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Hypoxia induces angiogenic factors in brain microvascular endothelial cells

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Abstract

Hypoxia is increasingly recognized as an important contributing factor to the development of brain diseases such as Alzheimer's disease (AD). In the periphery, hypoxia is a powerful regulator of angiogenesis. However, vascular endothelial cells are remarkably heterogeneous and little is known about how brain endothelial cells respond to hypoxic challenge. The objective of this study is to characterize the effect of hypoxic challenge on the angiogenic response of cultured brain-derived microvascular endothelial cells. Brain endothelial cell cultures were initiated from isolated rat brain microvessels and subjected to hypoxia (1% O₂) for various time periods. The results showed that hypoxia induced rapid (≤ 0.5 h) expression of hypoxia-inducible factor 1 α (HIF-1 α) and that cell viability, assessed by MTT assay, was unaffected within the first 8 h. Examination of brain endothelial cell cultures for pro- and anti-angiogenic proteins by western blot, RT-PCR and ELISA revealed that within 0.5 to 2 h of hypoxia levels of vascular endothelial growth factor and endothelin-1 mRNA and protein were elevated. The expression of heme oxygenase-1 also increased but only after 8 h of hypoxia. In contrast, similar hypoxia exposure evoked a decrease in endothelial nitric oxide synthase and thrombospondin-2 levels. Exposure of brain endothelial cell cultures to hypoxia resulted in a significant ($p < 0.001$) decrease (94%) in tube length, an *in vitro* index of angiogenesis, compared to control cultures. The data indicate, despite a shift toward a pro-angiogenic phenotype, hypoxia inhibited vessel formation in brain endothelial cells. These results suggest that in brain endothelial cells expression of angiogenic factors is not sufficient for the development of new vessels. Further work is needed to determine what factors/conditions prevent hypoxia-induced angiogenic changes from culminating in the formation of new brain blood vessels and what role this may play in the pathologic changes observed in AD and other diseases characterized by cerebral hypoxia.

Keywords

Hypoxia; brain microvascular endothelial cells; endothelin-1; vascular endothelial growth factor; heme oxygenase-1; Alzheimer's; cerebral hypoperfusion

Introduction

The human brain, although only 2% of total body weight, accounts for 20% of oxygen consumption, reflecting its high rate of metabolic activity (Pimenta de Castro et al., 2010).

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The high energy demands of the brain render it especially susceptible to the deleterious effects of hypoxia. Hypoxia is increasingly recognized as an important contributing factor to the development of neurodegenerative diseases in the brain (Kaur and Ling, 2008; Peers et al., 2007; 2009; Quaegebeur and Carmeliet, 2010). Emerging evidence suggests that hypoxia is an important risk factor for the development of dementias, since patients suffering from cerebral ischemia or stroke are much more susceptible to dementias, particularly Alzheimer's disease (AD) (Desmond et al., 2002; Kalaria, 2000; Kokmen et al., 1996; Moroney et al., 1997). Furthermore, cerebral hypoperfusion is one of the major clinical features of AD and pathological changes caused by chronic hypoxia in the CNS are similar to those observed in AD (de la Torre, 2000; Lee et al., 2011; Miklossy, 2003). Despite data which suggest a strong link between cerebral hypoxia and AD, the mechanisms whereby hypoxia contributes to neurodegenerative disease processes are unknown.

In the periphery, hypoxia is a powerful regulator of angiogenesis. The angiogenic process is complex and involves a sequence of discrete steps beginning with endothelial activation and culminating in the formation of new blood vessels. Genes involved in the different stages of angiogenesis have been shown to be responsive to hypoxia in tissue culture (Pugh and Ratcliffe, 2003). Hypoxia regulates angiogenesis by modulating a large number and variety of pro- and anti-angiogenic factors (Enholm et al., 1997; Liu et al., 1995; Oh et al., 1999). Regulation of genes that encode proteins involved in angiogenesis occurs via activation of hypoxia-inducible factor (HIF). HIF, a heterodimeric complex consisting of an oxygen-destructible α subunit and an oxygen-indestructible β subunit, is a sequence-specific DNA-binding protein that affects transcription of a broad range of genes (Brahimi-Horn and Pouyssegur, 2009). The oxygen sensitive α subunit ensures a quick response to minute changes in oxygen concentration by regulating proteasomal degradation of HIF-1 (Semenza, 1999). One of the most versatile angiogenic factors stimulated by hypoxia is vascular endothelial growth factor (VEGF) (Forsythe et al., 1996; Liu et al., 1995). VEGF is induced and regulated in a strictly dose-dependent manner by HIF-1 (Ferrara et al., 2003). Other angiogenic factors such as thrombin, endothelin-1 (ET-1), and heme oxygenase-1 (HO-1) have also been reported as factors associated with hypoxia (Landau et al., 2000; Motterlini et al., 2000; Yamashita et al., 2001). In contrast, anti-angiogenic factors such as thrombospondins (TSPs) are inhibited by hypoxia (Laderoute et al., 2000), although this varies depending on cell type (Phelan et al., 1998).

Despite advancements in research on hypoxia-induced angiogenesis in the periphery, little is known about how brain endothelial cells respond to hypoxic challenge. The vascular endothelium is a remarkably heterogeneous organ. Endothelial cells have the ability to differentiate both in structure and function in response to the needs of diverse tissue environments (Craig et al., 1998; Molema, 2010). Endothelial cells from different vascular beds differ in their morphology, cellular behavior and responses to injury (Langenkamp and Molema, 2009). Brain endothelial cells are a highly differentiated, specialized blood-brain barrier endothelial phenotype possessing unique biochemical and structural features not found in other vascular beds (Abbott et al., 2006; Zlokovic, 2008). The cerebrovasculature is increasingly implicated as contributory to the development of neurodegenerative diseases such as AD (Bell and Zlokovic, 2009; Brown and Thore, 2011; Grammas, 2011; Nation et al., 2011). A growing literature shows biochemical and functional changes in the cerebrovasculature in AD including expression and/or release of proteins related to vascular endothelial activation and angiogenesis such as VEGF, ET-1, HIF-1 α , and thrombin (Grammas et al., 2006; Thirumangalakudi et al., 2006; Luo and Grammas, 2010; Yin et al., 2010). However, the function of angiogenic proteins in the AD brain is unknown and their significance is controversial (Bell and Zlokovic, 2009; Grammas, 2011; Paris et al., 2010., Wu et al., 2005). Understanding the effects of hypoxia on brain endothelial cells is important to determining mechanisms of hypoxia-induced damage in the brain and its links to AD.

