

Hypoxia-Ischemia Stimulates Hippocampal Glutamate Efflux in Perinatal Rat Brain: An *In Vivo* Microdialysis Study¹

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ABSTRACT. We used *in vivo* microdialysis to determine the impact of a focal hypoxic-ischemic insult on hippocampal amino acid efflux in the immature brain. Microdialysis probes were inserted into the right hippocampus of postnatal d 7 rats. To induce hypoxic-ischemic injury, the right carotid artery was ligated and animals were exposed to 8% oxygen for 2 h ($n = 6$, histologically verified). Ten 20-min dialysis fractions were collected from each animal: three sequential 20-min baseline samples, six samples during hypoxia, and a recovery sample in room air. Eight amino acids were detected in dialysates with a HPLC assay. There was marked intra- and interanimal variation in glutamate efflux; mean glutamate efflux increased from 17 pmol/min in baseline samples to 51 in the 2nd h of hypoxia ($p = 0.002$, Kruskal Wallis test). There was a concurrent decline in glutamine efflux (310 to 207 pmol/min, $p = 0.0005$). Alanine efflux doubled during hypoxia ($p = 0.015$). There were no changes in efflux of the other five amino acids analyzed. In this experimental model of perinatal stroke, during the acute evolution of hypoxic-ischemic brain injury, transient large increases in glutamate efflux and corresponding declines in glutamine efflux were detected. These data support the hypothesis that synaptic concentrations of the endogenous excitatory amino acid glutamate increase during the evolution of hippocampal ischemic injury. (*Pediatr Res* 30: 587-590, 1991)

Abbreviations

EAA, excitatory amino acids
PND, postnatal day
ECF, extracellular fluid

Considerable experimental evidence indicates that endogenous EAA contribute to the pathogenesis of ischemic hippocampal injury (1-4). In perinatal brain, the immaturity of hippocampal synaptic circuitry might suggest that the contribution of endogenous EAA to the evolution of ischemic neuronal injury would be reduced, as compared with the adult. However, because PND 7 rats are susceptible to excitotoxic neuronal damage (5, 6), elevated synaptic concentrations of endogenous glutamate or aspartate could be neurotoxic at this developmental stage. In a perinatal rodent stroke model, *i.e.* stroke induced by right carotid artery ligation followed by exposure to moderate hypoxia in

PND 7 rats (7), considerable data suggest that endogenous glutamate plays an important role in the pathogenesis of brain injury; for example, striatal ECF glutamate content peaks transiently during the evolution of injury (8), and systemically administered glutamate antagonists attenuate the severity of tissue damage (9-11).

The hippocampus is particularly susceptible to ischemic neuronal injury. In this perinatal model, the propensity for selective CA1 pyramidal cell damage that characterizes adult ischemia models (12) is not observed (6), and the distribution of hippocampal histopathology is more variable.

In this study we used *in vivo* microdialysis to examine hippocampal amino acid efflux during the evolution of hypoxic-ischemic injury. With *in vivo* microdialysis, sequential changes in ECF content of amino acids can be readily estimated within defined brain regions (13, 14). ECF accumulation of neurotransmitter amino acid reflects regional synaptic concentrations; increased concentrations of EAA could result from increased release [from neurotransmitter or metabolic pools (1-4, 15, 16)] or suppression of neuronal and glial reuptake (17, 18). Calculated efflux rates provide estimates of regional ECF content of compounds of interest.

We measured hippocampal amino acid efflux to determine if EAA efflux increased during the evolution of a focal hypoxic-ischemic injury in the immature brain.

MATERIALS AND METHODS

Microdialysis probes were implanted into hippocampi of PND 7 Sprague-Dawley rats, using a modification of our previously described methods for striatal dialysis (19, 20). Probes had a concentric design and a 2-mm dialyzing tip (diameter 250 μ m, molecular weight cutoff 6000; Spectra-Por, Spectrum Medical, Los Angeles, CA). In preliminary experiments, we attempted to optimize stereotaxic coordinates for probe insertion. The site of probe placement was verified by postmortem examination of sequential horizontal brain sections and was considered acceptable if a discrete (probe-induced) anatomical defect was identified within the right hippocampus.

