

HIF-1 α -Deficient Mice Have Increased Brain Injury after Neonatal Hypoxia-Ischemia

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Key Words

Brain injury · Transcription factor · Hypoxia-inducible factor 1 α · Neonatal hypoxia-ischemia

Abstract

Evidence suggests that the activation of the transcription factor hypoxia-inducible factor 1 α (HIF-1 α) may promote cell survival in hypoxic or ischemic brain. To help understand the role of HIF-1 α in neonatal hypoxic-ischemic brain injury, mice with conditional neuron-specific inactivation of HIF-1 α underwent hypoxia-ischemia (HI). Mice heterozygous for Cre recombinase under the control of the calcium/calmodulin-dependent kinase II promoter were bred with homozygous 'floxed' HIF-1 α transgenic mice. The resulting litters produced mice with a forebrain predominant neuronal deletion of HIF-1 α (HIF-1 $\alpha^{\Delta/\Delta}$), as well as littermates without the deletion. In order to verify reduction of HIF-1 α at postnatal day 7, HIF-1 $\alpha^{\Delta/\Delta}$ and wild-type mice were exposed to a hypoxic stimulus (8% oxygen) or room air for 1 h, followed by immediate collection of brain cortices for determination of HIF-1 α expression. Results of Western blotting of mouse cortices exposed to hypoxia stimulus or room air confirmed that HIF-1 $\alpha^{\Delta/\Delta}$ cortex expressed a minimal amount of HIF-1 α protein compared to wild-type cortex with the same hypoxic stimulus. Subsequently, pups underwent the Vannucci

procedure of HI at postnatal day 7: unilateral ligation of the right common carotid artery followed by 30 min of hypoxia (8% oxygen). Immunofluorescent staining of brains 24 h after HI confirmed a relative lack of HIF-1 α in the HIF-1 $\alpha^{\Delta/\Delta}$ cortex compared to the wild type, and that HIF-1 α in the wild type is located in neurons. HIF-1 α expression was determined in mouse cortex 24 h after HI. Histological analysis for the degree of injury was performed 5 days after HI. HIF-1 α protein expression 24 h after HI showed a large increase of HIF-1 α in the hypoxic-ischemic cortex of the wild-type compared to the hypoxic only cortex. Histological analysis revealed that HI injury was increased in the neuronally deficient HIF-1 $\alpha^{\Delta/\Delta}$ mouse brain ($p < 0.05$) and was more severe in the cortex. Genetic reduction of neuronal HIF-1 α results in a worsening of injury after neonatal HI, with a region-specific role for HIF-1 α in the setting of neonatal brain injury.

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Introduction

Hypoxia-inducible factor 1 α (HIF-1 α) is a transcription factor that is essential for the activation of over 40 hypoxia-inducible genes, such as erythropoietin, glucose transporters, glycolytic enzymes, and vascular endothelial growth factor (VEGF) [1, 2]. Thus, HIF-1 α has a wide-

ranging role in the cellular response to hypoxia, and in some pathological circumstances, such as tumor growth, its role is increasingly well established [3, 4]. In the brain, however, where oxygen homeostasis is of critical importance, the role of HIF-1 α , whether physiologic or pathologic, is less well understood. Hypoxia induces many of the genes regulated by HIF-1 α in the brain [5, 6], and preconditioning with hypoxia may induce HIF-1 α and promote cell survival in the subsequently hypoxic or ischemic brain [7].

Increased expression of HIF-1 α -dependent genes, such as VEGF [8, 9], several glycolytic enzymes [10], glucose transporter 1 [10] and erythropoietin [11–14], is thought to be important in recovery from hypoxia-ischemia (HI) and stroke in the neonatal brain. In fact, evidence suggests that hypoxia preconditioning protection depends on gene expression induced by activation of HIF-1 α , since the functions of the HIF-1 α target genes are to increase blood flow, promote angiogenesis, stimulate glucose and lactate delivery to cells, enhance metabolism and promote growth and repair in neurons [4, 15–17]. An ischemic insult can also lead to induction of HIF-1 α and its target gene, VEGF, as has been reported following neonatal focal ischemia in the rat [18]. In addition, it has been shown in vitro that the neuroprotective effects of deferoxamine are mediated, at least in part, through induction and stabilization of HIF-1 α [19]. Subsequently, we showed that deferoxamine treatment after middle cerebral artery occlusion in neonatal rats is protective and prolongs the expression of HIF-1 α compared to saline-treated rats. The HIF-1 α target gene erythropoietin was also upregulated in the protected brains [9].