The objective of this study is to characterize the effect of hypoxic challenge on the angiogenic response of cultured brain-derived microvascular endothelial cells

Methods

Treatment of endothelial cell cultures

Rat brain endothelial cell cultures were isolated from rat brain microvessels, as previously described (Diglio et al., 1993). The purity of these cultures was confirmed using antibodies to the endothelial cell surface antigen factor VIII. Endothelial cells used in this study (passages 8 to 15) were maintained in Dulbecco's modified Eagle's medium (DMEM, Sigma-Aldrich, St Louis, MO) supplemented with 10% fetal bovine serum (FBS), 1% antibiotic/antimycotic and 2 mM glutamine. Confluent endothelial cell cultures were washed 3 times with Hank's balanced salt solution (HBSS, Gibco, Grand Island, NY) and then incubated with serum-free DMEM. Cells were exposed to hypoxia (1% O₂) for different periods of time in a humidified incubator at 37 °C. Cells grown under normoxic conditions (21% O₂) served as a positive control.

Measurement of cell survival by MTT assay

Cell viability was determined using the MTT reagent (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide, Promega, Madison, WI) as follows. Treatment medium was replaced with fresh treatment medium containing 20 µl/ml of the Cell Titer 96 Aqueous One Solution and incubated for 10 min at 37°C after which optical density was measured at 490 nm using a microplate reader. The quantity of soluble formazan product, as measured by the amount of absorbance, was directly proportional to the number of viable cells. The number of viable cells after treatment was determined by measuring optical density and viability expressed as percent of untreated controls.

Measurement of tube formation

Geltrex™ reduced growth factor basement membrane matrix (Invitrogen, Carlsbad, CA, #12760) was added to wells of a 24-well plate (150 µl/well) and incubated at 37 °C for 30 min. After Geltrex™ matrix was polymerized, brain microvascular endothelial cells were seeded onto the matrix at 10⁵ cells/well, and maintained in DMEM supplemented with 10% FBS. Plates were exposed to hypoxia (1% O₂) or normoxia (21% O₂) at 37°C for 4 h (Kourembanas et al., 1994; Loboda et al., 2006). After staining with fluorescent dye Calcein (Invitrogen, #C3099) for 30 min, the tube-like structures were visualized and captured using Olympus IX71 microscope at 10X magnification. Tube length was analyzed and quantitated using image processing software (ImageJ) available from the National Institutes of Health.

Western blot analysis

Total protein from cell cultures was extracted using lysis buffer (0.1% SDS, 1% Triton X-100, and 0.5% phenylmethyl-sulfonyl fluoride), and protein concentration determined by the Bradford method using Bio-Rad protein reagents (Bio-Rad, Hercules, CA). Samples were resolved in 10% poly-acrylamide gel (25 – 30 µg/lane) using SDS-PAGE and transferred to PVDF membranes. The blots were blocked with 5% nonfat milk in Tris-buffered saline containing 0.1% Tween 20 (TBST) for 1 h. Membranes were then incubated with primary antibodies (HIF-1α: Abcam, Cambridge, MA, #1; VEGF: Abcam, #1316; TATA-binding protein (TBP): Abcam, #818; β-actin: Abcam, #6276; endothelial nitric oxide synthase (eNOS): Abcam, #66127; inducible nitric oxide synthase (iNOS): Santa Cruz Biologicals, Santa Cruz, CA, #651) diluted 1:500 in TBST for 2 h, washed 3 times with TBST, and incubated with peroxidase-conjugated secondary antibodies (HIF-1α, VEGF, TBP and β-actin: anti-mouse antibody, Bio-Rad, #170-6516; eNOS and iNOS: anti-rabbit

antibody, Bio-Rad, #172-1019) in TBST (1:2000 dilution) for 1 h. Membranes were washed 3 times with TBST, developed with chemiluminescent reagents and visualized on film. The average intensities over the area of the bands were measured using Quantity One software (Bio-Rad).

RT-PCR analysis of mRNA expression

Total RNA was extracted using the Trizol method (Invitrogen), and 4 µg of RNA were reverse transcribed into cDNA using random primers (Roche, Indianapolis, IN) according to manufacturer's instructions. cDNA strands (0.2 µg) were amplified by PCR using the gene specific primers listed in Table 1. The reaction was performed by GoTaq Green master mix (Promega, #M7123) for 30 cycles, each at 95°C for 40 s, 55°C for 40 s and 72°C for 1 min. The PCR products were visualized on a 1.2 % agarose gel using UV transillumination.

Detection of VEGF or ET-1 by ELISA

Endothelial cell culture medium was collected and concentrations of VEGF or ET-1 in the medium were determined using ELISA kits from R&D Systems (Minneapolis, MN). According to manufacturer's instructions, 100 µl of standard or sample were added to each well. After incubation at room temperature on a shaker for 2 h, wells were washed 4 times and then incubated with 200 µl conjugate/detection antibody for 2 h at room temperature on the shaker. For VEGF detection, the wells were coated with the capture antibody prior to adding the samples. For ET-1, plates pre-coated with the capture antibody were used. After streptavidin-HRP was conjugated to target molecules, it was detected by addition of 200 µl of O-phenylene diamine-H₂O₂ (Pierce, Rockford, IL) for 20 min. Optical density was measured at 450 nm using a microplate reader.

