Antibody Binding to Fas Ligand Attenuates Inflammatory Cell Infiltration and Cytokine Secretion, Leading to Reduction of Myocardial Infarct Areas and Reperfusion Injury

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SUMMARY: Fas ligand (FasL) induces apoptotic cell death when bound to Fas antigen. The engagement of FasL has anti-inflammatory effects through the prevention of cell proliferation and cytokine secretion. However, the role of FasL in myocardial ischemia/reperfusion (MI/R) injury is unclear. We examined the expression of FasL mRNA in the myocardium of MI/R rats by ligating the left coronary artery for 30 minutes and allowing reperfusion to occur for 0, 1, 3, and 24 hours. The expression of FasL mRNA was enhanced 1 hour after reperfusion, and enhanced levels were consistently seen after 24 hours of reperfusion. FasL immunostaining was observed on neutrophils, macrophages, T cells, and vascular endothelial cells. We then assessed the potential role of FasL in the cell proliferation and cytokine production seen in MI/R injury after 24 hours of reperfusion. Rats were divided into three groups; Group A, without treatment; Group B, treated with nonspecific rabbit IgG; and Group C, treated with anti-FasL antibody. Anti-FasL antibody or rabbit IgG were administered intravenously before coronary artery occlusion. In Group C, interleukin-1 β and interleukin-2 mRNA levels were decreased, and neutrophil and T cell accumulation was attenuated. The infarct area determined by triphenyltetrazolium chloride staining was significantly smaller in Group C (18 ± 4%) than in Group A (34 ± 2%) or Group B (33 ± 4%) ($\rho < 0.0001$). However, there was no significant difference in the prevalence of terminal deoxynucleotidyltransferase-mediated dUTP-biotin nick end-labeling-positive cardiomyocytes among the three groups. These findings suggest that the cardioprotective effect of anti-FasL antibody is due to its anti-inflammatory action, rather than antiapoptotic action. The Fas/FasL system may be involved in the development of MI/R injury. (*Lab Invest 2002, 82:1121–1129*).

E arly coronary artery reperfusion plays a key role in reducing myocardial tissue injury associated with acute myocardial infarction. Nevertheless, reperfusion itself can cause damage to the myocardium, termed myocardial ischemia/reperfusion (MI/R) injury (Braunwald and Kloner, 1985). In MI/R injury, several kinds of inflammatory cells, including neutrophils, macrophages, and T cells, accumulate in the reperfused tissue (Birdsall et al, 1997; Dreyer et al, 1991; Ono et al, 1999). Accumulating neutrophils may contribute to production of myocardial damage by releasing oxygen

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free radicals, proteases, and leukotrienes (Mehta et al, 1988). Macrophages elaborate proinflammatory cytokines, such as interleukin (IL)-1 β , TNF- α , and IL-6, which mediate the acute inflammatory response and subsequent remodeling after MI/R injury (Finkel et al, 1992; Herskowitz et al, 1995; Yue et al, 1998).

Fas ligand (FasL, CD95L) is a 40-kd type II membrane protein member of the TNF/nerve growth factor family (Suda et al, 1993) that induces apoptosis by binding to its membrane receptor Fas (Nagata and Golstein, 1995). In Langendorff perfusion, the Fas/ FasL system contributes to cell death in cardiac cells in response to MI/R injury (Jeremias et al, 2000). However, recent reports indicate that FasL has other important functions as well as this well-known apoptosis-promoting activity. First, FasL itself has chemotactic activity and triggers inflammation. Enforced FasL expression was found to facilitate rejection by triggering a neutrophil-mediated inflammatory response. This response was observed in a variety of cell lines and tissues, namely, islets (Allison et al, 1997; Kang et al, 1997b), myoblasts (Kang et al, 1997a), and

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tumor cells (Arai et al, 1997; Miwa et al, 1998; Seino et al, 1997). Furthermore, a soluble type of FasL (sFasL) was a potent chemoattractant for human neutrophils (Ottonello et al, 1999). These reports suggest that FasL has strong proinflammatory properties. Second, FasL itself transduces signals independently of Fas signals. FasL engagement by a recombinant form of sFas, FasFc, stopped CD4⁺T cell proliferation and cytokine secretion. When engaged by FasFc, CD4⁺T cells undergo cell cycle arrest and cell death (Desbarats et al. 1998). It is known that neutrophils, macrophages, and T cells express FasL (Kiener et al, 1997; Liles et al, 1996; Nagata, 1994). In this study, we analyzed the expression of FasL in MI/R and examined whether anti-FasL antibody can prevent inflammation and reduce the infarct size in a rat model of MI/R.