We explored the effect of neuron-specific reduction of HIF-1 α (HIF-1 $\alpha^{\Delta/\Delta}$) on neonatal hypoxic-ischemic brain injury. We employed a well-characterized model of neonatal HI in HIF-1 $\alpha^{\Delta/\Delta}$ compared to wild-type (Wt) littermates, and found increased cortical injury.

Materials and Methods

Animals

All animal research was approved by the University of California San Francisco Institutional Animal Care and Use Committee and was performed with the highest standards of care under the National Institutes of Health guidelines. Mice with conditional neuron-specific inactivation of HIF-1 α were generated using Cre/Lox technology. These mice have previously been well characterized [20, 21]. Briefly, the deletion was attained by breeding mice that have loxP-containing 'floxed' HIF-1 α alleles with mice expressing Cre recombinase under the control of the calcium/calmodulin-dependent kinase II promoter [22]. The mice

containing the Cre transgene used were of the 'R1ag#5' line [23]. The resulting litters produced mice with a forebrain predominant, neuron-specific deletion of HIF-1 α (HIF-1 $\alpha^{\Delta/\Delta}$), as well as littermates without the deletion. All mice negative by PCR for the Cre gene were considered 'wild-type (Wt)'. Genotyping was carried out by PCR on tail DNA samples using standard methods.

Hypoxic Stimulus Alone

At postnatal day 7 (P7), mice were placed in chambers maintained at 37°C and subjected to 1 h of 8% oxygen, or room air. Brain cortices were collected immediately after the hypoxia period, snap frozen and stored at –80°C for HIF-1 α protein determination by Western blot.

Hypoxia-Ischemia

At P7, mice underwent the Vannucci procedure of neonatal HI [24, 25]. Briefly, under isoflurane anesthesia, the right common carotid artery was permanently ligated. Following a 90-min recovery period, mice were exposed to hypoxia for 30 min (8% oxygen, balance nitrogen) while being maintained at 37°C. We have previously shown that the background mouse strain, C57Bl/6, while somewhat resistant to injury from HI, has high mortality [26]. By using 30 min of hypoxia, we had a minimal amount of mortality, with moderate injury. At either 24 h (for immunofluorescence) or 5 days after HI (for scoring of the degree of injury), mice were perfused with 4% paraformaldehyde in 0.1 M phosphate buffer, brains were removed, and processed histologically. The 5-day post-HI brains were analyzed for the degree of injury using a scoring system as previously described [26]. In addition, some mice were perfused with saline and a piece of contralateral cortex was obtained for Western blot to confirm the tail DNA PCR results. These brains were then immersion-fixed in paraformaldehyde prior to histological preparation as above.

Immunofluorescence

Flash-frozen mouse brains were cut on a cryostat and 12- μ m sections collected on glass slides. Brain sections were treated with 2 N HCl for 10 min at 37°C and then with 0.5 M boric acid (pH 8.4) for 10 min at room temperature. After a 2-min rinse with PBS, a blocking solution composed of PBS, 10% goat serum and 0.3% Triton X-100 was applied for 1 h. The primary antibodies against HIF-1 α (1:100, Santa Cruz Biotechnology) and NeuN (1:100, Millipore) in a similar blocking solution (2.5% goat serum) were applied overnight at 4°C. After washing, secondary antibodies (1:1,000, Alexa Fluor 488 and 568, Invitrogen) in a blocking solution were applied for 1 h at room temperature. After rinsing in PBS, DAPI staining was applied for 5 min and thereafter the sections were coverslipped with ProLong Gold (Invitrogen). Images were taken with a Nikon spectral confocal microscope at the Nikon Imaging Center, University of California San Francisco.

