

Anti-septic effects of dabrafenib on HMGB1-mediated inflammatory responses

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A nucleosomal protein, high mobility group box 1 (HMGB1) is known to be a late mediator of sepsis. Dabrafenib is a B-Raf inhibitor and initially used for the treatment of metastatic melanoma therapy. Inhibition of HMGB1 and renewal of vascular integrity is appearing as an engaging therapeutic strategy in the administration of severe sepsis or septic shock. Here, we examined the effects of dabrafenib (DAB) on the modulation of HMGB1-mediated septic responses. DAB inhibited the release of HMGB1 and downregulated HMGB1-dependent inflammatory responses by enhancing the expressions of cell adhesion molecules (CAMs) in human endothelial cells. In addition, treatment with DAB inhibited the HMGB1 secretion by CLP and sepsis-related mortality and pulmonary injury. This study demonstrated that DAB could be alternative therapeutic options for sepsis or septic shock via the inhibition of the HMGB1 signaling pathway. [BMB Reports 2016; 49(4): 214-219]

INTRODUCTION

Sepsis is systematic inflammatory response syndrome caused by infection and can lead to shock, multiple organ failure, and death if not treated promptly (1). Sepsis is the major causes of death in intensive care units (ICUs) and the tenth leading cause of death overall in high-income countries (1), despite ad-

vances in antibiotic therapy and intensive care (2, 3). Recombinant activated protein C (APC, Xigris, Eli Lilly) was licensed in 2001 by the Food and Drug Administration (FDA) and in 2002 by the European Medicine Agency for the treatment of severe sepsis and septic shock (4). In October 2011, however, the only FDA-approved drug for severe sepsis, APC was withdrawn from the market because of side effects and PROWESS and septic shock trials showed its inability to enhance survival rate on 28-day mortality (5). Therefore, we have no drug for sepsis or septic shock, currently. Therefore, developing more effective therapeutic options for sepsis are needed.

The pathogenesis of sepsis is rather complex, tumor necrosis factor (TNF)- α , interleukin (IL)-1, interferon- γ , and macrophage migration inhibitory factor were upregulated early by endotoxin (6). For instance, TNF- α neutralizing antibodies decreased lethality of mouse endotoxemic shock model (7). In contrast, the rapid kinetics of the TNF- α responses in clinical trial made its targeting difficult (7), which drive us to find another late septic mediator such as high mobility group box 1 protein (HMGB1) as potential therapeutic targets for sepsis. In mouse models of sepsis, HMGB1 is first detectable in the circulating blood 8 h after the onset of the disease, and finally increasing to plateau levels from 16 to 32 h (8). In contrast to TNF- α and other early proinflammatory cytokines, the circulating concentrations of HMGB1, as a late acting mediator of sepsis, proceeds and is similar to the severity of sepsis (8). HMGB-1, an important chromatin protein present in almost all eukaryotic cells, is known to interact with nucleosomes, transcription factors, and histones in the nucleus (9). HMGB1 can be secreted into the extracellular space through a passive process initiated by necrotic cells, thereby allowing the innate immune system to recognize damaged cells (10). The secretion of HMGB1 in the endothelium upregulates the expression of cell adhesion molecules (CAMs), such as vascular cell-adhesion molecule (VCAM), intercellular adhesion molecule (ICAM), and E-selectin, which promote inflammation via the recruitment of leukocytes (11). Enhanced blood concentrations of HMGB1 are

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quantitative in septic patients upto7 days after a diagnosis of sepsis, and these concentrations are associated with the severity of organ damage (12, 13). Large therapeutic options for clinical intervention for sepsis treatment were provided by HMGB1, as a late mediator of sepsis (14).

In our search for repositioning FDA-approved drugs (total 1,163), 327 drugs were selected which are related to vascular inflammation and infection. Among selected drugs, high contents screening system (PerkinElmer Operetta, Waltham, MA) was used to select the compounds which control HMGB1-mediated vascular barrier disruption, and we found that dabrafenib (DAB) inhibited HMGB1- or CLP-induced septic responses. Dabrafenib (brand name; Tafinlar™) is a kinase inhibitor indicated as a single agent for the treatment of patients with unresectable or metastatic melanoma with BRAF V600E mutation as detected by an FDA-approved test (15). Here, we showed the anti-septic effects of DAB on HMGB1-mediated vascular barrier disruption *in vitro* and *in vivo*.