Assessment of HIF-1α expression by immunofluorescence

Endothelial cells were fixed in cold acetone for 5 min and washed 3 times with phosphate buffered saline. After 30 min incubation with 1% BSA in phosphate buffered saline tween to block nonspecific binding, cells were incubated (4°C) simultaneously with two primary antibodies (HIF-1α: Abcam #1, 1:200; VWF: Santa Cruz #14014, 1:50) overnight. Cells were washed then incubated with two secondary antibodies HIF-1α: Alexa Fluor 488 goat anti-mouse IgG (H+L), Invitrogen #A11001; for VWF: Alexa Fluor 594 goat anti-rabbit IgG (H+L), Invitrogen #A11012) diluted 1:400 with antibody diluent for 1 h. After stringent washing with TBST, cells were counter-stained with nuclear staining DAPI solution for 20 min. After washing and drying at room temperature, samples were observed with an Olympus IX71 microscope.

Statistical analysis

Data from each experiment are expressed as mean ± standard deviation (SD). Comparisons between control and treatment groups were conducted using a two-tailed Student's *t*-test. For multiple comparisons among control and treatment groups, the one-way ANOVA followed by Bonferroni's multiple comparison test for multiple samples were used. Statistical significance was determined at $p < 0.05$.

Results

Time-dependent effects of hypoxia on cell survival and induction of HIF-1α in cultured brain endothelial cells

Confluent brain microvascular endothelial cell cultures were subjected to hypoxia (1% O₂) for various periods of time and cell viability measured by MTT assay. The results showed that hypoxic stress did not affect cell viability within the first 8 h. In contrast, exposure of

cells to hypoxia for ≥ 8 h evoked significant ($p < 0.001$) cell death (45%). After 8 h of hypoxia, reoxygenation of cell cultures for an additional 8 h significantly increased cell death an additional 40% ($p < 0.01$) compared to 8 h hypoxia-only treated cultures.

Exposure of cultured brain endothelial cells to hypoxia caused expression ($p < 0.01$) of HIF-1 α protein within 0.5 h (Fig. 1A). Expression of HIF-1 α was highly significant ($p < 0.001$) at 2 and 4 h of hypoxia. Induction of HIF-1 α protein expression was confirmed by immunofluorescent comparison of cultures exposed to normoxic or hypoxic conditions (4 h) (Fig. 1B). Hypoxia increase HIF-1 α mRNA expression (177%) but not until 8 h. Reoxygenation for 2 h significantly reduced the increase evoked by hypoxia on protein ($p < 0.01$) (45%) and mRNA ($p < 0.05$) (61%) levels, respectively.

Exposure of endothelial cells to hypoxia induces VEGF expression and secretion

The effect of hypoxia on the expression and release of VEGF from cultured brain microvascular endothelial cells was determined by Western blot, RT-PCR and ELISA. Both cell-associated protein (Fig. 2A) and mRNA (Fig. 2B) levels of VEGF were significantly increased by hypoxia treatment in a time-dependent manner. Two hours of reoxygenation restored VEGF expression to control levels at both protein and mRNA levels (Fig. 2). Similarly, exposure of cultured microvascular brain endothelial cells to hypoxia resulted in elevated release of VEGF into culture medium compared to normoxic controls at the same time points (Fig. 3).

Hypoxia induces an increase in ET-1 and a decrease in eNOS

In vascular endothelial cells, the regulation of ET-1 and nitric oxide (NO) is often coordinated (Boulanger and Luscher, 1990). Hypoxic stress evoked a significant increase of ET-1 released into culture medium compared to normoxic controls examined at the same time points (Fig. 3). A two-fold increase in mRNA levels of ET-1 was observed with hypoxia treatment as early as 0.5 h (Fig. 4A). With 2 h of reoxygenation ET-1 mRNA levels recovered to that of normoxic controls (Fig. 4A). In contrast to ET-1, 2 h of hypoxia treatment resulted in a significant ($p < 0.001$) decrease in expression of eNOS (Fig. 4B). Two hours of reoxygenation was able to partially blunt the reduction in eNOS protein levels but did not restore expression to control levels (Fig. 4B). Exposure of endothelial cells to hypoxia did not affect expression of iNOS (Fig. 4B).

Hypoxia induces expression of HO-1 and reduces levels of TSP-2

Exposure of cultured endothelial cells to hypoxia did not affect expression of mRNA for HO-1 until 8 h (Fig. 5A). At 8 h of hypoxia there was a significant ($p < 0.001$) increase in levels of HO-1. Reoxygenation (2 h) was able to restore expression of HO-1 to control levels (Fig. 5A).

A significant ($p < 0.01$) decrease in TSP-2 mRNA expression was detectable at 2 h of hypoxia and by 8 h the level of TSP-2 was less than 25% ($p < 0.001$) of that demonstrable in normoxic controls (Fig. 5B). Reoxygenation (2 h) was unable to restore expression of TSP-2 (Fig. 5B). Exposure of endothelial cells to hypoxia did not affect expression of TSP-1 (Fig. 5B).

Hypoxia inhibits vessel formation in cultured brain endothelial cells

Experiments performed to determine the effect of hypoxia on brain endothelial cell vessel formation in culture showed that endothelial cells seeded onto extracellular matrix form vessels after 4 (Fig. 6A). In contrast, exposure of cultures to hypoxia significantly inhibited vessel formation (Fig. 6A). Quantification of stained cultures showed a 94% reduction in

tube length in hypoxia-exposed endothelial cell cultures compared to control cultures (Fig. 6B).

