

Interaction between expression of angiogenic factors and pathology of the microvasculature and cardiomyocytes in myocardial tissue of patients with diabetes mellitus

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Abstract

Sudden death with myocardial infarction has always been a challenging issue for the investigators and forensic pathologists. When a person suffering from angina or myocardial infarction is simultaneously suffering from diabetes mellitus issue becomes even more complex for the investigators as usual signs & symptoms of MI may not manifest so as to rouse a suspicion of MI. This study will help the pathologists to understand the microscopic changes of diabetic cardiomyopathy better in a case having MI with diabetes. This study was done in 47 cases to know the pathology of microvasculature and cardiomyocytes in myocardial tissue of diabetic patients and expression of angiogenic factors.

Keywords: Sudden death, diabetes mellitus, myocardial infarction.

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Introduction

The major cause of both morbidity and mortality in diabetic patients is cardiovascular disease. Diabetic cardiomyopathy is proposed to affect up to 50% of patients with Type 2 diabetes¹⁻⁴. Light and electron microscopic ultra structural changes of the cardiomyocyte have been quantified in a range of animal models of diabetes including the dog⁵, monkey⁶, rabbit⁷, mouse⁸ and rat⁹⁻¹². However, widely varying results have been reported with regard to the onset, type and severity of cardiomyocyte pathology¹⁰. Studies in humans are limited¹³.

Pathogenesis of cardiomyocyte involvement in diabetes mellitus involves vascular endothelial cell dysfunction and cardiomyocyte necrosis¹⁴. Whilst most structural studies of the myocardial vasculature have focused on coronary arteries and intramural arterioles, few studies have assessed myocardial capillary structure in either clinical or experimental diabetes showing significant ultrastructural changes in cardiomyocytes with or without concomitant vascular changes in the diabetic heart^{11,13,15}. These changes include thickening of endothelial cytoplasm, endothelial cell

hypertrophy, cytoplasmic bridging and irregular thickening of capillary basement membrane^{8, 11-13, 15-17}.

Vascular endothelial growth factor (VEGF) has been shown to contribute to the development of collateral vessels¹⁸ and is expressed in increased quantities in cardiomyocytes and arteriolar smooth muscle cells following MI in non-diabetic patients¹⁹. However, the expression of VEGF protein and mRNA, as well as its receptors, VEGFR1 and VEGFR2, are significantly decreased in the myocardium of both diabetic and insulin-resistant non-diabetic rats by 40-70%²⁰. In diabetic patients, the normal molecular processes which regulate angiogenesis may be impaired, although there are no direct studies to substantiate this to date.

Emerging evidence suggests that HIF-1 α and HIF-2 α have important and independent effects on pathological angiogenesis. Cells respond to hypoxic stress through multiple mechanisms, including the stabilization of hypoxia-inducible factors (HIFs), which directly regulate the expression of more than 150 target gene²¹.

Although functional changes in the diabetic heart are well established in the literature, the structural basis for these changes is not well defined. Most of the literature on ultra structural pathology of the micro vessels and cardiomyocytes is confined to animal models of diabetes, with studies in man being small and qualitative. It is therefore of considerable mechanistic and translational importance to define morphological and immunohistochemical changes in human diabetic myocardium. It was therefore decided to carry out a study with the objective to examine the interaction between expression of angiogenic factors and pathology of the microvasculature and cardiomyocytes in myocardial tissue of patients with diabetes mellitus.

