Loss of DPP4 activity is related to a prothrombogenic status of endothelial cells: implications for the coronary microvasculature of myocardial infarction patients

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Received: 16 August 2011 / Revised: 15 November 2011 / Accepted: 30 November 2011 / Published online: 14 December 2011
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Abstract Pro-coagulant and pro-inflammatory intramyocardial (micro)vasculature plays an important role in acute myocardial infarction (AMI). Currently, inhibition of serine protease dipeptidyl peptidase 4 (DPP4) receives a lot of interest as an anti-hyperglycemic therapy in type 2 diabetes patients. However, DPP4 also possesses anti-thrombotic properties and may behave as an immobilized anti-coagulant on endothelial cells. Here, we studied the expression and activity of endothelial DPP4 in human myocardial infarction in relation to a prothrombogenic endothelial phenotype. Using (immuno)histochemistry, DPP4 expression and activity were found on the endothelium of intramyocardial blood vessels in autopsied control hearts (n = 9). Within the infarction area of AMI patients (n = 73), this DPP4 expression and activity were significantly decreased, coinciding with an increase in Tissue Factor expression. In primary human umbilical vein endothelial cells (HUVECs), Western blot analysis and digital imaging fluorescence microscopy revealed that DPP4 expression was strongly decreased after metabolic inhibition, also coinciding with Tissue Factor upregulation. Interestingly, inhibition of DPP4 activity with diprotin A also enhanced the amount of Tissue Factor encountered and induced the adherence of platelets under flow conditions. Ischemia induces loss of coronary microvascular endothelial DPP4 expression and increased Tissue Factor expression in AMI as well as in vitro in HUVECs. Our data suggest that the loss of DPP4 activity affects the anti-thrombogenic nature of the endothelium.

Keywords DPP4 · Acute myocardial infarction · Tissue Factor · ROS · Coagulation · Platelet adhesion
Introduction

Research on coronary perfusion and restoration thereof after acute myocardial infarction (AMI) has largely been focused, for obvious reasons, on the larger branches of the coronary vasculature located on the epicardium. In the last two decades, however, the role of the smaller intramyocardial microvasculature has gained significant interest. Dysfunction of the intramyocardial microvasculature, specifically the endothelium, following AMI is suggested to relate to left ventricle remodeling, contractility and cardiac events, such as reinfarction and death [6, 9, 18, 40]. Also, hindered perfusion of the ischemic myocardium after restoration of coronary flow, known as no-reflow or coronary microvascular obstruction (MVO), might point to dysfunctional coronary microvasculature and associates with larger infarct size and poor clinical outcome [29, 39].

A multitude of factors may contribute to MVO including vasospasms, distal embolization, capillary plugging [24, 31]. A role for in situ thrombosis in MVO is debated [31], but the extrinsic (Tissue Factor-dependent) coagulation pathway was shown to play a role in MVO in rabbits [21, 22]. We have reported on structural and functional aberrations of the coronary microvasculature in AMI patients such as basement membrane thickening, adhesion molecule accumulation and products of non-enzymatic glycation (AGEs), as markers of a pro-inflammatory endothelium [3, 8]. It was also shown that a pro-inflammatory phenotype of the coronary microvasculature can induce cardiac fibrosis [28]. A post-AMI shift towards a more pro-inflammatory and pro-coagulant endothelial phenotype may therefore be involved in reduced coronary flow. During our studies we observed unexpectedly—as presented below—a marked decrease of microvascular endothelial dipeptidyl peptidase 4 (DPP4) expression in recently infarcted human hearts.

DPP4, also known as CD26, is expressed by microvascular endothelial cells in humans (i.e. in liver, spleen, lungs and brain) and in the hearts of rats [23, 34]. This serine exopeptidase terminally digests proteins and peptides containing proline in the penultimate position [10]. Its substrates include at least nine chemokines, neuropeptide Y and glucagon-type peptide-1 (GLP-1) [15, 23, 52]. Because it inactivates GLP-1, inhibition of DPP4 receives a lot of continuing interest as an anti-hyperglycemic therapy in type 2 diabetes [15]. In addition, recent publications suggest cardioprotective properties for GLP-1 analogues and pharmaceutical or genetic inhibition of DPP4 in different animal models of AMI [27, 47].