Results

Expression of FasL in Ischemic Reperfused Myocardium

Figure 1A shows representative results of RT-PCR of FasL mRNA. The expression of FasL mRNA in the reperfused myocardium was shown to increase 1 hour after reperfusion, and markedly increased levels were consistently seen after 24 hours of reperfusion (Fig. 1B). Figure 2 shows immunohistochemical staining for



Figure 1.

Fas ligand (FasL) mRNA expression in myocardial ischemia/reperfusion (MI/R) rats. A, Time course of FasL mRNA expression in the myocardium of rats subjected to 30 minutes of left coronary artery occlusion and killed 0, 1, 3, 24 hours after reperfusion. Control (*C*) represents FasL mRNA expression in the heart of a normal rat. B, Graphic representation of the time course of FasL mRNA expression normalized to the level of glyceraldehyde-3-phosphate dehydrogenase (GAPDH). The ratio of FasL/GAPDH in MI/R rats (n = 5) was higher than in control rats 1, 3, and 24 hours after reperfusion. *p < 0.0001 compared with control rats (n = 5).



Figure 2.

A

Immunohistochemical staining for FasL, ED1, and OX19 in MI/R rats. FasL was visualized with diaminobenzidine (brown) and ED1 and OX19 with Fast Blue (blue). A, Double staining using anti-FasL antibody and OX19 in reperfused rats (original magnification, \times 640). B, Double staining using anti-FasL antibody and ED1 in reperfused rats (original magnification, \times 640).

FasL, ED1, and OX19. FasL immunoreactivity was observed in neutrophils, ED1-positive inflammatory macrophages, ED2-positive resident macrophages, OX6-positive dendritic cells, OX19-positive T cells, and vascular endothelial cells (data not shown). The major populations of FasL-positive cells were neutrophils, ED1-positive macrophages, OX19-positive cells, and vascular endothelial cells 24 hours after reperfusion.

Regulation of Cytokine Expression after Administration of Anti-FasL Antibody

Figure 3 shows representative results of RT-PCR of IL-1 β , IL-2, IL-6, TNF- α , and interferon (IFN)- γ mRNA and graphical representations of the relative level of mRNA expression of each, normalized to the level of glyceraldehyde-3-phosphate dehydrogenase (GAPDH) mRNA. In the hearts of control rats, IL-1 β , TNF- α , and IFN- γ mRNA were weakly expressed, whereas IL-2 and IL-6 mRNA were not detected. Significant elevation of each cytokine was found during the 24-hour period after reperfusion, and administration of anti-FasL antibody



Figure 3.

Effect of anti-FasL antibody on IL-1 β , IL-2, IL-6, IFN- γ , and TNF- α mRNA expression. A, Cytokine mRNA expression from the myocardium 24 hours after reperfusion of untreated rats (*UT*) (n = 5), rats treated with nonspecific rabbit IgG (*IgG*) (n = 5), and rats treated with anti-FasL antibody (*Anti-FasL*) (n = 5). Control (*C*) corresponds to levels of cytokine mRNA in the hearts of normal rats. B, Relative level of IL-1 β , IL-2, IL-6, IFN- γ , and TNF- α mRNA (cytokine/GAPDH). *p < 0.05, **p < 0.01 versus untreated group and group treated with rabbit IgG.

resulted in a marked decrease of mRNA expression only in IL-1 β (p < 0.01) and IL-2 (p < 0.05).

