Effects of Neuroglobin Overexpression on Acute Brain Injury and Long-Term Outcomes After Focal Cerebral Ischemia

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Background and Purpose—Emerging data suggest that neuroglobin (Ngb) may protect against hypoxic/ischemic neuronal insults. However, the underlying mechanisms in vivo and implications for long-term outcomes are still not well understood.

Methods—Using our newly created Ngb overexpressing transgenic (Ngb-Tg) mice, we measured brain infarction on day 1 and day 14 after transient focal cerebral ischemia and performed neurobehavioral assessments in sensorimotor deficits on days 1, 3, 7, and 14. To test the hypothesis that Ngb may play a role in reducing oxidative stress after stroke, intracellular malondialdehyde levels were measured and compared in Ngb-Tg and wild-type mice.

Results—Increased Ngb mRNA and protein levels were identified in Ngb-Tg brains. Malondialdehyde levels in ischemic hemispheres of Ngb-Tg were significantly reduced compared with wild-type controls at 8 hours and 22 hours after transient focal cerebral ischemia. Compared with wild-type controls, brain infarction volumes 1 day and 14 days after transient focal cerebral ischemia were significantly reduced in Ngb-Tg mice. However, there were no significant improvements in sensorimotor deficits for up to 14 days after stroke in Ngb-Tg mice compared with wild-type controls.

Conclusions—Ngb reduces tissue infarction and markers of oxidative stress after stroke. Tissue protection by overexpressing Ngb can be sustained for up to 2 weeks. (Stroke. 2008;39:1869-1874.)

Key Words: neuroglobin ■ neuroprotection ■ oxidative stress ■ stroke

Neuroglobin (Ngb) is a recently discovered tissue globin with a high affinity for oxygen that is widely expressed in neurons of vertebrate central and peripheral nervous systems, retina, and endocrine tissues.1–5 It was recently shown that Ngb was elevated in both transcript and protein levels in cultured primary cortical neurons during the acute phases of hypoxia.6 An age-related decline in Ngb expression in rat brain was found and suggested that loss of this protein may have a role in increasing susceptibility to age-related neurological disorders.7

As a newly discovered member of the globin family, Ngb has been considered as the equivalent of brain or nerve tissue hemoglobin.8 The distribution is indicative of a function of Ngb in metabolically active, oxygen-consuming cell types. In general, tissue globins mediate multiple cellular and molecular responses to hypoxic/ischemic insults. For example, myoglobin in cardiomyocytes and oxidative skeletal myofibers helps facilitate oxygen transport, maintain nitric oxide homoeostasis, and scavenge reactive oxygen species.9–11 It is possible that Ngb has similar actions in brain. Greenberg and colleagues showed that overexpression of Ngb protects against hypoxic neuron injury in cell culture and reduces acute 1-day ischemic brain damage in vivo.6,12 However, the effects of Ngb overexpression on long-term neurological outcomes have not been validated and underlying neuroprotective mechanisms also remain unknown. In this study, we used our newly created Ngb-overexpressing transgenic mouse to examine neuroprotection of Ngb in both acute and prolonged times after focal cerebral ischemia and tested the hypothesis that Ngb promotes neuron survival in part by reducing oxidative stress.

Materials and Methods

All animal experiments were performed following protocols approved by the Massachusetts General Hospital Institutional Animal Care and Use Committee in compliance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals.
Generation of Neuroglobin-Overexpressing Transgenic Mice

To test our hypotheses, we generated Ngb transgenic (Ngb-Tg) mice following standard methods. In brief, the full-length cDNA sequence of mouse Ngb was cloned following bioinformatic analysis and the rapid amplification of cDNA ends technique. The full-length of Ngb cDNA (GenBank Accession No NM 022414) was used to amplify the cDNA sequence. To create a fusion of Ngb gene and the N-terminal hemagglutinin epitope tag, the mouse Ngb gene was subcloned into the Sal I and Kpn I sites of the MCS of pCMV-myc (Clontech) plasmid in frame with the hemagglutinin coding sequence. The subcloned eukaryotic expression plasmid, pCMV-HA-mouse-Ngb, capable of encoding mouse Ngb tagged with hemagglutinin, respectively, was used to release the Ngb gene. The full length of the pCMV-Ngb transgene containing the Ngb gene and polyadenylation signals in the 3' portion of the construct was released from the vector with SpI I and Mfe I. The purified plasmid was obtained by sucrose density gradient centrifugation and sequenced to determine the orientation of the subcloned Ngb product. The purified DNA fragment was used for generating Ngb-Tg mice in the Transgenic Core Facility at Massachusetts General Hospital. Pronuclear microinjection was performed by using standard techniques. Tg mice were initially created with B6C3F1 background and then backcrossed with C57BL/6J mice. All animals used in this study were 6 to 9 generations; they were Ngb-Tg mice and wild-type (WT) littermates.

