Functional, Cellular, and Molecular Characterization of the Angiogenic Response to Chronic Myocardial Ischemia in Diabetes

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Background—Ischemic heart disease is the most common cause of mortality in diabetic patients. Although therapeutic angiogenesis is an attractive option for these patients, they appear to have reduced collateral formation in response to myocardial ischemia. The aims of this study were to establish a large animal model of diabetes and chronic myocardial ischemia, evaluate the effects of diabetes on the angiogenic response, and elucidate the molecular pathways involved.

Methods and Results—Diabetes was induced in male Yucatan miniswine using a pancreatic β-cell specific toxin, alloxan (150 mg/kg; n = 8). Age-matched swine served as controls (n = 8). Eight weeks after induction, chronic ischemia was induced by ameroid constrictor placement around the circumflex coronary artery. Myocardial perfusion and function were assessed at 3 and 7 weeks after ameroid placement using isotope-labeled microspheres. Endothelial cell density and myocardial expression of angiogenic mediators was evaluated. Diabetic animals exhibited significant endothelial dysfunction. Collateral dependent perfusion and LV function were significantly impaired in diabetic animals. Diabetic animals also demonstrated reduced endothelial cell density (173 ± 14 versus 234 ± 23 cells/hpf, P = 0.03). Expression of VEGF, Ang-1, and Tie-2 was reduced, whereas antiangiogenic proteins, angiostatin (4.4 ± 0.9-fold increase, P < 0.001), and endostatin (2.9 ± 0.4-fold increase, P = 0.03) were significantly elevated in the diabetic myocardium.

Conclusions—Diabetes results in a profound impairment in the myocardial angiogenic response to chronic ischemia. Pro- and antiangiogenic mediators identified in this study offer novel targets for the modulation of the angiogenic response in diabetes. (Circulation. 2007;116[suppl I]:I-31–I-37.)

Key Words: diabetes ■ angiogenesis ■ endothelium ■ ischemia ■ growth factors

More than 35 million people in the United States are affected by diabetes and glucose intolerance. These individuals carry up to 8 times the risk of cardiovascular events compared with nondiabetic individuals, making cardiovascular disease the largest cause of mortality in this population.1 Diabetic patients experience accelerated atherosclerosis and also exhibit a diminished angiogenic response to myocardial ischemia as shown angiographically2 and in autopsy studies.3 Although, growth factor or cell-based angiogenic therapies are an attractive therapeutic option for these patients, a better understanding of pro- and antiangiogenic influences in diabetic patients is crucial to maximize the effects of such therapies.

Various mechanisms have been postulated to explain the impaired angiogenic response in diabetes. First, the presence of vascular dysfunction, characterized by both endothelial and vascular smooth muscle impairments, is a well-documented phenomenon in this setting.4–7 Second, exposure to chronic hyperglycemia leads to the nonenzymatic glycation of proteins and in particular, glycation of the extracellular matrix has been shown to impair the formation of new blood vessels.8,9 Lastly, the presence of diabetes is associated with abnormalities in growth factor signaling which have not been well characterized, particularly in the myocardium.10,11

An important limitation to the study of myocardial angiogenesis in the setting of diabetes is the lack of a validated large animal model that captures the functional, microcirculatory, and molecular abnormalities associated with diabetes, and is suitable for the induction of chronic myocardial ischemia. Although murine and rodent hind-limb ischemia models have been used to study angiogenesis, their applicability to the setting of myocardial angiogenesis is limited. The aims of this study were to (1) develop a large animal model of diabetes and chronic myocardial ischemia, (2) examine the effects of diabetes on the coronary microcirculation, myocardial angiogenesis, and myocardial function in response to chronic ischemia, and (3) to elucidate the molecular pathways involved. The aims of this study were to (1) develop a large animal model of diabetes and chronic myocardial ischemia, (2) examine the effects of diabetes on the coronary microcirculation, myocardial angiogenesis, and myocardial function in response to chronic ischemia, and (3) to elucidate the molecular pathways involved.
involved with the goal of finding novel targets to modulate the myocardial angiogenic response in diabetes.