Western Blot Analysis

We first determined HIF-1 α expression with and without a hypoxic stimulus, in which P7 HIF-1 $\alpha^{\Delta/\Delta}$ (n = 8) and Wt (n = 6) mice were exposed to hypoxia (8% oxygen) or room air for 1 h. Brain cortices were collected immediately after the hypoxia period, snap frozen and stored at –80°C until use. Another group of mice underwent HI at P7 and cortices were collected, as above, 24 h later. Western blots were subsequently performed for HIF-1 α : in brief, cortices were homogenized in ice-cold lysis buffer

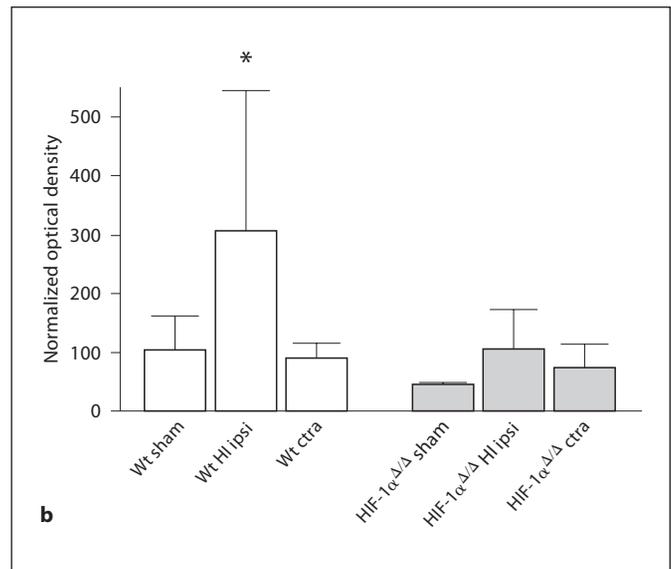
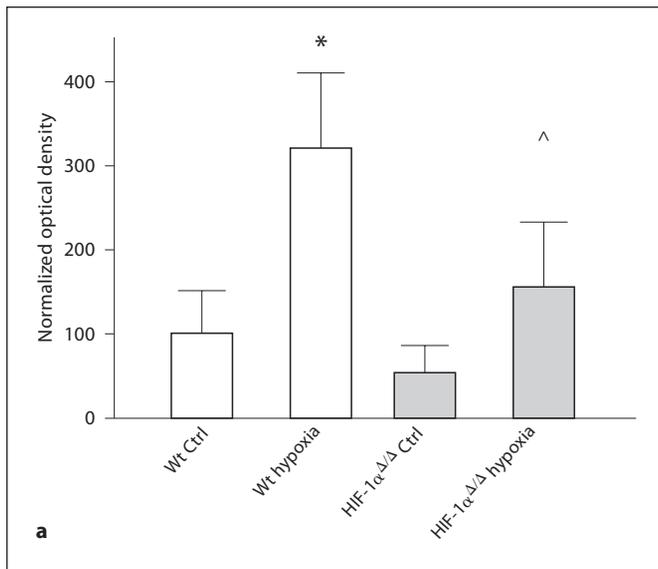


Fig. 1. HIF-1 α protein expression. **a** HIF-1 α protein expression in P7 mouse cortices, with or without 1 h of 8% oxygen exposure, by Western blot analysis. Optical densities are normalized to β -actin and values are expressed as mean \pm SD of optical densities normalized to Wt control (Ctrl). HIF-1 α protein expression is higher in hypoxic Wt (*) compared to all other groups: Wt naïve control ($p < 0.05$), HIF-1 $\alpha^{\Delta/\Delta}$ naïve control ($p < 0.01$), and HIF-1 $\alpha^{\Delta/\Delta}$ hypoxia ($p < 0.04$). In addition, HIF-1 $\alpha^{\Delta/\Delta}$ with hypoxia has higher HIF-1 α protein levels than HIF-1 $\alpha^{\Delta/\Delta}$ control ($^{\wedge} p < 0.05$). For Wt control, Wt hypoxia and HIF-1 $\alpha^{\Delta/\Delta}$, $n = 3$; for HIF-1 $\alpha^{\Delta/\Delta}$ hypoxia, $n = 5$. Analysis by ANOVA with the Bonferroni multiple comparison test. **b** HIF-1 α protein expression in mouse cortices 24 h after HI at P7. The Wt cortex ipsilateral (ipsi) to HI injury (* $n =$