Discussion

Hypoxic challenge to the brain is a characteristic feature of both acute brain injury and chronic neurodegenerative diseases (Hackett, 1999; Peers et al., 2009). The chain-of-events, initiated by hypoxia, which culminate in neuronal dysfunction and/or death remains unclear. However, a key role for the cerebral endothelium, which is responsible for tightly regulating the CNS milieu and is a primary sensor of blood oxygen levels, is likely (Madri, 2009). HIF-1 α , a master regulator of the cellular and physiological response to hypoxia (Semenza, 1999), is elevated in brain blood vessels in AD (Grammas et al., 2006). HIF-1 α levels are controlled by multiple transcriptional, post-transcriptional and post-translational mechanisms (Semenza, 1999). In this study we observe a time-dependent increase in accumulation of HIF-1 α protein inside the nucleus of brain microvascular endothelial cells challenged with hypoxia; consistent with the stabilization of HIF-1 α under hypoxic conditions (Brahimi-Horn and Pouyssegur, 2009). Brain microvascular endothelial cells produce more than a 2-fold increase in HIF-1 α protein levels compared to controls within a half hour of hypoxia exposure, suggesting that the initial HIF-1 α response to hypoxia is due to post-translational regulation of its expression. In contrast, there is no change in HIF-1 α mRNA until 8 h, indicating that transcriptional regulation is delayed in brain microvascular endothelial cells. In the periphery, hypoxia activates HIF-1 α which in turn initiates a series of gene expression changes in vascular endothelial cells that are consistent with angiogenesis involving VEGF, ET-1, eNOS, and HO-1 TSPs (Kourembanas et al., 1991; Liu et al., 1995; Strijdom et al., 2009; Sun et al., 2002; Tenan et al., 2000). Therefore, in the current study we examined the effect of hypoxic challenge on expression of these factors by cultured brain endothelial cells as discussed below.

VEGF, a multifunctional cytokine, induces endothelial cell migration and proliferation (Neufeld et al., 1999). Stimulation of VEGF gene expression by hypoxia is thought to be mediated by the specific binding of HIF-1 α to hypoxic response elements in the regulatory region of the VEGF gene (Liu et al., 1995). In brain microvascular endothelial cells we demonstrate that HIF-1 α stimulates both expression and secretion of VEGF. In the current study we show a significant increase in VEGF protein precedes an increase in mRNA which is consistent with studies in other cell types that document post-transcriptional regulation of VEGF levels (Shenberger et al., 2007). VEGF has been shown to regulate HO-1 expression and activity in vascular endothelial cells. The cytoprotective protein HO-1 is also pro-angiogenic (Suzuki et al., 2003). In one study, after 6 h of hypoxia HO-1 mRNA expression increases about two-fold while HO-1 protein levels are not significantly affected. Here we show an increase in HO-1 mRNA levels, but only after 8 h of hypoxia. Hypoxia-induced HO-1 expression in bovine aortic endothelial cells has also been documented (Mottlerini et al., 2000; Sun et al., 2002).

ET-1, a vasoconstrictor produced in vascular endothelial cells, is also recognized as an angiogenic factor (Knowles et al., 2005). In human umbilical vein endothelial cells, hypoxia induces ET-1 gene expression and secretion (Kourembanas et al., 1991) which is consistent with the results obtained herein from brain-derived microvascular endothelial cells. ET-1 stimulation by hypoxia is mediated by HIF-1 α (Hu et al., 1998) and antagonized by NO (Blanchard et al., 1992). It is well documented that NO and ET-1 regulate each other in the vascular endothelium and thus modulate vascular tone as well as response to injury (Bourque et al., 2011). In the current study, although there is no change in expression level of iNOS we document a dramatic loss in eNOS protein levels in brain microvascular endothelial cells after exposure to hypoxia for 2 h. Similarly, Strijdom et al. (2006) reported

a significant decrease in eNOS levels in rat hypoxic cardiomyocytes exposed to hypoxia. Our data showing an increase in ET-1 and decrease in eNOS are consistent with literature that shows differential regulation of these two mediators in vascular endothelial cells. The mechanism whereby hypoxia reduces eNOS is uncertain; however, it has been reported that hypoxia decreases eNOS at the message level by inducing changes in transcription kinetics and stability of eNOS mRNA (McQuillan et al., 1994). Our data indicate that hypoxia can also reduce eNOS at the protein level. These data are consistent with what has been published in human coronary artery and microvascular endothelial cells (Loboda et al., 2006; Olszewska-Pazdrak et al., 2009;). Finally, we examined expression of TSP-1 and TSP-2, important physiological inhibitors of angiogenesis (Vailhé and Feige, 2003), and find that in brain microvascular endothelial cells hypoxia decreases TSP-2 gene expression but does not affect expression of TSP-1. Taken together, these data demonstrate a pro-angiogenic shift in brain endothelial cells exposed to hypoxia *in vitro*. The increased expression of pro-angiogenic factors and the decrease in angiogenic inhibitors favor the formation of new blood vessels. However, our data indicate that despite this pro-angiogenic phenotypic shift in brain endothelial cells, vessel formation is inhibited by hypoxia. In our study, culture of brain-derived endothelial cells on a cellular matrix that favors angiogenesis results in tube formation by 4 h. In contrast, in cultures exposed to hypoxia formation of tubes is inhibited. Quantitation of tube formation length shows significantly ($p < 0.001$) less tube formation in hypoxia-exposed cultures compared to normoxic control cultures. These data are in contrast to considerable literature that documents an increase in angiogenesis in response to hypoxia (Yamakawa et al., 2003; Tang et al., 2004). However, an angiogenic response to hypoxia is not a universal finding, as some studies demonstrate that hypoxia inhibits vessel/tube formation (Olszewska-Pazdrak et al., 2009). In this regard, Isner (2002) finds that in patients with myocardial ischemia, angiogenic responses to hypoxia are defective or absent. In a study where human coronary endothelial cells are exposed to 1% O₂, similar to the conditions of our study, there is a decrease in both basal and VEGF-mediated tube formation as well as in active eNOS (Olszewska-Pasdrak et al., 2009).