**Methods**

**General Experimental Sequence and Diabetes Induction**

Sixteen Yucatan miniswine of male sex (Sinclair Research Inc, Colombia, Mo) were used for the studies. Diabetes was induced when animals were 8 months old using a single intravenous injection of alloxan (150 mg/kg). Only animals that achieved and maintained blood glucose levels greater than 200 mg/dL were included in the diabetic group. Animals with blood glucose levels exceeding 400 mg/dL were treated with daily insulin (70% amorphous, 30% crystalline) given intramuscularly to maintain blood glucose levels between 250 and 400 mg/dL. Fasting blood glucose was monitored every 2 days early after diabetes induction and weekly thereafter by obtaining a small blood sample (50 μL) from a capillary bed in the ear. Diabetes was maintained for 8 weeks before surgical instrumentation. Age matched miniswine (n=8) served as controls.

All animals underwent an identical experimental protocol involving 3 separate procedures on each animal. Anesthesia was performed as reported previously, and animals received humane care in compliance with the Harvard Medical Area Institutional Animal Care and Use Committee and the National Research Council’s *Guide for the Care and Use of Laboratory Animals*, prepared by the Institute of Laboratory Animals and published by the National Institutes of Health (NIH publication No. 5377-3 1996). Briefly, for all surgical procedures, anesthesia was induced with ketamine (10 mg/kg IM), thiopental (5 to 10 mg/kg IV), and thiopental 2.5% and isoflurane at 0.75 to 3.0%. The animals were intubated and mechanically ventilated at 12 to 20 breaths/min.

The first procedure, performed via small left anterolateral thoracotomy at 8 weeks after diabetes induction, consisted of the placement of a 1.75-mm ameroid constrictor around the proximal circumflex artery during the second procedure. After euthanasia, 10 circumferential, transmural left ventricular sections were collected for ILM assays in each animal, weighed, and dried. Each sample was exposed to neutron beams and microsphere densities measured in a γ counter. Adjusted myocardial blood flow (at rest and with pacing), reflecting changes in lateral myocardial perfusion, was determined from the 2 myocardial samples which showed the lowest count of gold microspheres by using the following equations:

\[
\text{Adjusted blood flow} = \frac{\text{Crude blood flow (tissue sample)}}{\text{Reference blood sample}}
\]

**Assessment of Myocardial Perfusion**

Myocardial perfusion was assessed during each procedure with isotope-labeled microspheres (ILM; Biophysics Assay Laboratories) using methods previously reported. Isotope-labeled microspheres, 15 μm in diameter, of different isotopic mass were used at each experimental stage. Gold-labeled microspheres were injected during temporary circumflex occlusion at the time of ameroid placement to identify myocardial samples that originated from the circumflex coronary distribution (those with the lowest count of gold-labeled microspheres). Lutetium and Europium-labeled ILM were used during the second procedure to determine baseline blood flow at rest and with atrial pacing at 150 bpm. Samarium and Lanthanum-labeled ILM were injected at rest and during pacing during the third procedure. Reference blood samples were obtained from the femoral artery during the second and third procedures. After euthanasia, 10 circumferential, transmural left ventricular sections were collected for ILM assays in each animal, weighed, and dried. Each sample was exposed to neutron beams and microsphere densities measured in a γ counter. Adjusted myocardial blood flow (at rest and with pacing), reflecting changes in lateral myocardial perfusion, was determined from the 2 myocardial samples which showed the lowest count of gold microspheres by using the following equations:

\[
\text{Adjusted blood flow} = \frac{\text{Crude blood flow (3rd Surgery)}}{\text{Crude blood flow at baseline (2nd Surgery)}}
\]

**Immunohistochemistry**

Myocardial sections from the circumflex territory of control and diabetic animals were stained with anti–PECAM-1 (CD-31) antibody diluted to 1:600 (BD Biosciences) as previously described. The sections were counterstained with methyl green, and examined in isolated organ chambers, as described previously. The responses to sodium nitroprusside (SNP; 1 nmol/L to 100 μmol/L), an endothelin-independent cGMP-mediated vasodilator, as well as adenosine 5’-diphosphate (ADP; 1 nmol/L to 10 μmol/L), substance P (1 fmol/L to 1 nmol/L), and VEGF (1 fmol/L to 1 nmol/L), 3 endothelin-dependent receptor-mediated vasodilators that act via bioavailable NO, were studied after precontraction by 20% to 50% of the baseline diameter and cell density in a triplicate, blinded fashion from 600×440 μm (0.264 mm²) cross-sectional fields randomly selected from the center of circumflex territories.