RESULTS AND DISCUSSION

Drug repositioning for development of new drug by using existing FDA-approved drugs has gained significant attention because of the time and cost benefits (16-18). Here we report the results of DAB on the basis of the inhibition of HMGB1-mediated septic activity *in vitro* and *in vivo*.

Effects of DAB on LPS and CLP-mediated release of HMGB1

HMGB1 is actively released by immune cells such as monocytes and macrophages in response to LPS, as well as by necrotic cells (11, 19-21). HMGB1 levels rise slowly after 8 h and are corresponded to the severity of sepsis (22). Thus, we tested whether DAB could reduce HMGB1 secretion by LPS in HUVECs. As shown in Fig. 1A, DAB reduced LPS-mediated HMGB1 release by dose-dependent manner, with a maximal effective concentration of 2 μ M. In order to confirm the inhibitory effects of DAB on HMGB1 secretion *in vivo*, DAB was evaluated for its ability to inhibit HMGB1 release using a mouse model of CLP-induced sepsis. Data showed that DAB also inhibited CLP-induced HMGB1 secretion (Fig. 1B). Because the total volume of blood in a mouse is 72 ml/kg (23) and the average weight of used mouse in this study is 27 g, the amount of DAB (260 or 520 μ g/kg) injected yielded a maximum concentration of 5, or 10 μ M in the peripheral blood. We next determined underlying molecular mechanism of the inhibitory effects of DAB on the secretion of HMGB1 on LPS-treated HUVECs. To do this, the effects of DAB on the transcriptional regulation of HMGB1 by LPS in HUVECs were tested. Thus, we determined the effect of DAB on LPS-induced HMGB1 mRNA levels using real time qRT-PCR. Data showed that LPS increased DAB treatment suppressed increased expressions of HMGB1 mRNA by LPS (Fig. 1C). We next assessed the effects of DAB and HMGB1 on the levels of the HMGB1 receptors (TLR2, TLR4, and RAGE) were evaluated.

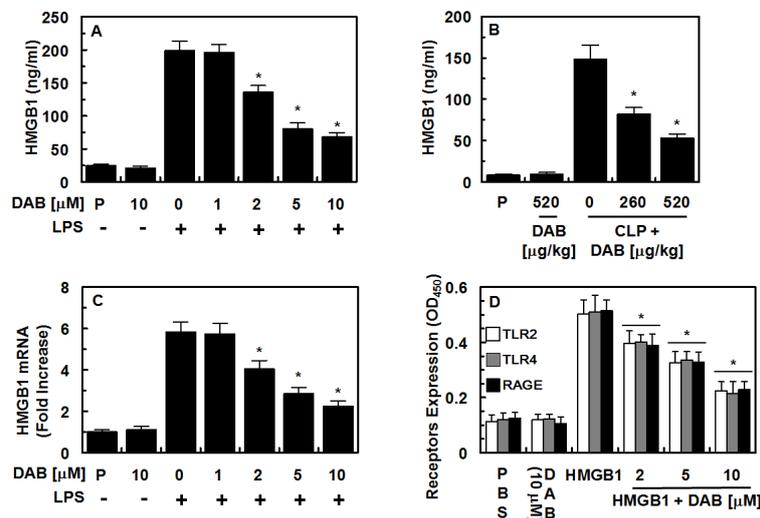


Fig. 1. Effects of DAB on HMGB1 release and expression of HMGB1 receptors. (A) HUVECs were treated with the indicated concentrations of DAB for 6 h, after stimulation with LPS (100 ng/ml, 16 h), and HMGB1 release was measured by ELISA. (B) Male C57BL/6 mice that underwent CLP were administered indicated DAB (intravenously 12 h after CLP, n = 5). Mice were euthanized 24 h after CLP. Serum HMGB1 levels were measured by ELISA. (C) The same as (A) except that real time qRT-PCR analysis was performed using specific primers for HMGB1 and GAPDH, as described in the materials and methods section. (D) Confluent HUVECs were activated with or without HMGB1 (1 μ g/ml, 16 h), followed by incubation with DAB for 6 h. Expression of TLR2 (white bar), TLR4 (gray bar) or RAGE (black bar) was determined by cell-based ELISA. The results shown are mean \pm SEM from three separate experiments in different days with triplicate wells. P = PBS is the vehicle control. *P < 0.05 versus LPS alone (A, C), CLP alone (B) or HMGB1 alone (D).