The data obtained in the current study may explain, in part, some paradoxical findings regarding hypoxia in the AD brain. Cerebral hypoperfusion is one of the major clinical features in AD and likely contributes to the clinical and pathological manifestations in this disease (de la Torre, 2000; Kalaria, 2000; Miklossy, 2003). Genome-wide expression profiling in the AD brain has identified a marked upregulation of genes that promote angiogenesis (Pogue and Lukiw, 2004). We have shown AD brain microvessels express or release inflammatory proteins, including thrombin, VEGF, angiopoietin-2, tumor necrosis factor- α , transforming growth factor- β , interleukin (IL) IL-1 β , IL-6, IL-8, monocyte chemoattractant protein-1, matrix metalloproteinases, and integrins (Grammas and Ovase, 2001; 2002; Grammas et al, 2006, Thirumangalakudi et al, 2006; Yin et al., 2010), all of which have been implicated in angiogenesis. Despite increases in several pro-angiogenic factors in the AD brain, evidence for increased vascularity in AD is lacking. On the contrary, it has been suggested the angiogenic process is delayed and/or impaired in aged tissues, with several studies showing decreased microvascular density in the AD brain (Buee et al., 1994; 1997; Edelber and Reed, 2003; Paris et al., 2010,). Therefore, lack of vessel formation despite the increase in pro-angiogenic factors evoked by hypoxia suggests these angiogenic factors are not sufficient for the completion of the angiogenic process and the development of new vessels by brain endothelial cells. Further work is needed to determine what factors/conditions prevent hypoxia-induced angiogenic changes from culminating in the formation of new brain blood vessels.

Our results showing that brain-derived endothelial cells respond differently to hypoxia than most endothelial cells derived from peripheral vessels may reflect the functional heterogeneity of endothelial cells (Aird, 2003). Brain endothelial cells are highly

differentiated and specialized in their blood-brain barrier function (Zlokovic. 2008; Grammas et al., 2011). Although the effect of astroglial-derived paracrine factors on maintenance of blood-brain barrier properties has been extensively documented (Abbott et al., 2006), other data also indicate there is an intrinsic endothelial identity, depending on vascular bed origin, that is stable and heritable (Aird, 2003; Chi et al., 2003). Indeed, there is evidence certain site-specific properties of endothelial cells are epigenetically programmed such that their maintenance is no longer dependent on signals from the extracellular milieu. For example, DNA microarray studies of multiple passaged endothelial cells cultured from different sites reveal differences in transcriptional profiles (Chi et al., 2003). The brain vasculature appears to be especially sensitive to hypoxia and oxidative stress. This sensitivity may in part be due to higher levels of NAD(P)H-oxidase in brain endothelial cells compared to endothelial cells in peripheral vessels (Closhen et al., 2010). In this regard, the inflammatory protein CRP evokes NAD(P)H-oxidase dependent functional derangements in brain- but not aorta-derived endothelial cells (Closhen et al., 2010). In addition, the higher concentration of mitochondria in cerebrovascular brain endothelial cells relative to other endothelia (Oldendorf et al., 1977) may render them more susceptible to the effects of hypoxia.

The angiogenic factors produced by brain endothelial cells in response to hypoxia have pleiotropic effects in the brain and likely important consequences for brain function. For example, in the brain HO-1 is a mediator with both beneficial and deleterious properties. The mechanisms responsible for excessive iron deposition and mitochondrial insufficiency in the aging and degenerating nervous system remain poorly understood; HO-1 has been implicated in this process (Schipper et al., 2009; Schipper, 2011). In rat astroglia transfected with the human HO-1 gene, mitochondrial iron trapping is abrogated by HO-1 inhibitors. Furthermore, HO-1 immunoreactivity is enhanced greatly in neurons and astrocytes of the hippocampus and cerebral cortex of AD brains as well as in the substantia nigra in Parkinson's disease (Schipper, 2000). These results suggest HO-1 over-expression contributes to the pathological iron deposition and mitochondrial damage documented in these neurodegenerative disorders.

VEGF has direct neurotrophic effects and can protect neurons from exogenous injury reducing neuronal cell death in vitro evoked by hypoxia, glutamate or serum-deprivation (Tolosa et al., 2008; Tolosa et al., 2009; Wick et al., 2002). In contrast to these neuroprotective effects, VEGF overexpression is associated with many CNS disorders. In the AD brain, VEGF is deposited in the walls of intraparenchymal vessels as well as in clusters of reactive astrocytes (Kalaria et al., 1998; Thirumangalakudi et al., 2006). In addition, intrathecal levels of VEGF in AD are related to clinical severity and to intrathecal levels of A β (Tarkowski et al., 2002). Because VEGF has a potentiating effect on CNS inflammation and increases in vascular permeability, continuous upregulation of VEGF at sites of brain injury may drive chronic neuroinflammation. Similarly, ET-1 has been shown to be both neuroprotective as well neurotoxic (Luo and Grammas., 2010). Whether ET-1 is neurotoxic or neuroprotective may depend on the expression of other inflammatory mediators as well as the timing of exposure. In this latter regard, we have shown that the ET-1 significantly increases neuronal survival when cells are challenged with oxidative stress (H₂O₂) or thrombin, but that this neuroprotective effect requires pretreatment (Luo and Grammas., 2010). The angiogenic factors produced by brain endothelial cells in response to hypoxia have pleiotropic effects in the brain. A thorough understanding of the identity, regulation and function of these proteins could shed light on the basis for cell dysfunction and death in the brain in conditions characterized by cerebral hypoxia.

Conclusions

The results of the current study indicate that despite a shift toward a pro-angiogenic phenotype hypoxia inhibited vessel formation in brain endothelial cells. These results suggest that in brain endothelial cells expression of angiogenic factors is not sufficient for the development of new vessels. Further work is needed to determine what factors/conditions prevent hypoxia-induced angiogenic changes from culminating in the formation of new brain blood vessels and what role this may play in the pathologic changes observed in AD and other diseases characterized by cerebral hypoxia.

