reconstruction, which may help improve understanding regarding cerebral infarction on a molecular level. Alterations in Glu metabolism may be associated with CCD; however, this requires further experimentation.

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