Coordinates used to guide probe insertion were (referenced to bregma): 3.5 mm lateral, 3.5 mm posterior, and 3 mm depth. Although these coordinates were accurate for intrahippocampal stereotaxic injections, they were not as reliable for sustained probe placement; the overall success rate for completion of sampling with verification of probe placement within hippocampus was <50%. Figure 1 demonstrates the most common site of probe placement within midhippocampus, adjacent to the superior blade of the dentate gyrus; however, using the same coordinates, probe placement sites were also identified more medially, adjacent to the CA1 pyramidal cell region and more laterally adjacent to CA3. The curvature of the skull surface made stabilization of the probe more difficult than for striatal insertions, and in about one fourth of the animals mechanical problems

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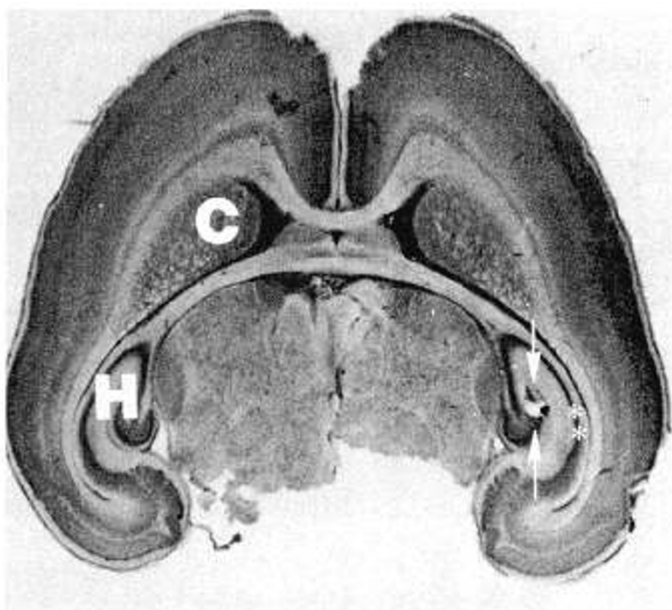


Fig. 1. Photograph of a cresyl violet-stained 50- μ m horizontal brain section prepared from an unlesioned PND 7 rat that underwent *in vivo* hippocampal microdialysis. The anatomical defect resulting from probe insertion is evident in the superior blade of the dentate gyrus (arrowheads) of the right hippocampus. C, caudate; H, hippocampus; **, CA1 pyramidal cell layer.

prevented completion of experiments. Based on these factors and on our considerable striatal microdialysis data that demonstrated stable amino acid efflux over up to 4 h, we limited analysis to hypoxic-ischemic animals.

In ether-anesthetized animals, a midline scalp flap was excised and the right carotid artery ligated in about 5 min. Two h later, pups were reanesthetized and probes were inserted. Probes were perfused with filtered modified Ringer's solution (NaCl 147 mM, KCl 4 mM, CaCl₂ 3.4 mM, pH 7.2) at 1.5 μ L/min using a microinfusion pump (CMA 100, BAS). Sampling was initiated after a 2-h recovery period, and 10 sequential 20-min fractions were collected (on ice).

After three baseline samples were collected, animals were exposed to 8% oxygen/92% nitrogen for 2 h; then, for the final sample, animals were returned to room air. Surface body temperature was monitored and was maintained at about 35.5°C. Animals were killed by decapitation at the end of sampling, and normoglycemia was verified (by Dextrostix \geq 80) of blood welled at the severed neck vessels. The microdialysis protocol was approved by the University of Michigan Committee on Use and Care of Animals.