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Highlights

- Effects of hypoxia on brain microvascular endothelial cells (ECs) were examined
- Hypoxia resulted in the increased expression and release of VEGF protein and mRNA
- Increased ET-1 was released into the culture medium under hypoxic conditions
- Hypoxia resulted in a decrease in expression of eNOS as well
- This study suggest hypoxia evokes a pro-angiogenic phenotype in ECs

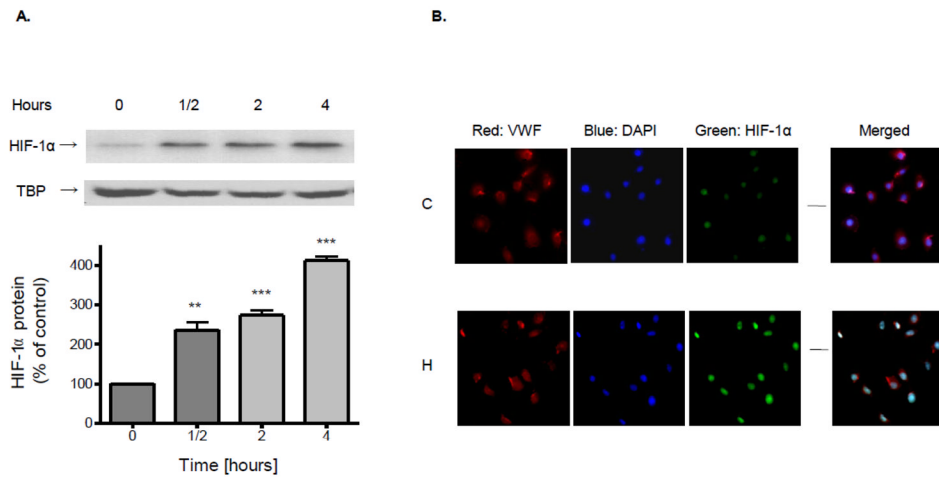


Figure 1.

Confluent brain microvascular endothelial cell cultures were subjected to hypoxia (1% O₂) for various periods of time. (A) Nuclear proteins were extracted and resolved using SDS-PAGE. HIF-1α or TBP protein was detected by Western blot probed with corresponding antibodies. Data in the bottom panel are the means ± SD of 3 experiments and expressed as percent of untreated control. ** $p < 0.01$, *** $p < 0.001$ vs. 0 h. (B) After exposure to normoxia (C) or 1% O₂ hypoxia (H) for 4 h, brain microvascular endothelial cell cultures were processed for immunofluorescence. Cultures were fixed, incubated with antibodies to VWF and HIF-1α and counter stained with DAPI.

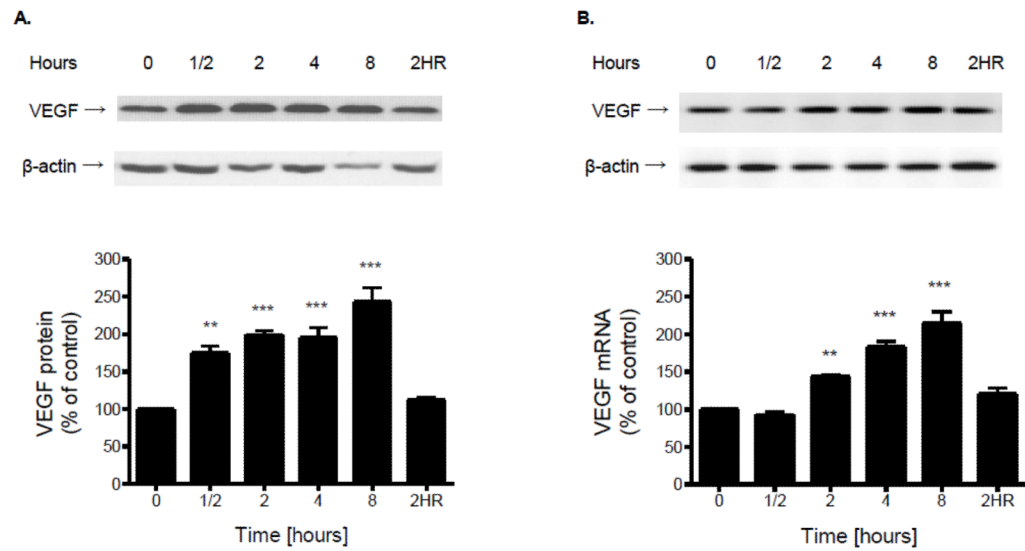


Figure 2.

Confluent brain microvascular endothelial cell cultures were subjected to hypoxia (1% O₂) for various periods of time. (A) Total protein was extracted and resolved using SDS-PAGE. VEGF or β -actin protein was detected by Western blot probed with corresponding antibodies. Protein or mRNA levels of VEGF were determined by normalizing the densities of their bands to those of β -actin. (B) Total RNA was extracted, reverse transcribed and amplified with gene specific primers for VEGF or β -actin. Data in the bottom panel are the means \pm SD of 3 experiments and expressed as percent of untreated control. 2HR: 2 h hypoxia followed by 2 h reoxygenation. ** $p < 0.01$, *** $p < 0.001$ vs. 0 h.