Materials and methods

Biopsies from right atrial appendix were obtained of the forty seven patients underwent various cardiac procedures between June 2003 and January 2004 at the cardiothoracic surgical department, Manchester Royal Infirmary. Before surgery, all patients underwent cardiac catheterization and coronary angiography. This study was approved by the local Ethics Committee, and all patients gave their written informed consent before participation. Patients were classified into three clinical groups:

1. Diabetic patients undergoing CABG alone (DIHD) (n=18).
2. Diabetic patients undergoing cardiac valve replacement with normal coronary angiography (D-IHD) (n=7).
3. Non-diabetic patients undergoing cardiac valve replacement with normal coronary angiography (control group) (n=22)

Formalin fixed paraffin embedded tissue were sectioned and stained with H&E for routine examination and primary and secondary antibody for immunohistochemical examination after antigen retrieval. Blocking was done using 1% hydrogen peroxide solution. All the tissues were stained by a panel of primary antibodies including vWF(M)+antiCD31a (monoclonal), vWF(P)+antiCD31 (polyclonal), antiVEGFb (polyclonal), antiVEGFR2b (monoclonal) and antiHIF-1 α (monoclonal). Biotin conjugated secondary antibody was applied at the recommended dilution in appropriate non-immune serum in TBS. Streptavidin-HRP and Biotin were

diluted in TBS (pH 7.6) for 30 min. Microscopic assessment of the intensity of HIF-1 α , VEGF and VEGFR2 staining on blood vessels and cardiomyocytes was semi quantified using a scale from 1 to 10 according to the intensity of staining. Statistical analysis was performed using SPSS for Windows 14.0. All data are expressed the mean and standard error of mean (SEM). Clinical data were compared between groups by Kruskal-Wallis test or Mann-Whitney test, as appropriate. Linear regression analyses were performed to determine correlation between various pathological changes and angiogenic factor expression. Statistical significance was accepted for P< 0.05 (two-tailed).

Results

HIF-1 α expression

Intensity and percentage of HIF-1 α expression

There were significant differences in the intensity and percentage of HIF-1 α expression in cardiomyocytes between the three groups (P= 0.001, P= 0.01). Thus, intensity of HIF-1 α was

Table1: Intensity and percentage of HIF-1 α expression in controls and diabetic patients with and without ischemia

Parameters	Controls	D-IHD	DIHD	P-value
Intensity of HIF-1 α staining	1.5 \pm 0.3	2.8 \pm 0.4	3.5 \pm 0.4	0.001
Percentage of HIF-1 α staining	30.7 \pm 5.1	34.1 \pm 5.7	48.2 \pm 3.6	0.01

significantly increased in diabetic patients with and without ischemia compared to controls (P= 0.000, P= 0.03). In addition, the percentage of HIF-1 α was increased in DIHD patients compared to D-IHD patients (P= 0.05) and controls (P= 0.007) (Table 1)

Intero-relation between HIF-1 α expression and distal myocardial capillary pathology

There was a significant positive correlations between HIF-1 α intensity with number of endothelial nuclei (P= 0.02). However, HIF-1 α intensity correlated negatively with lumen area (P= 0.04). In addition, HIF-1 α percentage showed a positive correlation with endothelial area (P=0.03), basement membrane

Table 2: Correlation between HIF-1 α expression and distal myocardial capillary pathology

	Lumen area	Endothelial area	Pericyte area	BM area	Vessel area	Endothelial cell profile	No of endothelial cell nuclei	No of pericyte nuclei
HIF-1 α intensity	R = 0.31 P = 0.04	R = 0.12 P = 0.32	R = 0.05 P = 0.86	R = 0.15 P = 0.24	R = 0.08 P = 0.7	R = 0.21 P = 0.2	R = 0.35 P = 0.02	R = 0.02 P = -0.62
HIF-1 α percentage	R = -0.09 P = 0.45	R = 0.34 P = 0.03	R = 0.08 P = 0.7	R = 0.4 P = 0.01	R = 0.2 P = 0.18	R = 0.19 P = 0.35	R = 0.3 P = 0.02	R = 0.01 P = 0.87

P= P-value, R= Correlation Coefficient.

area (P= 0.01) and number of endothelial nuclei (P= 0.02) (Table3).