Genotyping of Neuroglobin Transgene by Polymerase Chain Reaction

To identify the Ngb transgene, mouse tail DNA was used for polymerase chain reaction (PCR) analysis with primers flanked by the cytomegalovirus (CMV) and Ngb sequence (5' tcaagtctccac-cgccattgacg 3' and 5'-tggtcactgcagcatcaatca-3'). A 940-bp band of PCR production was the indicator for the presence of the mouse Ngb transgene.

Quantitative Real-Time Polymerase Chain Reaction Analysis

mRNA levels of Ngb were measured by real-time reverse transcription–PCR analysis following a standard method with minor modification. In brief, total brain tissue RNA was prepared for real-time PCR using the RNeasy Lipid Tissue Mini Kit (Qiagen) according to the manufacturer’s instructions. Reverse transcriptase was performed using Superscript III Reverse Transcriptase (Invitrogen) to obtain cDNA. The primers for mouse Ngb were as follows: 5'- tcaagtctccacgctcag 3' and 5'-tggtcactgcagcatcaatca-3'. Real-time PCR was performed on an ABI prism 7000 sequence detection systems (Applied Biosystems). Data were analyzed according to the comparative threshold cycle method with glyceraldehydes-3-phosphate dehydrogenase expression for sample normalization. Melting curves for each PCR reaction were generated to ensure the purity of the amplification products.

Western Blot and Immunohistochemistry

To examine Ngb protein expression levels, customized polyclonal anti-Ngb antibody (GenScript Corporation) was produced by immunizing rabbits with 2 synthetic peptides (KLH coupled, FQYNGQFSPPEDC and IRQSWRVVVSRSINGLE) corresponding to mouse Ngb. The antibody was purified by protein A and peptide affinity chromatography. Western blot for Ngb was performed following the protocol as previously described. Relative Ngb protein expression levels were assessed by quantification of optical density of Ngb protein bands with National Institutes of Health Image software. For immunohistochemistry, mice were transcardially perfused with ice-cold phosphate-buffered saline (pH 7.4) and 4% paraformaldehyde. Brains were fixed in 4% paraformaldehyde. Brains were fixed in 4% paraformaldehyde. Image software. For immunohistochemistry, mice were transcardi-
ipsilateral and contralateral limbs. Mice were suspended by its forelimbs on a wire stretched between 2 posts 60 cm above a foam pillow. The time (in seconds) until the mouse fell was recorded, and the posture of the 4 limbs and tail and whether the mouse could walk were recorded as different scores (0 to 5). A score of zero was assigned if the mouse fell immediately and 60 seconds was the time-out period. Three trials were performed for each testing day.

**Statistical Analysis**

For parametric and continuous variable measurements, we used analysis of variance followed by Tukey-Kramer post hoc tests. For measurements taken over time, we used repeated-measures analysis of variance and Fisher partial least-square difference test. For nonparametric ordinal data (eg, functional outcomes), we used nonparametric Kruskal-Wallis followed by post hoc Mann-Whitney tests. Overall, \( P < 0.05 \) was considered significant.

**Results**

**Production and Characterization of Neuroglobin-Overexpressing Transgenic Mice**

PCR analysis with primers flanking CMV and Ngb sequence detected a 940-bp band of PCR production indicates the presence of mouse Ngb transgene. DNA markers (Marker) appeared on the first lane; pCMV-Ngb plasmid served as a positive control (P). Brain RNA was extracted for measuring Ngb mRNA levels by quantitative real-time PCR. There was an approximately 2.6-fold increases of Ngb mRNA levels in Ngb-Tg mouse brains compared with WT and WT littermates (WT-L); similar Ngb mRNA levels were detected in WT (C57BL/6) and WT-L mouse brains. All animals used were 22 to 26 males between 10 and 12 weeks of age. Mean ± SEM, \( n = 6 \) per group, \( * P < 0.05 \). Top panel is a representative Western blot showing the increase of Ngb protein in Ngb-Tg mouse brains (Tg) compared with WT controls. Actin served as equal loading controls. Lower panel shows relative Ngb protein expression levels quantified by optical density of Ngb protein bands. Mean ± SEM, \( n = 4 \) per group, \( * P < 0.05 \). In cerebral cortex, immunohistochemistry showed Ngb immunoreactivity was mainly colocalized with neuron marker neuronal nuclei (D, A–C), not colocalized with astrocyte marker glial fibrillary acidic protein (D, D–F) in WT mice. By comparison, Ngb immunoreactivity was enhanced in neurons (D, G–I), some colocalized with astrocyte marker glial fibrillary acidic protein as pointed out by arrows (D, J–L) in Ngb-Tg mice. Bar: 50 µmol/L.
Table 1. Cerebral Blood Flow Measurements

