Site-specific induction of intestinal hypoxia-inducible factor-1α after hemorrhagic shock

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Abstract. The intestine is a major target organ in hemorrhagic shock (HS)-induced tissue injury. Hypoxia-inducible factor (HIF)-1α is the primary transcription factor responsible for regulating cellular response to changes in oxygen tension. Since HS is an acute hypoxic insult, the present study examined changes in the gene expression of HIF-1α in various regions of the intestine, as well as the distribution of HIF-1α protein in the intestinal cells of a rat model of HS. Levels of HIF-1α mRNA were marginally detectable in the intestine of sham-operated control animals, but obviously induced following HS. Duodenal, jejunal and colonic levels of HIF-1α mRNA robustly increased and reached a maximum during the early ischemic phase of HS, indicating that the regulation of HIF-1α expression in the intestinal mucosa is site-specific. These findings suggest that the intestine adaptively responds to hypoxic insult by HS in a regiospecific manner.

Introduction

Hemorrhagic shock (HS) induces a systemic inflammatory response that results in multiple organ failure with significant mortality and morbidity (1). The intestine is considered to contribute to the establishment of multiple organ damage, both as an instigator of systemic inflammation as well as a victim of end-organ injury (2).

Hypoxia-inducible factor (HIF)-1α is a basic helix-turn-helix transcription factor that controls cellular adaptation to hypoxia, including erythropoiesis, angiogenesis, vasodilatation and anaerobic metabolism (3). Previous studies have indicated that hypoxic insult activates HIF-1α mRNA and protein expression in various organs in a tissue-specific manner (4-6). However, few studies have investigated the effect of hypoxia on the expression of intestinal HIF-1α.

Here, we examined the effect of HS on HIF-1α expression in the intestine at the transcriptional and protein level in various regions of the intestine, because HS results in tissue hypoxia. HIF-1α mRNA and protein expression was found to be increased in the mucosal cells of the duodenum, jejunum and colon, but not in the mucosal cells of the ileum during the early ischemic phase of HS, indicating that the regulation of HIF-1α expression in the intestinal mucosa is site-specific. These findings suggest that the intestine adaptively responds to hypoxic insult by HS in a regiospecific manner.

Materials and methods

Animals. All procedures conformed to the guidelines for the care and use of laboratory animals established by the Animal Use and Care Committee of Okayama University Medical School. Male Sprague-Dawley rats weighing 380-420 g purchased from Charles River Laboratories Japan Inc. (Yokohama, Japan) were housed in a temperature-controlled (25°C) room with an alternating 12-h light/12-h dark cycle, and were given water and a chow diet ad libitum until the start of the experiments.

Experimental protocol. Rats anesthetized with intraperitoneal sodium pentobarbital (50 mg/kg) underwent a sham procedure or HS as described (7). In brief, heparinized polyethylene tubes were inserted into the left femoral artery to measure blood pressure and into the left femoral vein to induce hemorrhage by bleeding into a heparinized syringe (10 units/ml) over a period of 15 min to achieve a mean arterial blood pressure of 30±5 mmHg. This level of blood pressure was maintained for 60 min by further blood withdrawal or by re-infusion with the...
shed blood. At this point, the animals were resuscitated over a period of 15 min by returning all shed blood followed by sterile saline as required to restore blood pressure to the baseline level. The sham group underwent all instrumentation procedures, but blood was not collected. The animals breathed spontaneously and naturally throughout the experiment. To maintain body temperature within physiological range, all manipulations were conducted on a heating pad, while body temperature was continuously monitored rectally. The rats were also monitored by continuous electrocardiography.

At 0, 10, 20, 30, 40, 50 and 60 min after the onset of HS, and at 0.5, 2 and 4 h after the start of resuscitation, the entire intestine was excised and the duodenum, jejunum, ileum and colon were dissected for Northern blotting as described in Materials and methods. (A) Northern blots of total RNA (20 μg) hybridized with [α-32P]dCTP-labeled HIF-1α cDNA. Loading control, same total RNA stained with ethidium bromide. Filled arrowhead, 28S ribosomal RNA; open arrowhead, 18S ribosomal RNA; S, sham-operated control animals. Results are typical of three independent experiments. (B) Levels of HIF-1α mRNA expressed as ratios (%) of respective sham-operated controls. Filled circles, duodenum; gray circles, jejunum; shaded boxes, ileum; dotted boxes, colon. Insets: maximal levels of HIF-1α mRNA during and after HS in various regions of the intestine expressed as arbitrary densitometric units. Data are presented as the means ± SEM (n=3). *p<0.05 vs. ileum.