5) has increased HIF-1 α protein expression compared to all other groups: Wt contralateral (ctr) to HI ($p < 0.01$, $n = 5$) Wt sham ($p < 0.01$, $n = 7$), HIF-1 $\alpha^{\Delta/\Delta}$ sham ($p < 0.01$), HIF-1 $\alpha^{\Delta/\Delta}$ ipsilateral to HI injury ($p < 0.01$) and HIF-1 $\alpha^{\Delta/\Delta}$ contralateral to HI injury ($p < 0.001$). There are no significant differences among the three HIF-1 $\alpha^{\Delta/\Delta}$ groups. However, there is a trend for higher HIF-1 α protein expression in the HIF-1 $\alpha^{\Delta/\Delta}$ cortex ipsilateral to HI injury ($n = 9$) compared to HIF-1 $\alpha^{\Delta/\Delta}$ sham ($p < 0.06$, $n = 4$) and contralateral to HI ($p = 0.10$, $n = 9$). Optical densities are normalized to β -actin and values are expressed as optical densities normalized to Wt sham (mean \pm SD). Analysis by ANOVA with the Bonferroni multiple comparison test.

containing sodium chloride (150 mM), Tris-HCl (50 mM, pH 8.0), NP-40 (1%), DOC (0.5%), SDS (0.1%), PMSF (0.5 mM), and the protease inhibitors aprotinin and leupeptin. The homogenates were centrifuged at 14,000 rpm for 30 min at 4°C, and the protein concentration of the supernatants determined by BCA (Pierce, Rockwell, Md., USA). Protein samples were diluted in SDS-PAGE sample buffer, boiled for 5 min and separated by SDS-PAGE. Proteins were transferred to polyvinylidene fluoride membranes, which were blocked in 5% non-fat dry milk in TBS, with 0.05% Tween 20 for 1 h at room temperature. After blocking, membranes were incubated with mouse anti-HIF-1 α monoclonal antibody (Santa Cruz Biotechnology, Santa Cruz, Calif., USA, 1:200) for 1 h at room temperature, then overnight at 4°C. Rabbit anti- β -actin polyclonal antibody (Sigma, St. Louis, Mo., USA, 1:200) was detected on the immunoblots to ensure equal protein loading. Following washes, the membranes were incubated with HRP-conjugated goat anti-mouse IgG (Santa Cruz, 1:3,000) in blocking solution for 1 h. After further washes, the signal of bound antibody was captured on X-ray film with enhanced chemiluminescence (Pierce, Rockford, Ill., USA). NIH Image was used to measure the densities of the protein signals after being scanned. Optical density of the protein bands was measured following

subtraction of the film background and protein levels were normalized to β -actin. All experiments were repeated at least three times to ensure reproducibility of results.

Statistical Analysis

For the histological scoring of brain injury, significance was determined by the Mann-Whitney test. For the Western blots, ANOVA with the Bonferroni test for multiple comparisons was used. Significance was set at $p < 0.05$. Analysis was done with Prism 4.0 (Graphpad Software Inc.).

Results

HIF-1 α Protein Expression

We confirmed that HIF-1 α is upregulated in the cortex immediately after a hypoxic insult. Specifically, HIF-1 α protein expression is increased in Wt cortex after a hypoxic stimulus compared to Wt naïve control ($p < 0.05$) and to HIF-1 $\alpha^{\Delta/\Delta}$ naïve control ($p < 0.01$). There is also an