VEGF expression

VEGF expression in different blood vessel regions and cardiomyocytes

There was a significant difference in the expression of VEGF on vascular endothelium between the three groups (P= 0.002). Thus, it was significantly reduced in D-IHD patients compared to controls (P= 0.000). In addition, there was a non-significant trend towards a reduction in VEGF expression on the endothelium of DIHD patients compared to controls (P=0.06). No significant difference was found in VEGF expression on endothelium between DIHD and D-IHD patients (P= 0.11). There was no significant difference in VEGF expression on cardiomyocytes, basement membrane and pericytes between groups (P= 0.18, P= 0.09, P= 0.95) (Table4).

Inter-correlation between VEGF expression and distal myocardial capillary pathology in diabetic patients

There was a significant positive correlation between VEGF expression on endothelium and endothelial

cell profile no./capillary (P= 0.01). In addition, VEGF expression on cardiomyocytes showed positive

Table 3: VEGF expression on cardiomyocytes, endothelium, basement membrane and pericytes in controls and diabetic patients with and without ischemia

VEGF Expression	Controls	D-IHD	DIHD	P Value
Cardiomyocyte	2.85 ± 0.21	2.66 ± 0.15	2.18 ± 0.31	0.18
Endothelium	2.62 ± 0.24	1.32 ± 0.13	1.62 ± 0.23	0.002
Basement Membrane	0.84 ± 0.09	0.79 ± 0.35	0.92 ± 0.11	0.09
Pericyte	0.12 ± 0.03	0.12 ± 0.06	0.08 ± 0.02	0.95

Correlation with endothelial cell profile no./capillary (P= 0.04) (Table4).

Table 4: Correlations between VEGF expression and myocardial capillary pathology

VEGF Expression	Lumen area	Endo area	Pericyte area	BM area	Vessel area	Endo cell profile	No of endothelial cell nuclei	No of pericyte nuclei
Cardiomyocyte	R = 0.09 P = 0.8	R = 0.13 P = 0.32	R = 0.19 P = 0.4	R = 0.09 P = 0.7	R = 0.002 P = 0.9	R = 0.31 P = 0.04	R = 0.12 P = 0.5	R = 0.2 P = 0.12
Endothelial cell	R = -0.23 P = 0.009	R = 0.2 P = 0.18	R = 0.08 P = 0.63	R = 0.04 P = 0.65	R = 0.14 P = 0.4	R = 0.4 P = 0.01	R = 0.05 P = 0.74	R = 0.12 P = 0.3
Basement Membrane	R = 0.2 P = 0.1	R = 0.08 P = 0.6	R = -0.13 P = 0.12	R = 0.12 P = 0.8	R = -0.19 P = 0.2	R = 0.1 P = 0.3	R = 0.03 P = 0.8	R = 0.16 P = 0.3
Pericyte	R = 0.09 P = 0.86	R = 0.14 P = 0.3	R = 0.07 P = 0.9	R = 0.24 P = 0.1	R = 0.21 P = 0.17	R = 0.05 P = 0.9	R = 0.02 P = 0.8	R = 0.002 P = 0.96

P= P-value, R= Correlation Coefficient

Discussion

Epidemiological and clinical trial data confirm the higher incidence and prevalence of heart failure in diabetes. Although functional changes in the heart of diabetic patients are well established in the literature, the structural basis for these changes is not well defined. In the present study the expression of key angiogenic factors (HIF-1 α , VEGF, and VEGFR2) has been quantified in relation to both pathology and key clinical variables. We have shown a significant reduction in capillary density in diabetic patients with ischemia. This finding is in accordance with the finding of other researchers. It is well known that in diabetic myocardium there is an inadequate response to ischemia which results in poor collateralization^{18,22, 23}. The impaired angiogenic response in diabetes has been explained by various mechanisms. It could be as a result of the presence of vascular dysfunction, characterized by endothelial and vascular smooth muscle impairment²⁴⁻²⁶.