Figure 1. Effect of hemorrhagic shock (HS) on hypoxia-inducible factor (HIF)-1α gene expression in various intestinal regions. Rats were sacrificed at 0, 10, 20, 30, 40, 50 and 60 min after the onset of HS, and at 0.5, 2 and 4 h after the start of resuscitation. Intestines were excised and the duodenum, jejunum, ileum and colon were dissected for Northern blotting as described in Materials and methods. (A) Northern blots of total RNA (20 μg) hybridized with [α-32P]dCTP-labeled HIF-1α cDNA. Loading control, same total RNA stained with ethidium bromide. Filled arrowhead, 28S ribosomal RNA; open arrowhead, 18S ribosomal RNA; S, sham-operated control animals. Results are typical of three independent experiments. (B) Levels of HIF-1α mRNA expressed as ratios (%) of respective sham-operated controls. Filled circles, duodenum; gray circles, jejunum; shaded boxes, ileum; dotted boxes, colon. Insets: maximal levels of HIF-1α mRNA during and after HS in various regions of the intestine expressed as arbitrary densitometric units. Data are presented as the means ± SEM (n=3). *p<0.05 vs. ileum.

RNA isolation and Northern blot analysis. Total RNA isolated from rat tissues using Tri-Reagent™ (Sigma Chemical Co., St. Louis, MO) according to the manufacturer’s protocol was subjected to Northern blotting as described (8), then 20 μg of the total RNA was resolved by electrophoresis on 1.2% (w/v) agarose gels containing 6.5% (v/v) formaldehyde. Resolved bands were blotted onto a Bio-Rad Zeta-Probe™ membrane (Bio-Rad Laboratories, Richmond, CA), hybridized with [α-32P]dCTP-labeled mouse HIF-1α cDNA (9) constructed in the pBlueScript Vector (Stratagene, La Jolla, CA), then washed under stringent conditions. The membrane was exposed to radiography film (Fujiﬁlm Medical, Tokyo, Japan) with an intensifying screen at -70˚C, and autoradiographs and levels of 18S ribosomal RNA were quantified using an image scanner (GelPrint™ 2000i, Genomic Solutions, Ann Arbor, MI) and computerized image analysis software (Basic Quantifier™, version 3.0, Genomic Solutions). Relative amounts of radiolabeled cDNA that hybridized to the blots were normalized to 18S ribosomal RNA levels for loading errors.

Immunohistochemistry. Sections were immunohistochemically analyzed as described (8). Intestinal tissue was fixed in 10% neutral buffered formalin, embedded in paraffin and cut into 4- to 6-μm sections. The enzymatic activities of endogenous peroxidases in deparaffinized and dehydrated sections were blocked with 3% hydrogen peroxide. Subsequently, the sections were incubated overnight at 4˚C with a rabbit anti-
rat HIF-1α polyclonal antibody (Novus Biologicals Inc., Littleton, CO). Antigen-antibody reactions were detected using a dextran polymer reagent conjugated with peroxidase and a secondary antibody (Dako EnVision™+, DakoCytopharmaion A/S, Copenhagen, Denmark), then color was developed with 3,3'-diaminobenzidine and the sections were counterstained with Mayer’s hematoxylin. The control for non-specific staining included normal rabbit serum. The number of HIF-1α-positive cells was counted in five non-consecutive sections from each rat at a magnification of x200 by two independent investigators (H.S. and E.O.) in a blinded manner. Statistical analysis. Data were statistically evaluated by the analysis of variance followed by Scheffé’s F-test using Statview software (Abacus Concepts, Berkeley, CA). Differences were considered significant at p<0.05. Data are presented as the means ± SEM.