Whole-cell lysates were isolated from the homogenized myocardial samples with a RIPA buffer (Boston Bioproducts) and centrifuged at 10 000 g for 10 minutes at 4°C to separate soluble from insoluble fractions. Protein concentration was measured spectrophotometrically at a 595-nm wavelength with a DC protein assay kit (Bio-Rad). Forty to 80 micrograms of total protein were fractionated by 4% to 20% gradient, SDS polyacrylamide gel electrophoresis (Invitrogen) and transferred to PVDF membranes (Millipore). Each membrane was incubated with specific antibodies as follows: anti-VEGF antibody (dilution 1:250; Calbiochem), anti-eNOS antibody (1:2500;...
BD Bioscience), anti–phospho-Akt (Ser473) antibody (1:1000), anti-Akt antibody (1:1000), anti–phospho-eNOS (Ser1177) antibody (1:1000; Cell Signaling), anti–Tie-2 antibody (1:200; Santa Cruz), anti–endothelin antibody (1:1000; Upstate), anti–angiostatin antibody. Then the membranes were incubated for 1 hour in diluted appropriate secondary antibody (Jackson Immunolab). Immune complexes were visualized with the enhanced chemiluminescence detection system (Amersham). Bands were quantified by densitometry of radioautograph films.

Data Analysis
Data are reported as means±SEM. Microvessel responses are expressed as percent relaxation of the preconstricted diameter and were analyzed using 2-way repeated measures analysis of variance examining the relationship between vessel relaxation, log concentration of the vasoactive agent of interest, and the experimental group (SAS Version 9.1). Immunoblots are expressed as a ratio of protein to loading band density and were analyzed after digitization and quantification of x-ray films with ImageJ 1.33 (National Institutes of Health, USA). Blots and ILM data were analyzed with 2-tailed analyses of variance. Bonferroni corrections were applied to multiple tests, and probability values of less than 0.05 were considered statistically significant.

Statement of Responsibility
The authors had full access to the data and take full responsibility for its integrity. All authors have read and agree to the manuscript as written.

Results
Experimental Model
Diabetes was successfully induced in 8 of 10 miniswine (80%). Four animals required small doses of intramuscular insulin (10 to 20 u/d) to keep blood glucose levels below 400 mg/dL at certain points during the experimental period. Blood glucose levels were 319±31 mg/dL in diabetic animals compared with 69±2 mg/dL in control animals (P<0.001). All animals survived the entire experimental protocol. Gross examination of harvested tissues did not show any evidence of myocardial hypertrophy, fibrosis, or scarring. Histological examination, using hematoxin and eosin staining, did not reveal any major abnormalities.

Coronary Microvessel Reactivity
Results of microvessel relaxation studies are summarized in Figure 1. U46619 was used at a dose of 1×10⁻⁶ M to preconstrict coronary microvessels by an average of 40±4% of the baseline diameter. Compared with the control group, diabetic animals demonstrated mild impairment in microvessel relaxation to endothelium-dependent vasorelaxants, ADP, and substance P (both P<0.001) in the ischemic territory, suggesting reduced NO bioavailability. Furthermore, microvessel relaxation in response to VEGF was markedly impaired in diabetic animals (P<0.001). Endothelium-independent relaxation, in response to sodium nitroprusside, was similar between groups (P=0.13).

Myocardial Perfusion
Crude circumflex territory myocardial blood flow was similar between control and diabetic animals at 3 weeks after ameroid placement, both at rest (0.43±0.07 versus 0.33±0.02 mL/min/g respectively, P=0.14) and with pacing (0.34±0.04 versus 0.39±0.03 mL/min/g, P=0.39; Figure 2A). At 7 weeks, whereas the control group had an increase in circumflex territory perfusion at rest (+0.23±0.07 mL/min/g), diabetic animals demonstrated a reduction in perfusion (−0.18±0.02 mL/min/g; P<0.001). These differences were also evident during rapid atrial pacing with the baseline-adjusted circumflex flow significantly higher in the control group (+0.21±0.05 versus −0.22±0.04 mL/min/g; P<0.001; Figure 2B).
Myocardial Function
Global left ventricular function was measured at the time of the final procedure, 7 weeks after ameroid placement, and systolic and diastolic function were quantified using maximum positive dP/dt (first derivative of LV pressure) and minimum negative dP/dt, respectively. Diabetic animals demonstrated reduced LV contractility as well as impaired LV relaxation (both \( P < 0.05 \); Figure 3).