Effect of Anti-FasL Antibody on Neutrophils, OX19-, and ED1-Positive Cell Infiltration

Rats were divided into three groups: Group A, without treatment; Group B, treated with nonspecific rabbit IgG; and Group C, treated with anti-FasL antibody. Examples of the histologic appearance of the ischemia/reperfusion (I/R) myocardium stained with naphthol AS-D chloroacetate esterase (Leider stain) from Group A and Group C are shown in Figure 4. Dense neutrophilic infiltrates were present in the I/R myocardium of rats from Group A. In contrast, the neutrophilic infiltrates in the myocardium were markedly reduced in rats from Group C. The number of neutrophils, OX19-, or ED1-positive cells, which were also predominantly immunoreactive for FasL, was calculated in each group. The numbers of neutrophils in Group A, Group

B, and Group C were 677 \pm 32 cells/mm², 680 \pm 55 cells/mm², and 397 \pm 25 cells/mm², respectively. Group C had significantly less neutrophilic infiltration than Group A and Group B (p < 0.0001). The numbers of OX19-positive cells in Group A, Group B, and Group C were 947 \pm 94 cells/mm², 931 \pm 66 cells/mm², and 755 \pm 28 cells/mm², respectively. The number of OX19-positive cells in Group C was smaller than in Group A and Group B (p < 0.05). However, the number of ED1-positive cells was similar among the three groups (Group A, 674 \pm 51 cells/mm²; Group B, 666 \pm 38 cells/mm²; Group C, 679 \pm 76 cells/mm²) (Fig. 5).

Apoptotic Index

Terminal deoxynucleotidyltransferase-mediated dUTPbiotin Nick End-labeling (TUNEL)-positive nuclei were detected in scattered myocytes in reperfused area lesions (Fig. 6). The apoptotic index in Group A, Group B,



Figure 4.

Infiltrating neutrophils stained with naphthol-ASD chloroacetate esterase 24 hours after reperfusion. Representative sections of myocardium of A, an untreated rat (original magnification, \times 640) and B, a rat treated with anti-FasL antibody (original magnification, \times 640).

and Group C was $6.7 \pm 1.7\%$, $6.8 \pm 1.0\%$, and $7.2 \pm 1.8\%$, respectively. There was no significant difference among the three groups (Fig. 7).

Effect of Anti-FasL Antibody on Myocardial Infarct Size

Figure 8 shows representative heart sections of rats from each group. Anti-FasL antibody reduced the infarct size. The ratios of infarct area to left ventricular area in Group A, Group B, and Group C were 34 \pm 2%, 33 \pm 4%, and 18 \pm 4%, respectively. The ratio in Group C was significantly smaller than in Group A and Group B ($\rho < 0.0001$) (Fig. 9).

Discussion

In this report, we clearly demonstrated a cardioprotective action of intravenous administration of anti-FasL antibody resulting from attenuation of neutrophil infiltration and cytokine secretion in a rat model of MI/R injury. It has been shown that inflammatory cells, including neutrophils, macrophages, T cells, and proinflammatory cytokines are important regulators in

reperfusion injury (Birdsall et al, 1997; Drever et al, 1991; Finkel et al, 1992; Herskowitz et al, 1995; Mehta et al, 1988; Ono et al, 1999; Zwacka et al, 1997). Neutrophils cause myocardial tissue injury through mechanical obstruction of capillary vessels, production of vasoactive substances, or release of cvtotoxic agents (Mehta et al, 1988). Macrophages and T cells also contribute to the pathogenesis of reperfusion injury (Birdsall et al, 1997; Ono et al, 1999; Zwacka et al, 1997). An important role of these cells is secretion of proinflammatory cytokines such as IL-1 β , TNF- α , IL-6. IL-2. and IFN- γ (Herskowitz et al. 1995; Nathan, 1987; Salgame et al, 1991). Proinflammatory cytokines produced by inflammatory cells promote further inflammatory cell infiltration and influence acute tissue injury and later tissue remodeling or progression of heart failure (Finkel et al, 1992; Herskowitz et al, 1995; Yue et al. 1998).