<table>
<thead>
<tr>
<th>Measurements</th>
<th>WT</th>
<th>Ngb-Tg</th>
</tr>
</thead>
<tbody>
<tr>
<td>Hydrogen clearance</td>
<td>83.3±3.2</td>
<td>88.5±8.9</td>
</tr>
<tr>
<td>Resting cerebral blood flow, mL/100 g/min</td>
<td></td>
<td></td>
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<tr>
<td>Laser Doppler fl owmetry</td>
<td></td>
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<tr>
<td>Ischemia</td>
<td></td>
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<tr>
<td>Reperfusion</td>
<td></td>
<td></td>
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<tr>
<td>Cerebral blood flow, % of preischemia baseline</td>
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</tr>
</tbody>
</table>
| Hydrogen clearance was used to measure resting cerebral blood flow; platinum wire was positioned under cortex caudate putamen nuclei. Laser Doppler flowmetry was used to measure cerebral blood flow during ischemia and reperfusion in Ngb-Tg and WT mice. Results are expressed as mean±SEM. N=5 per group for hydrogen clearance and n=10 per group for laser Doppler flowmetry measurement, respectively. All animals used were 23- to 27-g male mice. In both measurements, there were no statistically significant differences between WT and Ngb-Tg mice.

Measurements of Cerebral Blood Flow

Hydrogen clearance was used to compare resting cerebral blood flow, and laser Doppler flowmetry was used to measure reduction of perfusion levels during ischemia and reperfusion. In both measurements, there were no statistically significant differences between WT and Ngb-Tg mice (Table 1).

Reduction of Acute Ischemic Infarction in Neuroglobin Transgenic Mouse Brains

To test the effect of Ngb overexpression in acute ischemic damage, brain infarction was examined at 24 hours after a transient 2-hour focal cerebral ischemia by standard 2,3,5-triphenyltetrazolium hydrochloride staining. Total infarct volumes were significantly smaller in Ngb-Tg mice compared with WT controls with the main reductions occurring mostly in cortical regions (Figure 2A). As expected, there was also some hemispheric swelling in ischemic brain, but there was no detectable difference between WT versus Ngb-Tg brains: 9.54±2.35% for WT and 7.44±3.1% (mean±SEM, n=8) for Ngb-Tg mice, respectively. Physiological parameters, including blood pressure (mm Hg), arterial pH, pCO2, and pO2 (mm Hg), were monitored in randomly selected animals; there was no significant difference between the 2 animal groups (Table 2).

Reduction of Late Ischemic Infarction in Neuroglobin Transgenic Mouse Brains

Because ischemic lesion evolution may continue beyond 24 hours after stroke, hematoxylin & eosin staining analysis was used to measure brain infarction at 14 days after a 1-hour period of transient focal cerebral ischemia. Infarction volumes in Ngb-Tg mice remained significantly smaller compared with WT controls even at this delayed time poststroke (Figure 2B).

Reduction of Malondialdehyde Production in Ngb-Tg Mouse Brains After Stroke

MDA levels were measured as a surrogate biomarker of reactive oxygen species generation after stroke. In nonischemic control brains, MDA levels were not significantly different between Tg and WT mice indicating similar baseline conditions. However,

Table 2. Physiological Parameter Measurements

<table>
<thead>
<tr>
<th>Parameters</th>
<th>pO2</th>
<th>pCO2</th>
<th>pH</th>
<th>Blood Pressure</th>
</tr>
</thead>
<tbody>
<tr>
<td>Preischemia</td>
<td>126.2±9.7</td>
<td>31.3±1.4</td>
<td>7.38±0.05</td>
<td>88.2±4.4</td>
</tr>
<tr>
<td>Postischemia</td>
<td>95.4±10.5</td>
<td>46.9±1.7</td>
<td>7.34±0.08</td>
<td>90.2±5.1</td>
</tr>
<tr>
<td>Ngb-Tg</td>
<td>Preischemia</td>
<td>123.1±9.2</td>
<td>30.5±1.5</td>
<td>7.37±0.06 89.7±4.5</td>
</tr>
<tr>
<td></td>
<td>Postischemia</td>
<td>98.3±8.9</td>
<td>47.3±2.4</td>
<td>7.35±0.07 86.8±5.7</td>
</tr>
</tbody>
</table>

Physiological parameters, including arterial pO2 (mm Hg), pCO2, pH, and blood pressure (mm Hg), were monitored before focal cerebral ischemia (Preischemia) and 1 hour after ischemia (Postischemia) in randomly selected WT and Ngb-Tg mice. Data are mean±SD, n=4 per group.
MDA production was significantly reduced in Ngb-Tg mice compared with WT controls at both early 8-hour and late 22-hour time points after transient focal cerebral ischemia (Figure 3).