Samples were analyzed by HPLC with electrochemical detection, and precolumn derivitization (modified from Ref. 21). Each sample (30 μ L) was mixed with 22.5 μ L of a solution containing 27 mg *o*-phthalaldehyde in 2 mL of 100 mM borax with 20 μ L mercaptoethanol, pH 9.5, for 30 s. The internal standard homoserine (22.5 μ L) was included. Samples (70 μ L) were injected onto a C18 reverse phase column (5 μ m ODS); the mobile phase was a 0.1 M NaH₂PO₄, pH 6.5 buffer with 32% methanol. A glassy carbon electrode connected to an electrochemical detector (Bioanalytical Systems) was used (sensitivity 20 nA, electrode potential 0.7 V *versus* Ag/AgCl to optimize the number of quantifiable peaks and minimize baseline noise). Values were quantified from peak heights, compared with peak heights of external standards assayed concurrently, and normalized using the internal standard homoserine. Values were also corrected for the recovery efficiency for each compound (recovery range 10–20%).

Efflux values were expressed both in pmol/min and as a

percentage of baseline; means (\pm SEM) for each sampling period were calculated. Values were also grouped for analysis into four time periods: 1) baseline, 2) 1st h of hypoxia, 3) 2nd h of hypoxia, and 4) recovery. Because variances were not equal, nonparametric tests were used for statistical analysis with a microcomputer-based statistical package (Statview).

RESULTS

Glutamate, aspartate, glutamine, glycine, asparagine, taurine, serine, and alanine were detected in hippocampal dialysate samples, as were several unidentified peaks; γ -aminobutyric acid was not consistently detected (despite an assay threshold of 2–3 pmol/20 μ L sample). There was considerable inter- and intrainimal variability in efflux values. Baseline efflux values were in a range similar to those obtained in striatum, using the same methods (Table 1) (20); hippocampal glutamate, aspartate, and taurine efflux values were higher than corresponding striatal values. Of interest, baseline hippocampal glutamate efflux was higher than peak values detected in striatum during hypoxia-ischemia (17 *versus* 7 pmol/min); efflux values appear to estimate relative changes in ECF amino acid content more accurately than their absolute synaptic concentrations.

Preliminary analysis indicated that glutamate efflux peaked transiently and variably in the 2nd h of hypoxia. Because previous studies in this model suggested that the time threshold for onset of irreversible injury occurs during the 2nd h of hypoxic exposure (22), values from the 1st h of hypoxia (fractions 4–6) and from the 2nd h of hypoxia (fractions 7–9) were grouped for analysis.

Figure 2 compares the widely divergent efflux patterns of the eight amino acids analyzed over these four time periods. Glutamate efflux increased during hypoxic exposure; in the 2nd h of hypoxia, mean values were three times as high as at baseline (51 \pm 9 *versus* 17 \pm 3 pmol/min, p = 0.002, Kruskal Wallis test comparing all values in each time period), and values declined toward baseline in the final recovery period in room air. In contrast, glutamine efflux declined progressively to 60% of baseline in the final sample (311 to 182 pmol/min, p = 0.0005, Kruskal Wallis test). Hypoxic exposure stimulated a sustained increase in alanine efflux (from 31 pmol/min at baseline to 63 pmol/min in the 1st h of hypoxia). No significant changes in efflux of the five other amino acids analyzed were detected.

To facilitate more detailed comparison of sequential changes in glutamate and glutamine efflux, for each animal values were normalized and expressed as a percentage of baseline (averaged from first three fractions) (Fig. 3). Mean glutamate efflux peaked early in the 2nd h of hypoxia, whereas glutamine efflux declined progressively.

Table 1. Comparison of baseline hippocampal and striatal amino acid efflux in PND 7 rats*

Amino acid	Efflux values (pmol/min)	
	Hippocampus	Striatum
Glutamate	17.3 \pm 3.1	2.1 \pm 0.5
Aspartate	6.2 \pm 1.2	1.3 \pm 0.5
Taurine	592 \pm 125	13.3 \pm 0.2
Asparagine	2.6 \pm 0.8	17.5 \pm 3.5
Glutamine	311 \pm 20	384 \pm 47
Serine	45.4 \pm 3.4	75.2 \pm 7.8
Alanine	30.9 \pm 2.5	40.2 \pm 4.5
Glycine	44.1 \pm 5.2	35.6 \pm 2.4

* Results were obtained from *in vivo* microdialysis experiments in PND 7 rats (see Materials and Methods). Baseline hippocampal efflux values (\pm SEM) were averaged from the first three dialysis fractions collected in each animal (n = 6). Striatal values, obtained using the same methods, were reported previously (20).