In this study, expression of FasL mRNA was enhanced in the heart after reperfusion, and FasL immunoreactivity was predominantly localized in neutrophils, macrophages, T cells, and vascular endothelial cells. Separate from its apoptotic activity, there is growing evidence that FasL has a strong proinflammatory activity. The forced expression of FasL in several cell types has been shown to induce an effusive neutrophil-mediated inflammatory response, as documented in vivo by either tissue transplant infiltration (Allison et al, 1997; Arai et al, 1997; Kang et al. 1997a. 1997b: Miwa et al. 1998: Seino et al. 1997) or neutrophil extravasation to the peritoneal cavity (Miwa et al, 1998). The neutrophil inflammatory response to membrane-bound FasL presented by tumor cells was blocked by the administration of anti-FasL antibody (Miwa et al, 1998). sFasL may be a direct chemoattractant for neutrophils. In vitro studies involving Boyden chamber migration assays suggested that the neutrophil response could be triggered by the establishment of sFasL chemoattractant gradient (Ottonello et al, 1999). Therefore, membrane-bound FasL and sFasL produced from these inflammatory cells may function as chemokines for neutrophils in the early stage of postreperfusion, and in this study, anti-FasL antibody neutralized this chemotactic activity, leading to a 41% decrease in neutrophil infiltration. Administration of anti-FasL antibody also achieved a 19% decrease in T cell infiltration and reduced IL-2 and IL-1 β mRNA expression. FasL engagement by FasFc leads to the prevention of IL-2 secretion and cell death in FasL-expressing CD4⁺T cells (Desbarats et al, 1998). Activated T cells produce IL-2 and mediate T cell proliferation and differentiation in an autocrine fashion. Therefore, FasL engagement by anti-FasL antibody may suppress IL-2 production from vanguard T cells in the myocardium, leading to suppression of T cell proliferation. In other conditions, FasL engagement suppresses T cell infiltration in the myocardium. For example, anti-FasL antibody reduced infiltration of inflammatory cells, including lymphocytes, macrophages, neutrophils, and giant cells,



Figure 5.





Figure 6.

Apoptosis was estimated with TUNEL staining (original magnification, \times 640). TUNEL-positive nuclei visualized with diaminobenzidine (brown) were detected in scattered myocytes stained with eosin.

in experimental autoimmune myocarditis mediated by CD4⁺T cells (Ishiyama et al, 1998).

Attenuation of IL-1 β and IL-2 expression may be associated with decreased numbers of inflammatory cells, because these cytokines are produced by several inflammatory cells, including macrophages (Herskowitz et al, 1995; Nathan, 1987), T cells (Salgame et al, 1991), and neutrophils (Lindemann et al, 1988). Cytokine mRNA expression, TNF- α , IL-1 β , IL-2, IL-6, and IFN- γ , was observed in rats with MI/R injury (Herskowitz et al, 1995). However, among these cyto-

and anti-FasL antibody (Anti-FasL).

kines, only IL-1 β and IL-2 mRNA were reduced in this study. Therefore, FasL engagement may suppress IL-1 β and IL-2 production in inflammatory cells, as IL-2 production from CD4⁺ T cells is suppressed by FasFc.

In our model, no significant difference in the number of ED1-positive macrophages was found between the anti-FasL antibody-treated group and the IgG-treated group. Therefore, other chemokines may mediate macrophage infiltration in the reperfused myocardium. Monocyte chemoattractant protein-1 (MCP-1) is an important mediator of monocyte recruitment to inflammatory sites (Robinson et al, 1989; Rollins et al, 1989). MCP-1 mRNA is consistently produced in the myo-



Figure 8.

Macroscopic view of the sliced heart after 30 minutes of ischemia and then by 24 hours of reperfusion. A, Infarcted heart without treatment. B, Infarcted heart treated with IgG. C, Infarcted heart treated with anti-FasL antibody.

cardium, and its expression is augmented by MI/R (Ono et al, 1999). Anti-MCP-1 antibody decreased infiltration of macrophages and consequently reduced infarct size in MI/R rats, indicating that MCP-1 may play an important role in macrophage recruitment and myocyte injury in the reperfused myocardium.

Administration of anti-FasL antibody achieved a 46% decrease in the size of the infarct area. This effect of anti-FasL antibody was specific, because an equivalent dosage of normal rabbit IgG had no effect on the suppression of myocardial damage. Recent



Figure 9.