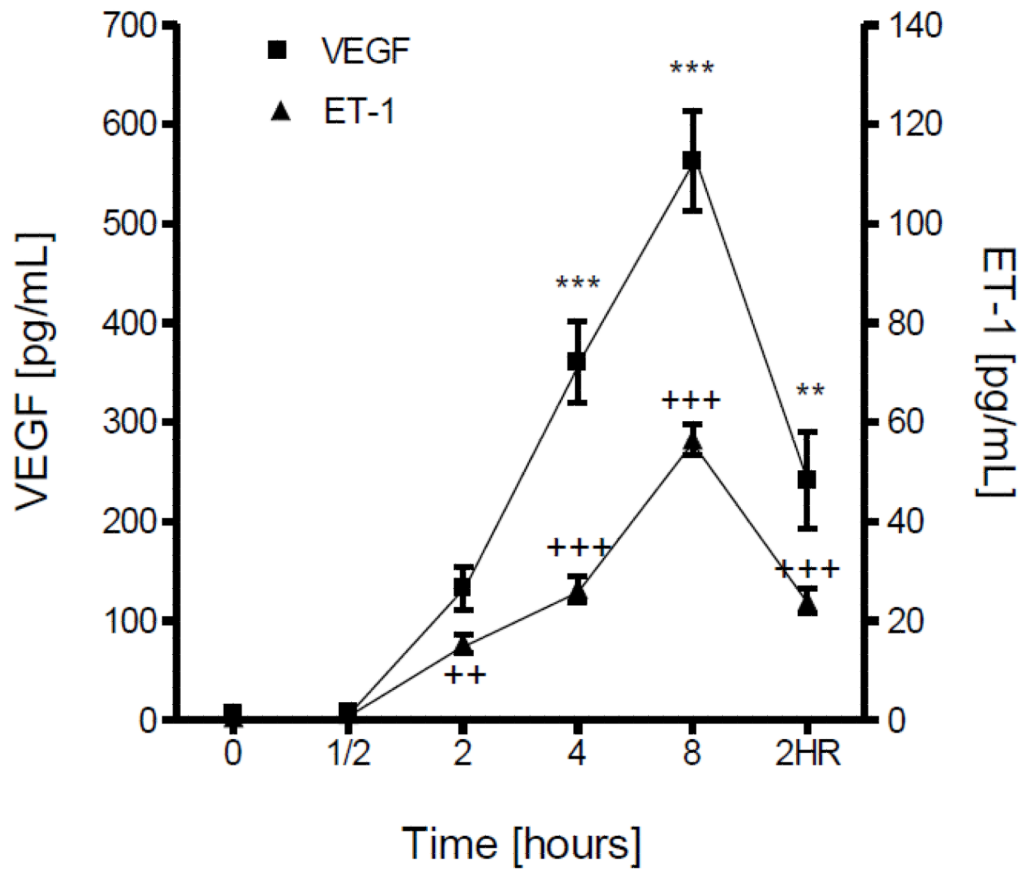
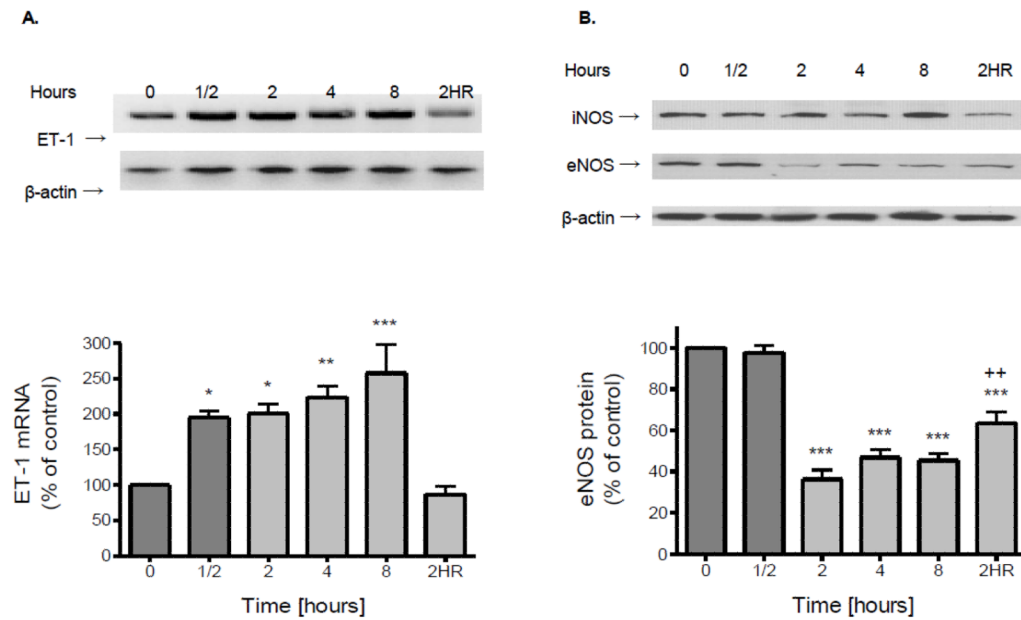


Figure 3. Confluent brain microvascular endothelial cell cultures were subjected to hypoxia (1% O₂) for various periods of time. ELISA was performed to determine VEGF or ET-1 concentration (pg/ml) in the culture medium. The amount of induced VEGF or ET-1 by hypoxia was compared to normoxia control cultures at the same time points. Results are means \pm SD of 3 experiments performed in triplicate. 2HR: 2 h hypoxia followed by 2 h reoxygenation. **/+ p <0.01, ***/+ p <0.001 vs. 0 h.

**Figure 4.**

Confluent brain microvascular endothelial cell cultures were subjected to hypoxia (1% O₂) for various periods of time. (A) Total RNA was extracted, reverse transcribed and amplified with gene specific primers for ET-1 or β -actin. (B) Total protein was extracted and resolved using SDS-PAGE. eNOS or β -actin protein was detected by Western blot probed with corresponding antibodies. Expression levels of ET-1 or eNOS were determined by normalizing the densities of their bands to those of β -actin. Data in the bottom panel are the means \pm SD of 3 experiments and expressed as percent of untreated control. 2HR: 2 h hypoxia followed by 2 h reoxygenation. * p <0.5, ** p <0.01, *** p <0.001 vs. 0 h; ++ p <0.01 vs. 2 h.