Reduced expression of VEGF and its receptors, along with reduction of other angiogenic factors and increase in expression of anti angiogenic proteins have been demonstrated in animals^{20,27}. In the present study, HIF-1 α expression was detected in cardiomyocytes and the expression of HIF-1 α was increased in diabetic patients compared to controls, but was maximally expressed in DIHD patients. Our observations are in contrast to other studies conducted in experimental animals²⁸. Expression of HIF-1 α is an important process allowing the myocardium to adapt to a reduction in blood flow and oxygenation. Experimental studies demonstrate that it might be a marker of persisting ischemia which indicates the presence of viable myocardium reacting to revascularization for a long period after an MI^{29, 30}. This explains our findings that expression of HIF-1 α on cardiomyocytes of diabetic patients is increased and may therefore indicate ongoing ischemia, which is the main activator for myocardial HIF-1 α production. The increase of HIF-1 α in D-DIHD patients suggests that although these patients had normal coronary arteries on angiography they still had micro vascular abnormalities characterized by low capillary density, reduced capillary luminal size, and thickening of basement membrane which could lead to ischemia and subsequent activation of HIF-1 α . Another explanation for increased HIF-1 α expression in D-IHD patients is it could be related to oxidative stress caused by ROS as hyperglycemia causes an increase of ROS and a state of

pseudohypoxia which might activate HIF-1 α expression^{28, 31}.

We show a significant negative correlation between the percentage of HIF-1 α expression and arteriole, capillary and total vessel density suggesting that HIF-1 α is upregulated by tissue hypoxia and ischemia. Thus, when vascular density increases via angiogenesis myocardial ischemia and hypoxia may reduce which subsequently down regulates HIF-1 α expression. In the current study, HIF-1 α expression correlates negatively with capillary luminal area whereas it correlated positively with basement membrane area and number of endothelial nuclei. These findings again support our hypothesis that coronary microangiopathy characterized by reduced luminal area, endothelial hypertrophy and thickening of basement membrane are associated with myocardial ischemia which stimulates HIF-1 α expression.

The best characterized regulators of angiogenesis are VEGF/VEGF receptor system³². VEGF expression is upregulated in response to hypoxia^{33, 34} and during physiological (wound healing) or pathological (tumor growth) needed for angiogenesis³⁵. In the current study we define the expression VEGF and its receptor VEGFR₂ in atrial appendages of diabetic patients with and without ischemia. We show a reduction in VEGF expression on the vascular endothelium of diabetic patients compared to controls. Whereas, VEGFR₂ expression showed a trend towards an increase in diabetic patients compared to controls. This finding is in contrast to the finding of reduction in VEGF and VEGFR1 expression by 40-70%²⁰, absence of VEGFR1 and VEGFR2³⁶, and lowered VEGF, VEGFR1 and VEGFR2 in diabetic animals³⁷. It is well known that in diabetic myocardium there is inadequate collateral formation as a result of an impaired angiogenic response. Thus the reduction in vascular density that we observe in the current study could be related to an impaired angiogenic response mediated by reduced VEGF in diabetic patient compared to controls. Recent experimental studies of diabetic myocardium have shown that low levels of VEGF precede all other features of diabetic cardiomyopathy and are followed by several changes which include increased apoptosis of endothelial cells, reduced numbers of circulating endothelial progenitor cells, reduced capillary density, and impaired myocardial perfusion. These changes lead to apoptosis and necrosis of

cardiomyocytes along with fibrosis and progressive diastolic and then systolic dysfunction³⁸. Thus, downregulation of VEGF expression may initiate or aggravate diabetic cardiomyopathy. We show an increased expression of VEGFR₂ on vascular endothelium in diabetic patients compared to controls. This upregulation in VEGFR₂ expression in diabetic patients could be a compensatory mechanism for the reduction in VEGF expression. However, the compensation was not enough to mediate the growth action of VEGF.

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