HMGB1 induced the expression of HMGB1 receptors such as TLR2, TLR4, and RAGE in HUVECs, and DAB inhibited the expressions of HMGB1 receptors (Fig. 1D). However, DAB alone did not affect the expressions of HMGB1 receptors. We next tested the effects of DAB on the cell viability. To do this, cell viability assays were applied to probe the toxicity of DAB in HUVECs after 48 h. At the concentrations used (up to 20 μ M), DAB did not affect cell viability (data not shown). Collectively, prevention of LPS- or CLP-induced release of HMGB1 by DAB indicates the possible use of DAB in the management of sepsis.

Effect of DAB on HMGB1-mediated vascular integrity

It is well known that LPS and HMGB1 could cause the cleavage and disruption of the endothelial barrier (24). Therefore, a permeability assay was applied to test whether DAB could restore LPS- or HMGB1-induced vascular disruptive responses in HUVECs. HUVECs were treated with various concentrations of DAB for 6 h following the addition of LPS (100 ng/ml) or HMGB1 (1 μ g/ml). Data showed that DAB dose-dependently inhibited LPS- and HMGB1-mediated membrane disruption (Fig. 2A and 2B). The *in vivo* effects of DAB on vascular per-

meability were assessed to corroborate the *in vitro* results. Fig. 2C shows that DAB induced a marked inhibition of the peritoneal dye leakage, induced by HMGB1. HMGB1 elicits an inflammatory response through diverse pathways, including MyD88/IRAK/TRAF, which leads to NF- κ B activation; the Rac1/PI3K pathway; via the activation of ERK 1/2 and p38 MAPK downstream of TLR2/4; and the Ras/p38 pathway downstream of RAGE (25-27). Therefore, DAB could affect the p38 pathway, HUVECs were stimulated by HMGB1 and treated with DAB. HMGB1 upregulated the expression of phosphorylated p38, which was clearly reduced by treatment with DAB (Fig. 2D). In addition, treatment with DAB (5 or 10 μ M) inhibited the formation of HMGB1-induced paracellular gaps with the formation of dense F-actin rings (Fig. 2E). However, DAB alone did not affect the formation of dense F-actin rings. The reduction in HMGB1-induced permeability and p38 activation by DAB points towards a promising role as ant-sepsis drugs.

Effects of DAB on HMGB1-mediated expression of CAMs, and adhesion/migration of human neutrophils

Previous report indicated that the expressions of CAMs (ICAM-1, VCAM-1, and E-selectin) were increased by HMGB1,