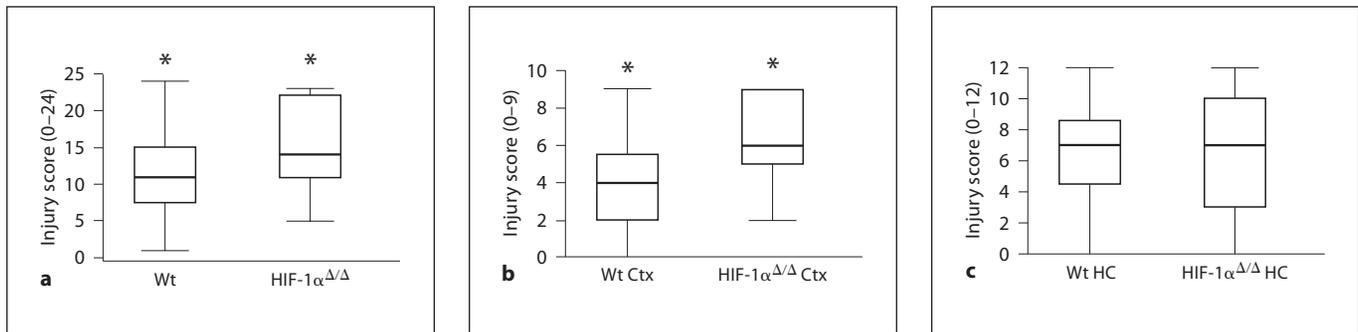


Fig. 2. HI brain injury 5 days after HI. **a** Total brain injury score (scale 0–24). HIF-1 $\alpha^{\Delta/\Delta}$ mice (median 14, $n = 11$) are more injured than Wt (median 11, $n = 27$) (* $p < 0.05$). **b** Cortical injury (scale 0–9). HIF-1 $\alpha^{\Delta/\Delta}$ cortices (Ctx) (median = 6) are more injured than Wt cortices (median = 4) (* $p < 0.01$). **c** Hippocampal injury (scale

0–12). There is no difference in hippocampal injury between groups (median = 7 for both, $p > 0.75$). The horizontal line within the box represents the median injury score for each group. Analysis by the Mann-Whitney test. HC = Hippocampus.

increase in protein expression in the HIF-1 $\alpha^{\Delta/\Delta}$ cortex with hypoxic stimulus compared to HIF-1 $\alpha^{\Delta/\Delta}$ naïve control ($p < 0.05$) (fig. 1a), indicating the presence of some cells that produce HIF-1 α .

Western blot analysis of brain cortices collected 24 h after HI (fig. 1b) shows that the Wt cortex ($n = 5$) has a level of HIF-1 α protein more than twice as high as the Wt sham cortex ($n = 7$; $p < 0.05$) and the Wt cortex contralateral to HI (hypoxic, $n = 5$; $p < 0.01$). The Wt HI cortex also had higher HIF-1 α protein levels than all HIF-1 $\alpha^{\Delta/\Delta}$ groups: sham ($p < 0.01$), ipsilateral to HI ($p < 0.01$) and contralateral ($p < 0.001$). In the HIF-1 $\alpha^{\Delta/\Delta}$ groups, there is a trend toward increased HIF-1 α levels in the HI cortex ($n = 9$) compared to sham ($n = 4$) ($p < 0.06$) and contralateral cortices ($p = 0.10$).

Histological Analysis for Degree of Injury

Histological analysis of brains 5 days after HI demonstrates an increase in injury in the HIF-1 $\alpha^{\Delta/\Delta}$ mouse brain compared to Wt brain (fig. 2a: median scores, Wt = 11, HIF-1 $\alpha^{\Delta/\Delta}$ = 14, $p < 0.05$). This injury is more severe in the cortex of the HIF-1 $\alpha^{\Delta/\Delta}$ brain compared to Wt brain (fig. 2b: median Wt = 4, HIF-1 $\alpha^{\Delta/\Delta}$ = 6, $p < 0.01$), than in the hippocampus, where there was no difference in degree of injury (fig. 2c: median Wt = 7, HIF-1 $\alpha^{\Delta/\Delta}$ = 7, $p > 0.75$). The total number of Wt mice was 27 (13 male, 14 female); the total number of HIF-1 $\alpha^{\Delta/\Delta}$ mice was 11 (8 male, 3 female). There is no effect on injury score associated with sex, although the number of females in the HIF-1 $\alpha^{\Delta/\Delta}$ group, while adequate, is small compared to males and thus may limit this analysis.

Immunofluorescence

Immunofluorescent staining of Wt (fig. 3a–d) and HIF-1 $\alpha^{\Delta/\Delta}$ (fig. 3e–h) brain 24 h after HI indicates a relative lack of HIF-1 α in the HIF-1 $\alpha^{\Delta/\Delta}$ cortex (fig. 3f) compared to Wt (fig. 3b). Double-labelling with NeuN demonstrates that HIF-1 α is localized primarily in neurons in the Wt cortex (fig. 3c). Double-labelling is virtually absent in the HIF-1 $\alpha^{\Delta/\Delta}$ brain (fig. 3g). A small amount of nonneuronal HIF-1 α appears to be present in Wt, and possibly to an even lesser extent in HIF-1 $\alpha^{\Delta/\Delta}$ brains (small red dots). The addition of DAPI shows the nucleus in relation to HIF-1 α and NeuN (fig. 3d, h).