**Sensorimotor Function Assessments**

Four assessments, including neurological score, Rotorod test, hanging wire, and foot fault test, were performed at day 1, 3, 7, and 14 days after ischemic onset. Two hours before focal ischemic injury, mice were assessed to obtain preinjury baselines. Body weight loss was also measured on each test day. Significant deficits were observed in all tests from day 1 to day 7 after ischemia; the deficits reached maximal levels at day 7 for the foot fault test, and day 3 to day 7 after body weight losses. All scores recovered close to preinjury baselines by day 14 after onset of ischemia. No statistically significant differences were detectable between WT and Ngb-Tg mice in all assessments during the 2-week stroke recovery period (Table 3).

### Discussion

Rapidly emerging data suggest that Ngb may function as an important neuroprotective molecule. However, the mechanisms underlying its actions remain to be fully elucidated. In the present study, we created a novel strain of Ngb-overexpressing transgenic mice and used this model system to show that Ngb is neuroprotective against cerebral ischemia in vivo. Acute focal infarction at 24 hours after transient 2-hour focal cerebral ischemia was significantly reduced approximately 30%, consistent with a most recent report from the Greenberg group. Importantly, our present study documented that tissue neuroprotection in the Ngb-Tg brains was sustained for up to 2 weeks. In terms of mechanisms, the reduction of MDA production in Ngb-Tg mouse brains suggests that Ngb promotes neuron survival in part by reducing oxidative stress after focal cerebral ischemia.

The functional role of Ngb in normal brain is not well understood. It has been suggested that Ngb may function as oxygen sensors associated with heme proteins and may respond to hypoxia by altering the rate of formation and release of reactive oxygen species that activate transcription factors. Furthermore, Ngb may function as a scavenger of nitric oxide or a regulator of reactive oxygen species. By altering these free radicals, Ngb may protect against oxidative stress in stroke. The present study demonstrated Ngb’s neuroprotection against ischemia might occur through decreasing oxidative stress. In addition to its use to dissect pathophysiology, our newly generated Ngb-Tg mouse may also provide a useful tool to elucidate the functional roles and mechanisms of Ngb under physiological conditions.

There are a few caveats in this study. First, our Ngb-Tg mouse is not tissue- or cell-specific. DNA fragments containing mammalian cell promoter CMV and Ngb full-length cDNA were extracted for generating Ngb transgenic mice. Further studies using conditional or neuron-specific Ngb-Tg mice may be required. A second related issue is that as a constitutive mutant, adaptive compensations may have occurred. Although we saw no differences in resting (hydrogen clearance) or ischemic (laser Doppler flowmetry) cerebral blood flow, other spatially resolved techniques such as autoradiography or anatomic mapping of capillary density may be required to fully address these blood flow questions. A third issue in the present study is that we failed to detect significant differences in sensorimotor function recov-