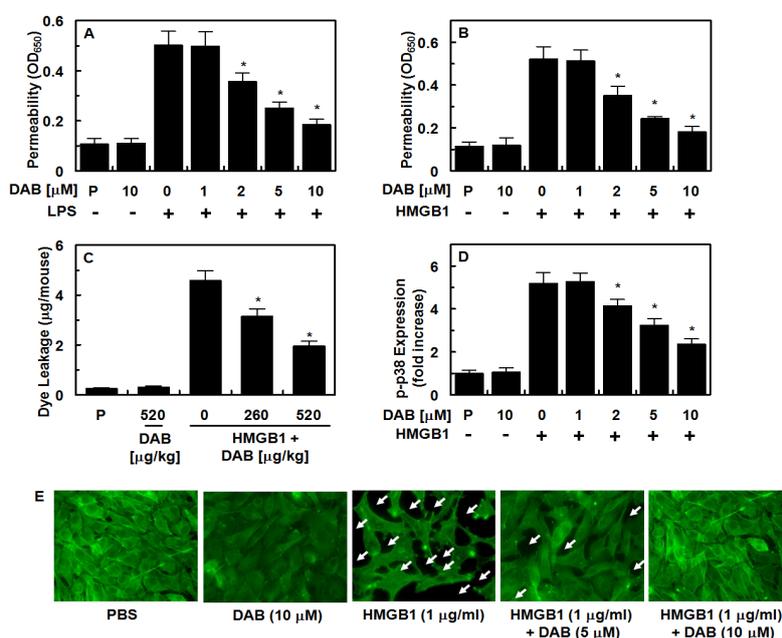


Fig. 2. Effects of DAB on HMGB1-mediated permeability *in vitro* and *in vivo*. Effects of treatment with different concentrations of DAB for 6 h on barrier disruption caused by LPS (A, 100 ng/ml, 4 h) or HMGB1 (B, 1 μ g/ml, 16 h) were monitored by measuring the flux of Evans blue-bound albumin across HUVECs. (C) The effects of DAB on HMGB1-induced (2 μ g/mouse, i.v.) vascular permeability in mice were examined by measuring the amount of Evans blue in peritoneal washings (expressed μ g/mouse, n = 5). (D) HUVECs were activated with HMGB1 (1 μ g/ml, 16 h), followed by treated with different concentrations of DAB for 6 h. The effects of DAB on HMGB1-mediated expression of phospho-p38 were determined by ELISA. (E) Staining for F-actin. HUVEC monolayers grown on glass coverslips were stimulated with or without HMGB1 for 1 h, and then treated with DAB (5 or 10 μ M) for 6 h, and stained for F-actin. Arrows indicate inter-cellular gaps. The representative images were from three separate experiments in different days with similar results. Results are expressed as the mean \pm SEM of three separate experiments in different days. *P < 0.05 versus LPS (A) or HMGB1 (B-D).

which is correlated with vascular inflammatory diseases (28). CAMs also support the behaviors of leukocytes (adhesion and transmigration toward endothelium to the site of inflammation (28). As can be seen in Fig. 3A, DAB reduced the expression of CAMs by HMGB1 signaling pathway. In addition to reduced CAM expression, DAB reduced the adherence of the human neutrophils to HUVECs and their subsequent migration (Fig. 3B-3D). These results were consistent with *in vivo* data by the suppression of HMGB1- or CLP-induced migration of leukocytes in the peritoneal space (Fig. 3E).

Effects of DAB on HMGB1-stimulated activation of NF-κB/ERK and production of IL-6/TNF-α

Vascular inflammatory mediators, such as TNF-α and IL-6, commit to progressions of the systemic severe vascular inflammatory responses and secreted by HMGB1 (29). As outlined above, HMGB1 stimulates the release of pro-inflammatory cytokines through various pathways (14, 27, 30), including ERK 1/2, ultimately leading to the activation of NF-κB and upregulation of inflammatory cytokine release. Therefore, we determined the effects of DAB on the activation of inflammatory signaling molecules and the secretion of TNF-α and IL-6, HUVECs were activated with HMGB1 for 16 h, followed by incubation with DAB for 6 h. Our results indicate that DAB significantly reduced TNF-α and IL-6 levels (Supplementary Fig. 1A and 1B). As shown in Supplementary Fig. 1C and 1D, DAB treatment (6 h) inhibited the phosphorylation of NF-κB and ERK 1/2 by HMGB1. And, we measured HMGB1-induced translocation of NF-κB from cytosol to nucleus by using p65 NF-κB and fluorescein isothiocyanate (FITC)-conjugated antibody. As a result, data (Supplementary Fig. 1E) indicated

that treatment with DAB inhibited the translocation of NF-κB p65 from cytoplasm into nucleus by HMGB1.

Protective effect of DAB in the CLP-induced septic lethality

It is well known that upregulated expressions of HMGB1 contribute the progression and severity of sepsis (14, 31). In this study, we showed the inhibitory and cytoprotective effects of DAM on HMGB1 and underlying signaling effects. This prompted us to validate the effects of DAB on the survival study of mice with sepsis following CLP surgery. To determine whether DAB protects mice from CLP-induced sepsis lethality DAB was administered to mice after CLP. A single administration of DAB (260 or 520 μg/kg, 12 h after CLP) did not prevent CLP-induced death (data not shown). Thus, we next administered same amounts of DAB twice (once 12 h after CLP and once 50 h after CLP). This increased the survival rate from 30 to 50%, according to the Kaplan-Meier survival analysis ($P < 0.00001$, Fig. 4A). The marked improvement in survival rate achieved by the administration of DAB indicated that the inhibitory effects of DAB on not only HMGB1 secretion but inflammatory responses were therapeutic strategy for management of sepsis and septic shock.