Capillary Endothelial Cell Density
Figure 4 shows the density of CD31+ capillary endothelial cells (cells/high power field) in the ischemic territory of pigs from both groups 7 weeks after ameroid placement. Diabetic animals had significantly reduced endothelial cell density compared with controls (173±14 versus 234±23; \( P = 0.03 \)).

Western Blotting
To elucidate molecular mechanisms for the impaired angiogenic response in diabetes, we examined the expression of pro- and antiangiogenic mediators in the myocardium of diabetic and control animals.

Proangiogenic Mediators: VEGF and Angiopoietin Pathways
Vascular endothelial growth factor and angiopoiein-1 are important proangiogenic growth factors that mediate their effects through binding to their respective receptors, flk-1 and Tie-2, leading to the downstream activation of Akt and endothelial nitric oxide synthase (eNOS; Figure 5A). The results of the Western blots and the densitometric analyses of proangiogenic mediators involved in the VEGF and Ang-1 pathway are summarized in Figure 5B and 5C. The expression of VEGF was markedly reduced in the diabetic myocardium compared with controls (−59±8%; \( P < 0.001 \)), whereas the difference in Ang-1 expression was not significant (\( P = 0.14 \)). Using Western blotting techniques, we were unable to measure the levels of flk-1. Tie-2 receptor expression, however, was significantly reduced in diabetic animals (−33±7%; \( P < 0.001 \)). Akt expression and phosphorylation as well as eNOS expression was similar between groups, whereas eNOS phosphorylation was significantly reduced in diabetic animals (−55±15%, \( P = 0.008 \)). These findings suggest that VEGF and Ang-1 signaling is affected at multiple levels in the diabetic myocardium.

Antiangiogenic Mediators: Angiostatin and Endostatin
Figure 6 displays the results of Western blots of antiangiogenic mediators, endostatin and angiostatin. The expression of both endostatin (2.9±0.4-fold, \( P = 0.03 \)) and angiostatin...
(4.4±0.9-fold, \(P<0.001\)) was profoundly increased in the myocardium of diabetic animals. Expression of these antiangiogenic proteins was similar between the ischemic circumflex and the nonischemic LAD territories.

### Discussion

The presence of diabetes is associated with accelerated coronary artery disease as well as alterations in the angiogenic response to ischemia. The study of the myocardial angiogenic response in diabetes is limited, in large part, because of the unavailability of a clinically relevant large animal model of chronic myocardial ischemia that captures the functional, microvascular, and molecular abnormalities associated with diabetes. In this study, we report the successful development of a porcine model of alloxan-induced diabetes and chronic myocardial ischemia, with a high rate of successful induction (80%) and maintenance of diabetes as well as excellent animal retention throughout the experimental protocol. We found that a 15-week exposure to diabetes resulted in coronary endothelial dysfunction, as shown by impaired coronary microvessel relaxation to ADP and substance P as well as a profound impairment in VEGF signaling in the coronary microvasculature. Diabetic animals also demonstrated a significantly impaired angiogenic response to

![Figure 4. Endothelial cell density. CD31+ endothelial cell density per high power field (0.264 mm²) was determined in the ischemic territory of control and diabetic animals. Diabetic animals had reduced endothelial cell density compared with controls (234±23 versus 173±14; \(^*P=0.03\)).](image)

![Figure 5. Myocardial expression of proangiogenic mediators. Expression of various proteins involved in VEGF and angiopoietin-1 signaling (A) was assessed using Western blotting. Diabetic animals had reduced myocardial expression of VEGF and Tie-2, but Ang-1 expression was not significantly different between groups (\(P=0.14\)). Akt expression and phosphorylation and eNOS expression was similar between diabetic and control animals, but eNOS phosphorylation was markedly reduced in the diabetic myocardium. VEGF indicates vascular endothelial growth factor; eNOS, endothelial nitric oxide synthase. \(^*P<0.01\).](image)
chronic myocardial ischemia. This was observed functionally as reduced perfusion of the collateral-dependent circumflex territory and morphologically as reduced endothelial cell density in the ischemic myocardium of diabetic animals. The reduction in the angiogenic response was associated with systolic and diastolic left ventricular dysfunction. Lastly, these functional alterations were associated with increased expression of antiangiogenic proteins, angiostatin and endostatin, as well as alterations in VEGF and Ang-1 signaling.

In summary, this study provides an in-depth characterization of the myocardial angiogenic response in diabetes at the functional, cellular, and molecular levels, and identifies molecular targets for the modulation of the myocardial angiogenic response in diabetes.