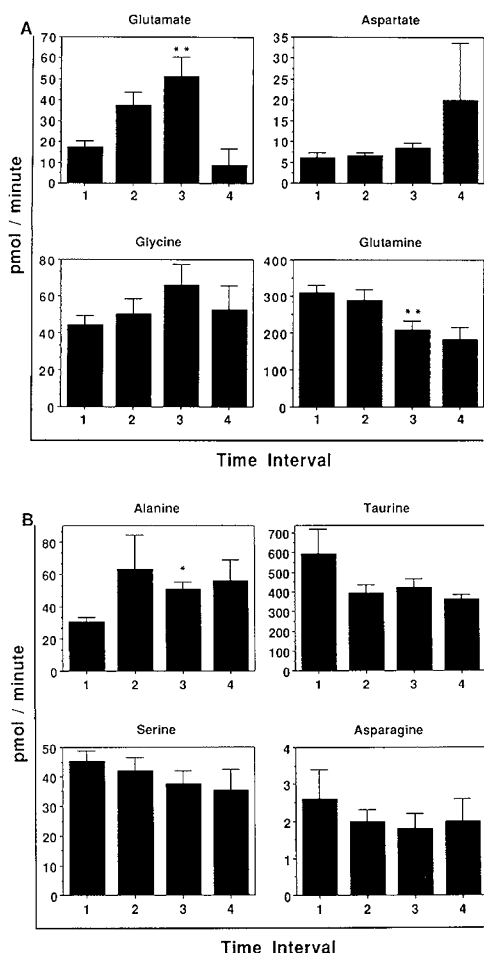


Fig. 2. *A* and *B*. These graphs compare efflux patterns of eight amino acids detected in hippocampal dialysates of PND 7 rats during hypoxia-ischemia. Microdialysis probes were inserted into the right hippocampus of PND 7 rats that underwent right carotid ligation, followed by exposure to 8% oxygen for 2 h ($n = 6$, see text for methods). Beginning 2 h after probe insertion, 10 sequential 20-min dialysis samples were collected from each animal. Samples were grouped for analysis into four time intervals: period 1, baseline, fractions 1–3; Period 2, h 1 of hypoxia, fractions 4–6; period 3, h 2 of hypoxia, fractions 7–9; and period 4, recovery, fraction 10. Amino acid content of dialysates was quantitated by an HPLC assay. Efflux values are means \pm SEM of all measurements in each time interval. **, $p < 0.005$; *, $p < 0.02$, comparison of all values within each time interval with Kruskal Wallis test.

DISCUSSION

Our results demonstrate that in this experimental model of perinatal stroke, which results in incomplete, reversible forebrain ischemia, hippocampal glutamate efflux rose markedly during the time period coinciding with the threshold for onset of irreversible injury.

Each efflux value reflects the cumulative influence of many neurons and glia over a prolonged sampling period relative to synaptic events (13, 14), and measurable changes in efflux indicate marked and sustained alterations in ECF glutamate accumulation. Yet, the extent of stimulation of glutamate efflux was considerably less than values reported in adult brain (3-fold compared with a >8 -fold rise in an adult stroke model) (3). This difference may reflect lower concentrations of synaptic glutamate than in adult brain, stimulation of release (and/or suppression of reuptake) in a smaller proportion of neurons, or a greater dilutional effect by the relatively larger ECF volume in the immature brain. Differences in the pathophysiology of ischemia induction (*e.g.* incomplete *versus* complete, focal *versus* global),