### Table 3. Sensorimotor Function Assessments

<table>
<thead>
<tr>
<th>Tests</th>
<th>Group</th>
<th>Day 0</th>
<th>Day 1</th>
<th>Day 3</th>
<th>Day 7</th>
<th>Day 14</th>
</tr>
</thead>
<tbody>
<tr>
<td>Neurological score (25th to 75th percentiles)</td>
<td>WT</td>
<td>0</td>
<td>1 (0.25–1)</td>
<td>1 (1–2)</td>
<td>1 (1–2)</td>
<td>1 (0.25–1)</td>
</tr>
<tr>
<td></td>
<td>Ngb-Tg</td>
<td>0</td>
<td>1 (1–2)</td>
<td>2 (1.5–2)</td>
<td>1 (0–1)</td>
<td>1 (0–1)</td>
</tr>
<tr>
<td>Rotorod test, % of day 0</td>
<td>WT</td>
<td>100</td>
<td>70.5±10.8</td>
<td>78.5±10.6</td>
<td>90.5±9.7</td>
<td>104.3±3.3</td>
</tr>
<tr>
<td></td>
<td>Ngb-Tg</td>
<td>100</td>
<td>68.7±12.8</td>
<td>78.1±10.9</td>
<td>84.5±17.4</td>
<td>110.7±2.5</td>
</tr>
<tr>
<td>Hanging wire median (25th to 75th percentiles)</td>
<td>WT</td>
<td>5 (4.75–5)</td>
<td>2.5 (1–4)</td>
<td>3.5 (2–4.25)</td>
<td>4 (4–5)</td>
<td>4.5 (4–5)</td>
</tr>
<tr>
<td></td>
<td>Ngb-Tg</td>
<td>5 (4.5–5)</td>
<td>3 (1.5–3)</td>
<td>4 (2.5–4)</td>
<td>4 (3.5–4)</td>
<td>4 (4–5)</td>
</tr>
<tr>
<td>Foot fault test median (25th to 75th percentiles)</td>
<td>WT</td>
<td>0 (0–1)</td>
<td>3 (0–5.75)</td>
<td>4 (0–5.25)</td>
<td>5 (4–5)</td>
<td>2 (1–2)</td>
</tr>
<tr>
<td></td>
<td>Ngb-Tg</td>
<td>1 (0–1.5)</td>
<td>2 (1–5)</td>
<td>4 (2–6)</td>
<td>4 (3.5–4.5)</td>
<td>1 (1–1.5)</td>
</tr>
<tr>
<td>Body weight, % of day 0</td>
<td>WT</td>
<td>100</td>
<td>91.7±0.57</td>
<td>82.1±1.21</td>
<td>81.8±3.04</td>
<td>91.3±1.54</td>
</tr>
<tr>
<td></td>
<td>Ngb-Tg</td>
<td>100</td>
<td>91.5±0.73</td>
<td>80.1±1.58</td>
<td>81.2±3.45</td>
<td>91.9±1.55</td>
</tr>
</tbody>
</table>

Sensorimotor deficits and body weight loss were assessed before (day 0) and at days 1, 3, 7, and 14 after transient (1-hour) focal cerebral ischemia. Results are expressed as median values and 25th to 75th percentiles for assessments of neurological score, hanging wire, and foot fault tests; and mean±SEM for assessments of Rotorod test and body weight loss. N=6 for Ngb-Tg and 7 for WT per group; 23- to 27-g male mice. In all assessments, there were no statistical significant differences between WT and Ngb-Tg groups.
ery in our Ngb-Tg mice. In part this is because 60 minutes middle cerebral artery occlusion might not be severe enough to cause long-term sensorimotor deficits because these smaller lesions are mainly localized in subcortex areas. Alternatively, we may have simply lacked power and larger numbers of test animals would have helped. Further investigation is warranted to carefully define the role of Ngb in both long-term sensorimotor and cognitive deficits after stroke. The final caveat is related to the inability to unequivocally separate reactive oxygen species effects versus mitochondrial effects of Ngb. There are multiple and probably unequivocally separate reactive oxygen species effects versus mechanisms. We acknowledge that it will likely be impossible to carefully define the role of Ngb in both long-term sensorimotor and cognitive deficits after stroke. However, for in vivo stroke models, it is difficult to unequivocally prove causality because reduced tissue damage may secondarily contribute to decreased oxidative stress and/or mitochondrial dysfunction.

In conclusion, this study documented the reduction of brain infarction in the Ngb-Tg brains was sustained for up to 2 weeks. This unique brain globin protects against ischemic injury by ameliorating oxidative stress. Further studies to test both sensorimotor and cognitive deficits are warranted in terms of fully evaluating the effect of Ngb in stroke long-term recovery.

Acknowledgments

We thank Dr David Greenberg for very helpful discussion and Drs Shuzhen Guo, Sun-Ryung Lee, and Changhong Xing for their excellent technical assistance.

Sources of Funding

This work was supported in part by National Institutes of Health grants R01-NS049476 (to X.W.), R01-NS37074, R01-NS48422, and P50-NS10828 (to E.H.L.), and a Scientist Development Grant, 050-NS10828 (to E.H.L.), and a Scientist Development Grant, grants R01-NS049476 (to X.W.), R01-NS37074, R01-NS48422, and R01-NS48422, from the American Heart Association (to X.W.).

Disclosures

None.

References


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*Stroke*. 2008;39:1869-1874; originally published online April 10, 2008; doi: 10.1161/STROKEAHA.107.506022

*Stroke* is published by the American Heart Association, 7272 Greenville Avenue, Dallas, TX 75231
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Print ISSN: 0039-2499. Online ISSN: 1524-4628

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