Protective effect of DAB in the CLP-induced tissues injury

We determined the effects of DAB on CLP-induced pulmonary injury to confirm the protective effects of DAB on CLP-induced death. We did not find the significant differences between the lungs of sham mice and sham mice treated with DAB (data not shown). In CLP-operating mice group, we found that the numbers of infiltrated inflammatory cells into the interstitium and alveolar spaces were increased and that the areas of damaged

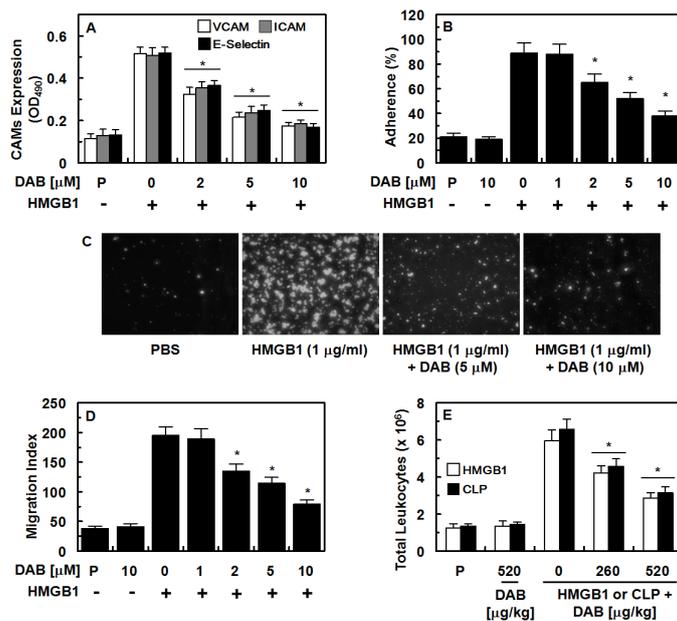


Fig. 3. Effects of DAB on HMGB1-mediated pro-inflammatory responses. (A-D) HUVECs were stimulated with HMGB1 (1 μg/ml) for 16 h, followed by treatment with DAB for 6 h. HMGB1-mediated (A) expression of VCAM-1 (white bar), ICAM-1 (gray bar), and E-selectin (black bar) in HUVECs, (B, C) adherence of human neutrophils to HUVEC monolayers, and (D) migration of human neutrophils through HUVEC monolayers were analyzed. (E) The effects of treatment with DAB on HMGB1-induced (2 μg/mouse, i.v., white box) or CLP-induced (black box) leukocyte migration into the peritoneal cavities of mice were analyzed. The representative images were from three separate experiments in different days with similar results. All results indicate the mean ± SEM of three separate experiments in different days. * $P < 0.05$ vs. HMGB1.