Results

Time courses of intestinal HIF-1α mRNA after HS. We examined levels of HO-1 mRNA in several regions of the intestine after HS. In sham-operated control animals, HIF-1α mRNA was only marginally detectable in the duodenum, jejenum, ileum and colon (Fig. 1). In contrast, HIF-1α mRNA levels were obviously increased after HS in the duodenum, jejenum and colon. Duodenal HIF-1α mRNA levels started to increase 10 min after HS, reached a maximum at 50 min, then rapidly decreased to 40% of the maximal level at 30 min after the start of resuscitation (Fig. 1). Jejunal and colonic HIF-1α mRNA levels similarly increased at 20 min after HS and reached a maximum at 30 and 40 min, respectively, then rapidly declined almost to basal levels 30 min after resuscitation (Fig. 1). Levels in the ileum hardly increased after HS. Levels of maximally induced HIF-1α mRNA in the duodenum, jejenum and colon were approximately 2.5-, 2- and 2-fold higher, respectively, than in the ileum (Fig. 1).

Intestinal HIF-1α protein expression after HS. Since Northern blot analysis revealed the expression of intestinal HIF-1α mRNA in HS animals, we examined the expression as well as the localization of HIF-1α protein in the intestinal tissue 40 min after the onset of HS by immunohistochemical analysis. Few HIF-1α-positive cells were detected in the intestines of sham-operated control animals (Fig. 2), but numbers were obviously increased in the duodenum, jejenum and colon after HS (Fig. 2). This finding was consistent with enhanced HIF-1α gene expression in HS animals. Mucosal cells emitted intense
HIF-1α signals, whereas the number of HO-1-positive cells did not increase in the ileum after HS (Fig. 2). Sections of the intestine of HS animals treated with non-immune rabbit serum were negative (data not shown).

Discussion

Our study demonstrated that HIF-1α mRNA and protein levels were increased in the intestinal mucosa of rats during the ischemic phase of HS. An important novel finding is that HIF-1α induction after HS was remarkably site-specific, in that it was obviously increased in the duodenum, jejunum and colon, but not in the ileum. These findings suggest that adaptation to hypoxic stress differs regionally in the intestine. Further studies should elucidate the mechanism and significance of such heterogeneous regiospecific induction.

Figs. 1 and 2 showed that HIF-1α mRNA and protein levels in the mucosal cells of the duodenum, jejunum and colon were rapidly and obviously increased during the early ischemic phase of HS. In contrast, these levels in the ileum were neither increased nor influenced by HS. Thus, HIF-1α expression was site-specific. Koury et al reported that ideal nuclear HIF-1α protein levels were increased after HS for 90 min plus laparotomy (10). However, their animal models underwent more serious insult than ours. Furthermore, they did not compare HIF-1α expression levels in the ileum with those of other intestinal regions, such as the duodenum, jejunum and colon. It therefore remains unclear whether ideal HIF-1α induction was greater than that in other intestinal regions.

Hemorrhage results in varying degrees of decrease in blood flow in subregions of the intestine (11). The rate of metabolism of glucose to lactate also differs distinctly depending on intestinal region (12). Thus, changes in regional blood flow and differences in cellular oxygen consumption may affect the degree of cellular hypoxia in various regions of the intestine. Moreover, previous studies have shown that tissue-specific changes in mRNA and protein synthesis are integral to survival under hypoxic/anoxic stress (13,14). Taken together, these factors might contribute to the site-specific induction of HIF-1α expression. Similar to the intestine, recent reports indicate that HIF-1α mRNA and protein expression induced by hypoxic insult is also regulated in a tissue-specific manner (4-6).

The biological significance of site-specific HIF-1α induction in the intestine after HS remains obscure. The expression of various genes that participate in cellular adaptation to hypoxia, including erythropoiesis, angiogenesis, vasodilatation and anaerobic metabolism, is regulated by HIF-1α (4). Among these genes is the one that encodes heme oxygenase-1 (HO-1) (15). The rate-limiting enzyme in heme catabolism that is also known as heat shock protein (16). The critical role played by HIF-1α in HO-1 induction during hypoxia was established based on the finding that hypoxia significantly increases the transcriptional rate of the ho-1 gene, while hypoxia-dependent HO-1 expression is abolished in mutant cells lacking HIF-1α DNA binding activity (16). We also demonstrated that HS results in the obvious induction of HO-1 in the mucosal cells of the duodenum, jejunum and colon, but not of the ileum (8), indicating that the site-specific induction and cellular localization of HIF-1α protein are entirely consistent with those of HO-1 induction after HS. Collectively, HIF-1α might be involved in the site-specific induction of HO-1 in the intestine. Thus, our findings suggest that HIF-1α induction constitutes a tissue-specific adaptive response to acute hypoxic insult by HS in the intestine.

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References