Endothelial dysfunction in the presence of diabetes has been demonstrated in various murine, rodent, and swine models as well as in coronary and noncoronary vasculature of humans with type I and type II diabetes. The majority of this information has been derived from noninvasive studies of the human forearm circulation or using rodent models of diabetes. As such, little is known about the effects of diabetes on the coronary microcirculation. In this study, diabetic animals demonstrated impaired coronary microvessel relaxation to ADP and substance P whereas relaxation to endothelin-independent vasorelaxant, SNP, was similar between groups, suggesting endothelial dysfunction and reduced NO bioavailability. Furthermore, the reduction in microvessel relaxation was greatly exaggerated in response to VEGF, suggesting impairment in VEGF signaling in addition to reduced NO bioavailability.

In addition to microvascular dysfunction, the presence of diabetes has been associated with a number of abnormalities that can impair the angiogenic response to myocardial ischemia. First, exposure to chronic hyperglycemia leads to the nonenzymatic glycation of protein leading to the formation of advanced glycation end products (AGEs). In vitro studies have demonstrated a diminished vascular tube formation in response to growth factors in a glycated collagen matrix. In a hindlimb ischemia model, Tamarat et al demonstrated reduced endogenous angiogenesis in diabetic mice which was reversed by the administration of aminoguanidine, an inhibitor of AGE formation.

Second, diabetes is associated with alterations in proangiogenic growth factor signaling. A number of studies have demonstrated alterations in the expression of various growth factors in plasma and noncardiac tissue in the setting of diabetes. Sasso et al studied the expression of VEGF and its downstream mediators in myocardial biopsies of patients with or without type II diabetes and found that although VEGF expression was increased, that VEGF receptor activation and downstream signaling was reduced. In concordance with this study, we also found that diabetes was associated with impaired growth factor factor signaling, which was manifest as reduced Tie-2 expression and decreased eNOS phosphorylation. Contrary to the above study, however, VEGF expression was reduced in the myocardium of diabetic swine and Ang-1 expression was similar between groups. The differences in VEGF expression between the 2 studies may be attributable to differences in the nature of diabetic disease as well as different treatments used in the patients examined.

Lastly, there is little available information on the role of antiangiogenic proteins in the context of diabetes. Angiostatin and endostatin, cleavage products of plasminogen and collagen XVIII, respectively, are formed through the proteolytic actions of various matrix metalloproteinases, and are potent antiangiogenic proteins. Weirauch et al demonstrated reduced collateral-dependent perfusion in chronically instrumented dogs under hyperglycemic conditions and demonstrated the critical role of angiostatin in inhibiting collateral vessel formation in response to repetitive coronary occlusion, both in vivo and in vitro. We have reproduced some of their findings in a clinically relevant model of diabetes and chronic myocardial ischemia by demonstrating a profound (>4-fold) increase in angiostatin expression in diabetic animals. In addition, diabetes was also associated with a 2-fold increase in endostatin expression. Inhibition of these antiangiogenic proteins may represent a novel target for the modulation and enhancement of the angiogenic response in diabetes.

Limitations
Our model of alloxan-induced diabetes largely mimics type I diabetes, ie, hypoinsulinemic hyperglycemia. Although various aspects of the disease are common between type I and type II diabetes including chronic exposure to hyperglycemia, endothelial dysfunction, increased oxidative stress and inflammation, accelerated atherosclerosis, and diminished angiogenesis, other features may be different and may limit the generalizability of this model to type II diabetes and insulin
resistance. Furthermore, this model exposes the animals to a relatively short duration (15 weeks) of severe hyperglycemia compared with the very long durations (many years) of mild to moderate hyperglycemia that diabetic patients experience. Lastly, although this model of chronic ischemia provides physiologically relevant measures of perfusion and function, it is limited by the fact that molecular markers were only assessed at a single time point, 7 weeks after ameroid placement, and therefore, acute changes in pro- and antiangiogenic mediators may not be adequately captured in this model.

Conclusions

Diabetes results in a profound impairment in endogenous myocardial angiogenesis in response to chronic ischemia. The associated alterations in growth factor signaling and increased expression of antiangiogenic mediators identified in this study may represent novel targets for the modulation of the diabetic angiogenic response. Finally, the establishment of this model will allow for the preclinical evaluation of growth factor or cell-based angiogenic therapies as well as effects of angiogenic modulating agents eg, insulin, l-arginine, statins, etc that may have therapeutic potential in patients with diabetes.

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Disclosures

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