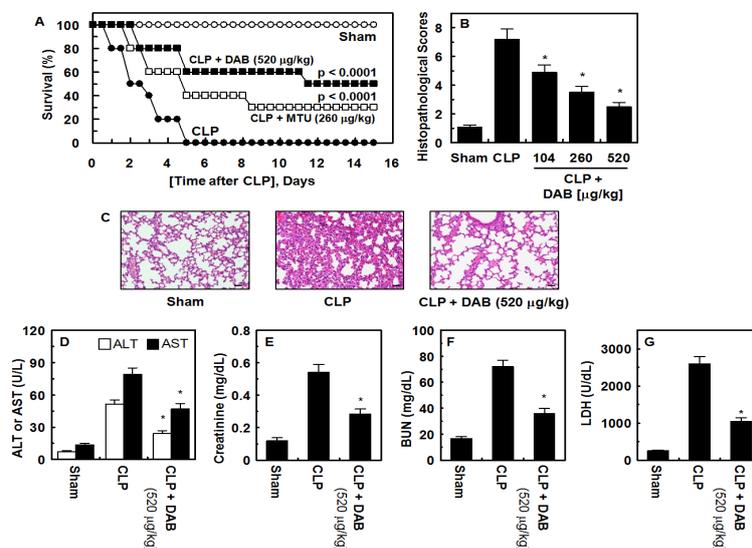


Fig. 4. Effects of DAB on lethality or pulmonary injury after CLP. (A) Male C57BL/6 mice ($n = 20$) were administered DAB at 260 $\mu\text{g}/\text{kg}$ (i.v. \square) or at 520 $\mu\text{g}/\text{kg}$ (i.v. \blacksquare) at 12 h and 50 h after CLP. Animal survival was monitored every 12 hours after CLP for 15 days. Control CLP mice (\bullet) and sham-operated mice (\circ) were administered sterile saline ($n = 20$). Kaplan-Meier survival analysis was used to determine the overall survival rates versus CLP treated mice. (B) Male C57BL/6 mice underwent CLP and were administered DAB intravenously at 12 h and 50 h after CLP ($n = 5$). Mice were euthanized 96 h after CLP. Histopathological scores for the lung tissue were recorded as described in methods. (C) Photomicrographs of lung tissues (H&E staining, $\times 200$). Sham group (grade 1); CLP group (grade 3); Right, CLP + DAB group, (grade 2). Illustrations indicate representative images from three independent experiments in different days with similar results. All results indicate the mean \pm SEM of three separate experiments in different days with similar results. (D-G) The same as (B, C) except that mice were bled and euthanized. AST, ALT, BUN, and creatinine level in plasma was measured. All results indicate the mean \pm SEM of three separate experiments. *indicates $P < 0.05$ vs. CLP.

pulmonary architecture were increased (Fig. 4B and 4C). However, DAB treatment decreased damaged pulmonary architecture and the lung injury score was decreased in CLP-operating mice (Fig. 4B and 4C). Multiple organ failure, such as liver and kidney, was found in severe vascular inflammatory conditions, sepsis (32). As shown in Fig. 4, treatment of DAB reduced CLP-induced plasma level of ALT and AST (markers of hepatic injury, Fig. 4D), and creatinine and BUN (markers of renal injury, Fig. 4E and F). DAB also reduced the general tissue damage marker, LDH, in CLP-operating mice (Fig. 4G).

The present work was undertaken to evaluate the protective effect of DAB on vascular barrier integrity. Management of vascular integrity is appearing as pivotal therapeutic strategies for the control of sepsis. HMGB1 is involved in the advancement of sepsis and associated with severe vascular inflammatory diseases (14). Here, our data indicated that the anti-septic responses of DAB occur via the suppression of HMGB1 release and HMGB1-induced vascular inflammatory responses. The molecular mechanism of anti-septic responses by DAB is attributed by the inhibition of HMGB1 secretion (Fig. 1A, 1B), the expressions of HMGB1 receptors (Fig. 1C) and HMGB1-induced vascular disruption (Fig. 3B, 3C, 3E) via suppression of the activation of p38 (Fig. 2D). In addition, DAB inhibited the adhesion and migration of leukocytes toward HUVECs by

suppression of the expressions of CAMs (Fig. 3). The inhibitory mechanism of DAB on septic condition is the downregulation of the production of inflammatory cytokines (TNF- α , IL-6, Supplementary Fig. 1A, 1B) and the activation of inflammatory transcriptional factor (NF- κ B, ERK1/2, Supplementary Fig. 1C, 1D). DAB also inhibited the translocation of NF- κ B from cytosol to nucleus (Supplementary Fig. 1E).

Taken together, the results of this study suggested that DAB inhibited both LPS- and CLP-mediated secretion of HMGB1, expressions of HMGB1 receptors, and HMGB1-mediated vascular disruption by increasing barrier integrity and suppression of the expression of CAMs. Further, DAB reduced the adhesion and migration of leukocytes toward HUVECs. The barrier protective effects of DAB were confirmed in a mouse model, in which treatment with DAB reduced CLP-induced mortality and pulmonary injury. Our findings indicate that DAB could be one of potential drug candidates for treatment of sepsis and septic shock.

MATERIALS AND METHODS

See supplementary information for this section.

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