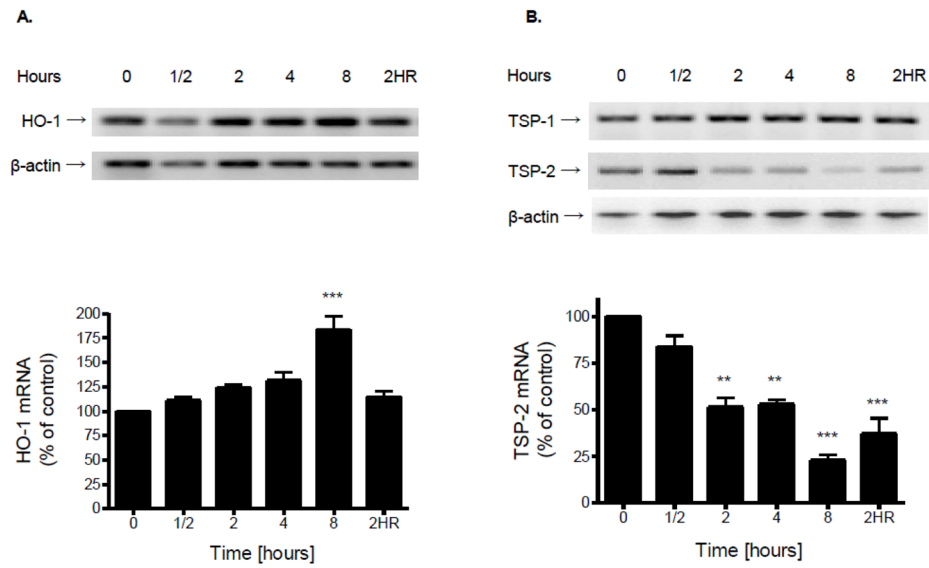


Figure 5. Confluent brain microvascular endothelial cell cultures were subjected to hypoxia (1% O₂) for various periods of time. Total RNA was extracted, reverse transcribed and amplified with gene specific primers for HO-1 (A); TSP-1 or TSP-2 (B). mRNA levels of HO-1, TSP-1, or TSP-2 were determined by normalizing their band densities to those of β-actin. Data in the bottom panel are the means ± SD of 3 experiments and expressed as percent of untreated control. 2HR: 2 h hypoxia followed by 2 h reoxygenation. ** $p < 0.01$, *** $p < 0.001$ vs. 0 h.

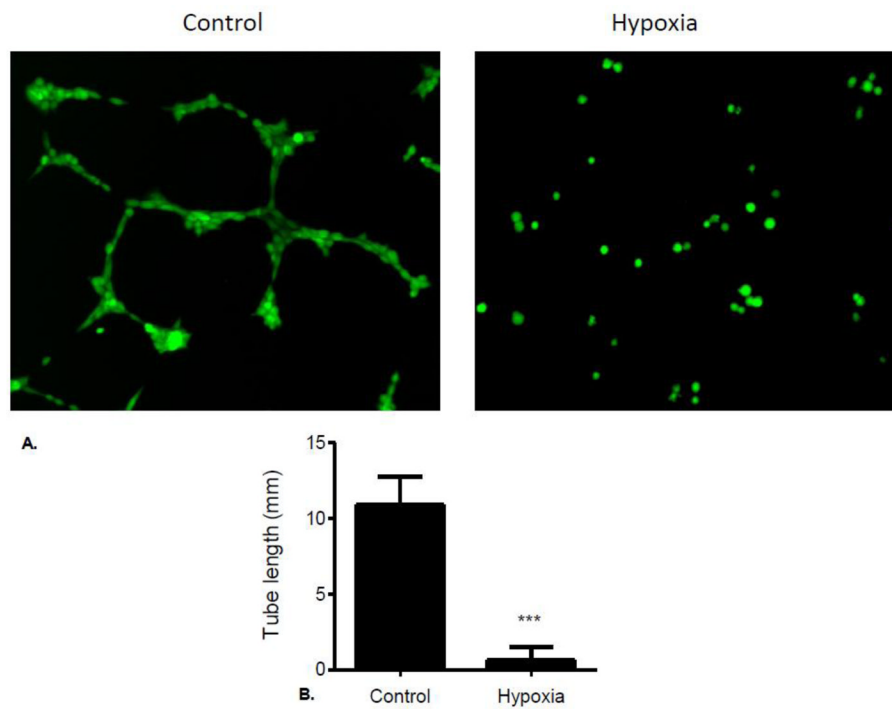


Figure 6. (A) Brain microvascular endothelial cells were seeded onto the layer of matrix at 10^5 cells/well, and maintained in DMEM supplemented with 10% FBS. Plates were incubated at 21% O_2 (Control) or 1% O_2 (Hypoxia) at 37°C for 4 h and then stained with fluorescent dye Calcein for 30 min. Tube-like structures were visualized and captured using Olympus IX71 microscope at 10x magnification. (B) Tube length was analyzed and quantitated using image processing software (ImageJ) available from the National Institutes of Health.

Table 1

Primers

Gene	Orientation	Sequence	Amplicon
HIF-1 α	Left primer	TGCATCTCCACCTTCTACCC	384 bp
	Right primer	CTGCTCCATTCCATCCTGTT	
VEGF	Left primer	GCCCATGAAGTGGTGAAGTT	360 bp
	Right primer	TTTCTTGCCCTTTCGTTTTT	
ET-1	Left primer	AGAAACAGCTGTCTTGGGAGCAGA	466 bp
	Right primer	TGCTGATGGCCTCCAACCTTCTTA	
HO-1	Left primer	TGCTCGCATGAACACTCTG	123 bp
	Right primer	TCCTCTGTCAGCAGTGCCT	
TSP-1	Left primer	TTCTGTTCATGTGTGGAAGC	696 bp
	Right primer	AAGGGTGAGAAGGACGTTGGTTGA	
TSP-2	Left primer	ATGAGTGTGCTGTGGTCGCAGATA	412 bp
	Right primer	TCCTGCCCAGAATTTGGCAGTTTG	
β -actin	Left primer	TGTCACCAACTGGGACGATA	165 bp
	Right primer	GGGGTGTGAAGGTCTCAA	