Discussion

We hypothesized that HIF-1 α serves to protect the neonatal brain from injury due to HI, and this study supports this hypothesis, since genetic reduction of HIF-1 α in the mouse forebrain leads to a greater degree of injury. While it is well established that HIF-1 α is the primary regulator of oxygen homeostasis [15], its role in injury and disease has yet to be fully understood. However, HIF-1 α may be especially important in the developing brain. The developing brain is particularly vulnerable to oxidative stress, with its high rate of oxygen consumption, low concentrations of antioxidants and availability of redox-active iron [27, 28]. The fact that HIF-1 α and its target gene, VEGF, have an important role in angiogenesis indicates potential for repair after brain injury. It has recently been shown in the adult ischemic brain, for example, that neural stem/progenitor cells have a role in vas-

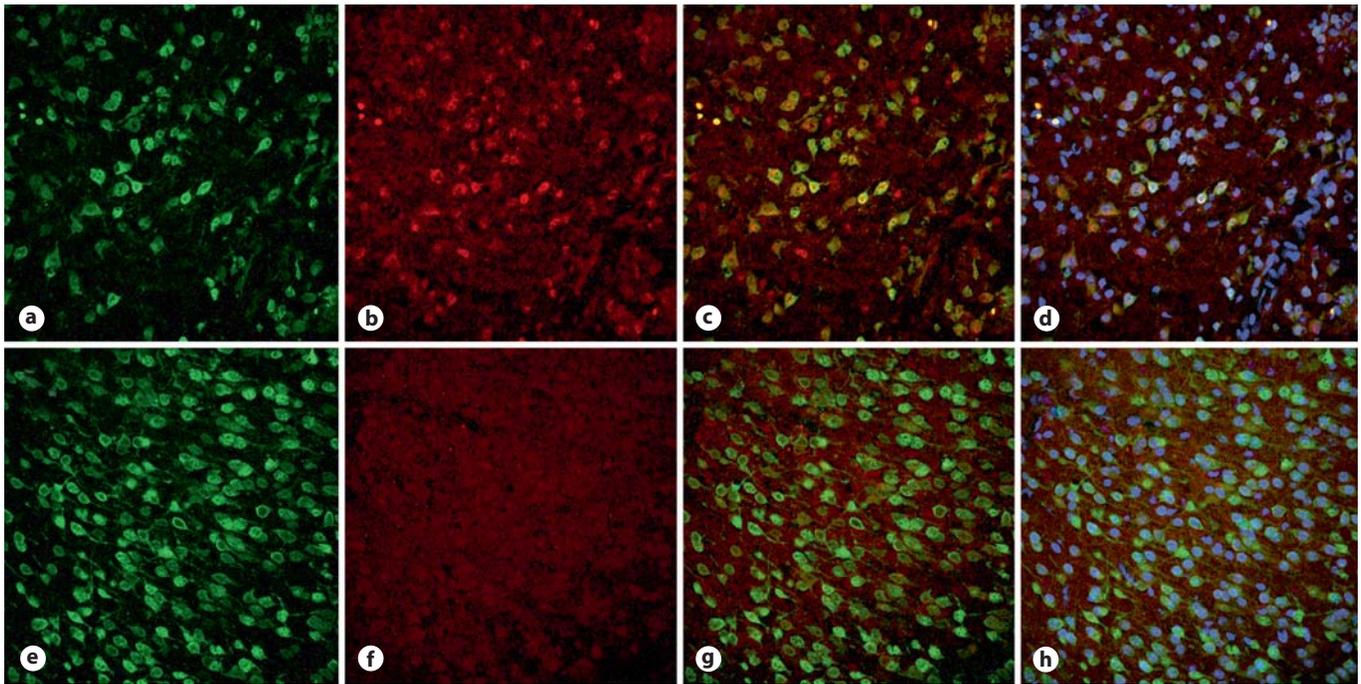


Fig. 3. Localization of HIF-1 α by immunofluorescent staining. Immunofluorescently labeled cells in the cortex of representative Wt (**a–d**) and HIF-1 $\alpha^{\Delta/\Delta}$ (**e–h**) brains 24 h after HI at P7.