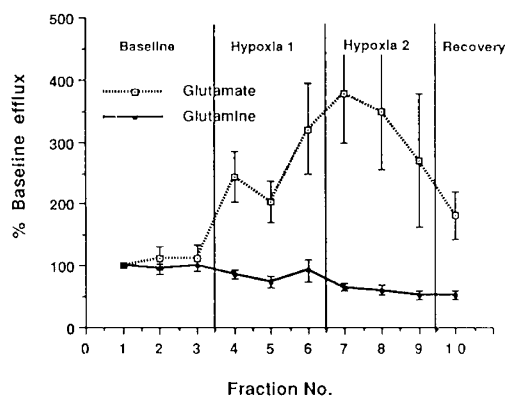


Fig. 3. this graph compares sequential hippocampal glutamate and glutamine efflux values, expressed as a percentage of baseline, in six animals that underwent right carotid artery ligation followed by 2 h of 8% oxygen exposure (see text for details of methods). Baseline values were defined as the averages from the first three fractions collected in each animal.

independent age-related effects, may also have contributed to the less-pronounced rise (8).

At least two major factors contributed to the wide range of efflux values observed. The extent of injury elicited by this lesioning method varies considerably; prominent lesions (with $>10\%$ reduction in ipsilateral hemisphere mass) evolve in about 70% of animals (9), and the severity of injury in individual animals undergoing microdialysis could not be determined. There were also unavoidable differences in anatomical placement of the probe within distinct functional regions of the hippocampus, in which both baseline and stimulated release might vary. In addition, small fluctuations in brain temperature (which we were unable to control) may have influenced neurotransmitter release (23).

In contrast with glutamate, glutamine efflux declined gradually; the diverging trends for the two amino acids suggest that systemic derangements such as dehydration were unlikely to account for the increase in glutamate efflux in the 2nd h of hypoxia. Inhibition of the glial enzyme glutamine synthetase, which converts glutamate to glutamine, has been reported in adult brain during reperfusion after cerebral ischemia; suppression of activity of this enzyme could result in both a reduction in glutamine efflux and synaptic glutamate accumulation (24).

In adult brain, aspartate efflux generally increases in parallel with changes in glutamate efflux during ischemia (3, 4); however, we found no consistent increase in hippocampal aspartate efflux. Similarly, in our previous microdialysis studies in PND 7 rats, neither hypoglycemia (20) nor hypoxia-ischemia (8) stimulated striatal aspartate efflux. This difference likely reflects lower synaptic concentrations of aspartate in immature brain; whether this developmental difference in EAA metabolism has pathophysiological significance as to mechanisms of neuronal injury is unknown. Awareness of the role of endogenous glycine in potentiating glutamate activation of N-methyl-D-aspartate receptors has prompted interest in analysis of glycine release (25); although no significant change in glycine efflux was detected, our data revealed a trend toward increased glycine efflux in the 2nd h of hypoxia.

Release of the endogenous potential neuromodulator taurine may increase in response to glutamatergic stimulation (26). In contrast with findings in hypoxic-ischemic striatum in which there was a close temporal correlation between glutamate and taurine efflux peaks (8), hippocampal taurine efflux did not rise. There may be regional functional differences in taurine metabolism; alternatively, because baseline hippocampal taurine efflux was much higher than in striatum, it is also possible that biologically important synaptic increases were not identifiable.

Baseline hippocampal γ -aminobutyric acid efflux was often

undetectable, and unlike in adult brain (3, 4), no increases were detected with hypoxia-ischemia; immaturity of γ -aminobutyric acid innervation may also be a contributing factor to the pathophysiological differences in the evolution of ischemic neuronal injury between immature and adult brain.

The observed increases in alanine efflux likely reflect increased glycolysis during hypoxia. Previously, in striatum, we found that both exposure to moderate hypoxia, which did not result in tissue damage, and hypoxia-ischemia stimulated alanine efflux; these findings suggested that increased alanine efflux was not an important contributing factor to the evolution of injury (8).

Our data do provide direct support of the hypothesis that extracellular glutamate accumulation may contribute to the pathogenesis of hypoxic-ischemic hippocampal injury in this model of perinatal stroke. It is also important to consider that recent studies in adult brain have demonstrated that glutamate efflux also rises during ischemia in brain regions that are relatively resistant to ischemic injury (27). Other factors, e.g. intrinsic variation in cellular susceptibility to excitotoxic injury and regional activity of endogenous neuromodulatory compounds, are likely to play critical roles in determining the extent of ischemic neuronal damage.

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