Effect of anti-FasL antibody on myocardial infarct size as a percentage of the total left ventricular area (n = 8). **p < 0.0001 versus heart from untreated rats (n = 8) and rats treated with rabbit IgG (n = 8).

studies have demonstrated that I/R induces apoptosis in the reperfused myocardium, and this might be important for delayed cardiomyocyte death (Gottlieb et al, 1997). A caspase inhibitor attenuated apoptosis of cardiac myocytes, resulting in a decrease of infarct size (Yaoita et al, 1998). One of the main mechanisms of caspase activation has been shown to involve the release of cytochrome c from the mitochondria to the cytosol, and another important pathway is through a mitochondrion-independent mechanism that uses cell death receptors (eq, Fas and TNF receptor). sFasL is released by the postischemic hearts early after the onset of reperfusion, and isolated hearts from Fasdeficient mice display reduction in cell death after MI/R injury (Jeremias et al, 2000). Therefore, Fas might be directly involved in cell death after myocardial ischemia. In this study, we assessed apoptotic myocardial cell death with TUNEL staining, which is widely used to detect apoptosis in various tissues. The percentage of TUNEL-positive cardiomyocytes (the apoptotic index) of Group A, B, and C were 6.7 \pm 1.7%, $6.8 \pm 1.0\%$, and $7.2 \pm 1.8\%$, respectively. There was no significant difference among the three groups. The results of this study suggest that the cardioprotective effect of anti-FasL antibody is due to its antiinflammatory action rather than antiapoptotic action in our model. Our results seem to contradict the study of Jeremias et al (2000) with regard to the contribution of apoptotic cell death. Therefore, further studies will be conducted to assess the cardioprotective effect of anti-FasL antibody in MI/R injury.

In conclusion, we have demonstrated that administration of anti-FasL antibody significantly reduces myocardial infarct size in a rat MI/R model. This cardioprotective effect could be attributed to reduced neutrophils and T-lymphocyte accumulation and attenuation of cytokine secretion in the reperfused myocardium. These results suggest that FasL plays an important role in inflammatory states in the I/R injury. Anti-FasL antibody might represent a potential therapeutic agent for I/R injury, not only in the myocardium, but also in other organs.

Materials and Methods

Experimental Ischemia/Reperfusion Model

This study was approved by the Animal Care Committee in accordance with Chiba University policies. Adult male Sprague-Dawley rats (250 to 300 g), purchased from Charles River Japan, Inc. (Kanagawa, Japan), were anesthetized with sodium pentobarbital (30 mg/ kg) by intraperitoneal injection. Rats were then intubated and ventilated with room air. Myocardial ischemia was produced by rapidly exteriorizing the heart and placing a 4.0 silk suture around the left coronary artery, approximately 2 mm from its origin. Ischemia was confirmed through elevation of the ST segment on an electrocardiogram and by change in the color of the myocardium. After 30 minutes of ischemia, the suture was removed, initiating reperfusion. The chest was then closed, pneumothorax was reduced by negative pressure, and the rats were returned to their cages.

RT-PCR

Total RNA was extracted from hearts of MI/R rats on 0, 1, 3, and 24 hours after reperfusion and from shamoperated rats, using the Qiagen RNeasy MIDI kit (Qiagen, Hilden, Germany), and was used for firststrand cDNA synthesis with random hexamer oligonucleotides (pd(N)₆; Pharmacia P-L Biochemical, Milwaukee, Wisconsin) and M-MLV reverse transcriptase (Life Technologies, Gaithersburg, Maryland), according to a standard protocol. Two micrograms of cDNA reaction mixture was used as a template for PCR. IL-1 β , IL-2, II-6, TNF- α , IFN- γ , FasL, and GAPDH mRNA were detected with primers and PCR conditions used previously (Damoiseaux et al, 1998; Herskowitz et al, 1995; Muschen et al, 1999).

The PCR reaction was carried out at three different cycle numbers to ensure it was performing in the linear range of amplification. RT-PCR products were electrophoretically separated on 3% agarose gels and visualized by ethidium bromide staining under ultraviolet transillumination. Densitometric analysis of bands was performed using NIH Image 1.59 software program (National Institutes of Health, Bethesda, Maryland). Each band was normalized to the relevant GAPDH control.