a, e NeuN. **b, f** HIF-1 α . **c, g** Merged NeuN and HIF-1 α . Yellow cells indicate localization of HIF-1 α , where present, to neurons. **d, h** NeuN, HIF-1 α and DAPI for visualization of nuclei.

cular stabilization, which is mediated via HIF-1 α -VEGF signaling pathways [29]. In neonatal HI in the rat, exogenously applied VEGF (via intracerebroventricular injection) is neuroprotective [30]. Increased expression and stabilization of HIF-1 α after neonatal brain injury may also be associated with neurogenesis and repair [9, 31].

While we hypothesized that a lack of HIF-1 α would exacerbate neonatal HI injury (but have confirmed that here in the developing mouse brain), there is conflicting evidence on whether HIF-1 α is beneficial or detrimental. In a different line of neural cell-specific HIF-1 α -deficient mice than we used, defective brain development was seen embryonically, leading to defects including hydrocephalus in the adult [32]. The mice we used do not express the deletion of HIF-1 α embryonically; the deleted HIF-1 α allele is detectable by Southern blot beginning at P5 [20]. In addition, all mice we used appeared normal both as neonates and adults. Reports from studies in adult ischemia have shown conflicting evidence in regard to the role of HIF-1 α in this same line of transgenic mice. In a model of global ischemia, which produces a distinct pattern of injury, HIF-1 $\alpha^{\Delta/\Delta}$ brains had less injury than Wt counterparts [21]. On the other hand, with transient focal cere-

bral ischemia, HIF-1 $\alpha^{\Delta/\Delta}$ brains had increased injury and reduced survival compared to Wt counterparts [20]. Recently, in a rat model of neonatal HI, acute, but not delayed, pharmacological inhibition of HIF-1 α resulted in reduced injury [33]. In this study, we chose to produce a moderate degree of HI injury for two reasons: to keep mortality low, and to avoid differential effects of severe injury on the induction of death mechanisms [34, 35]. With our model, genetic reduction of neuronal HIF-1 α resulted in a worsening of injury after neonatal HI, especially to the cortex. We did not measure HIF-1 α in the hippocampi collected from these same brains because HIF-1 α was measured in the nuclear fraction of the homogenate and the small size of the hippocampi prohibits this measurement without pooling samples. It has also been shown that calcium/calmodulin-dependent kinase II is absent in the hippocampal synapse at P2 and remains very low at P10, increasing dramatically at P35 [36]. Therefore, the lack of effect in the hippocampus may be due to the lack of expression of the promoter needed for deletion of the HIF-1 α gene. In addition, the hippocampus may express other isoforms or have differential susceptibility to the injury.

We have previously shown that HIF-1 α upregulation peaks at 8 h after transient ischemia in neonatal rats, and declines at 24 h [9]. Here, while it is clear that HIF-1 α is upregulated in Wt cortex immediately after hypoxia, and also at 24 h after HI, it remains for the future to be determined at what time point peak expression in the Wt cortex lies. There may also be a region-specific role for HIF-1 α in the setting of neonatal brain injury, since we found a difference in the degree of injury in the cortex, but not the hippocampus, when compared to Wt.

Given the large number of genes regulated by HIF-1 α and the disparate outcomes of HIF-1 α studies, based on different rodent species, ages, and models, it seems likely that a number of factors influence whether HIF-1 α has a deleterious or a protective effect. This may be due, in part, to the variable functions of different cell types and how they interact [37]. A recent study of neurons and astrocytes in co-culture showed that loss of HIF-1 α function in neurons reduced neuronal viability after hypoxia, yet selective loss of HIF-1 α function in astrocytes protected

neurons from hypoxia. This indicates that HIF-1 α may not only have different roles depending on cell type, but that role may be critical to survival [38].

Thus, the specific circumstances we have explored here may well have produced results specific to the developing brain. Whether HIF-1 α acts to promote or prevent injury is of critical importance in our quest to develop therapies for the human infant, as a greater understanding of the role of HIF-1 α in neonatal hypoxic-ischemic brain injury may lead to new avenues for therapies and improve outcome for babies affected by HI.

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