Antibody Treatment and Determination of Infarct Size

Rats were randomly divided into three groups; Group A (n = 8), without treatment; Group B (n = 8), treated with nonspecific rabbit IgG (0.1 mg/kg); and Group C (n = 8), treated with anti-FasL antibody (0.1 mg/kg; anti-P5, rabbit IgG) (Hakuno et al, 1996). Anti-FasL antibody or rabbit IgG was administered via the tail vein 1 hour before coronary artery occlusion. Twenty-four hours after reperfusion, rats were anesthetized with sodium pentobarbital (30 mg/kg, intraperitoneal injection), and the chest was reopened. The heart was excised, and the ventricle was cut into six transverse slices from the apex to the base. The slices were

incubated for 10 minutes in 1% triphenyltetrazolium chloride (TTC) solution at 37° C. Regions that failed to demonstrate brick-red staining with 1% TTC solution were considered to be infarcted myocardium. The slices were imaged by a color camera, and the images were stored on a Macintosh computer and analyzed by the NIH Image 1.59 software program. The results were expressed as the percentage of infarcted area to total left ventricular area. Heart tissues were fixed in 10% phosphate buffered formalin and embedded in paraffin for histologic studies.

Leider Stain for Neutrophils

Representative $4-\mu m$ thick paraffin-embedded sections were treated with the Leider stain using the naphthol AS-D chloroacetate esterase kit (Sigma, St. Louis, Missouri) in a standard manner to identify neutrophils in the I/R myocardium. The number of neutrophils in the I/R myocardium 24 hours after reperfusion was calculated from randomly selected fields, using CAS 200 (Becton Dickinson, San Jose, California). The resultant data were expressed as mean number \pm standard deviation per square millimeter.

Immunohistochemistry

Rat hearts were excised 24 hours after reperfusion. Freshly dissected left ventricles were frozen in optimal cutting temperature compound (OCT Compound: Miles Laboratories, Naperville, Illinois) in liquid nitrogen and stored at -80° C. Cryostat sections were cut sequentially at 4 μ m and fixed in acetone for 10 minutes. To identify the inflammatory cells, the slides were incubated with a mouse monoclonal antibody against OX6 (Serotec, Oxford, United Kingdom) for recognizing major histocompatibility complex (MHC) class II-expressing cells, including dendritic cells, some macrophages, and B lymphocytes; ED1 (PharMingen, San Diego, California), a marker for inflammatory macrophages; ED2 (BioSource International, Camarillo, California), a marker for resident macrophages; and OX19 (Serotec), a marker for pan-T cells (Ishiyama et al, 1997). After washing, the streptavidinbiotin complex method was performed according to a standard protocol for the Histfine kit from Nichirei Co. (Tokyo, Japan), and labeling was visualized with diaminobenzidine and hydrogen peroxide. The number of inflammatory cells was counted using a CAS 200 in the I/R myocardium and expressed as mean number ± standard deviation per square millimeter. Double staining using rabbit anti-rat FasL antibody and ED1, ED2, OX6, or OX19 was performed using a modification of Nakane's method (Nakane, 1968). Cytokines were localized by the streptavidin-biotin complex method. The antibodies were removed from the sections by glycine-hydrochloric acid buffer, pH 2.2, leaving the colored reaction product of diaminobenzidine. The second antigen was localized similarly, using the Fast Blue Kit (Nichirei) that develops reaction products of a color different from diaminobenzidine.

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TUNEL Assay

The TUNEL assay was performed using the ApopTag Kit (Oncor, Gaithersburg, Maryland). In brief, cryostat sections were fixed in 1% paraformaldehyde and permeabilized with the solvent (ethanol:acetic acid, 2:1) for 5 minutes at -20° C. Endogenous peroxidase was inactivated by incubation with 3% H₂O₂. Terminal deoxynucleotidyltransferase and digoxigenin-11dUTP were added to the sections according to the instructions provided by the manufacturer. The incorporated nucleotides were optimized for antidigoxigenin antibody binding. Nuclei were counterstained with hematoxylin. The sections were then stained with eosin to identify myocytes. Normal rat testis was used as the control tissue. The percentages of TUNELpositive nuclei of cardiomyocytes (apoptotic index) were calculated by counting 500 nuclei of myocytes at a high magnification (×1,000) per tissue per rat 3 hours after reperfusion.

Statistical Analysis

Data were expressed as mean \pm standard deviation. The values were tested by one-way ANOVA, and then by Scheffé's multiple comparison test. In all analyses, findings of p < 0.05 were considered statistically significant.

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