On the other hand, another known DPP4 substrate is fibrin and via cleaving N-terminal Gly-Pro from the fibrin z-chain, DPP4 can inhibit fibrin polymerization and clot formation [25, 42]. As such DPP4 may behave as an immobilized anti-coagulant on (microvascular) endothelium. Consequently, we asked ourselves whether the loss of coronary microvascular DPP4 after AMI relates to the endothelial phenotype shift towards a more pro-coagulant state and the induction of situ thrombosis as a possible inducer of MVO after AMI.

In the present study we evaluated a putative relation between DPP4 expression and pro-coagulant microvasculature in the infarcted human heart and the occurrence of thrombi in the coronary microvasculature. In addition, DPP4 activity was studied in relation to platelet adhesion in human umbilical vein endothelial cells (HUVECs).

Methods

Patients

A total of 82 autopsied patients were included in the study (Table 1). 73 Patients showed a recently developed left ventricular AMI [decreased lactate dehydrogenase (LDH) staining of the affected myocardium]. Nine patients who died without heart disease were included as controls. Sampling for (immuno)histochemical analysis was performed as in [3]. From the excised hearts 1 cm cross sections were made. In these cross sections the infarction area was determined using lactate dehydrogenase (LDH) staining. From each AMI patient, a sample of heart tissue was taken from the center of the LDH-determined infarction area and a sample was taken from the non-infarcted right ventricle as an internal control. In control patients a tissue sample was taken from the anterior wall of the left ventricle. The study protocol was approved by the ethics committee of the VU Medical Center, and executed according to the principles of the Declaration of Helsinki.

Microscopic criteria [36] were used to estimate infarct duration and to categorized into five phases of infarction: early phase, PMN (polymorphonuclear neutrophils) phase, chronic phase infarctions, early phase reinfarctions (chronic phase as well as early phase morphology) and PMN-phase reinfarctions (chronic phase as well as PMN-phase morphology) and corresponded with the clinical evaluation of AMI.

DPP4 enzymatic activity

A histochemical azo-coupling reaction was performed on 5 µm frozen heart tissue sections to identify DPP4 enzymatic activity. The slides were fixed in chloroform and acetone (1:1 v/v) and were incubated with the substrate glycyl-proline-4-beta-naphthylamide (Sigma St. Louis, MO, USA) dissolved in N,N-dimethylformamide (Merck, Haarlem, The Netherlands) and the azo-dye Fast Blue BB Salt (G.T. Gurr Ltd, London, UK) dissolved in PBS, fixed...
again in 4% formalin mounted in Immuno-mount (Shandon, Dreieich, Germany) and analyzed for the percentage of vessels positive for DPP4 activity.

Primary antibodies

For immunohistochemistry the following antibodies were used: mAb against DPP4 (1:100) (Serotec, Oxford, UK), mAb against CD31 (1:40) (Dako, Glostrup, Denmark), mAb against CD68 (1:400) (Dako), mAb against Tissue Factor (1:50) (Acris, Herford, Germany), and mAb against complement factor C3d (1:1,500) [36]. For digital imaging microscopy the following antibodies were used: mAb against DPP4 (1:100) (Serotec), mAb against Tissue Factor (1:50) (Acris) and pAb against nitrotyrosine (1:50) (Invitrogen Corporation, Carlsbad, CA, USA). For Western blot analysis the following antibodies were used: mAb against DPP4 (1:100) (Serotec), mAb against Tissue Factor (1:50) (Acris) and pAb against nitrotyrosine (1:50) (Invitrogen Corporation, Carlsbad, CA, USA). For all Abs PBS controls (Fig. 1a, right panel) and irrelevant Abs (IgG1), tested at similar concentrations, were negative.

Immunohistochemistry

Cryostat sections (5 μm) were fixed in chloroform and acetone (1:1 v/v) and blocked by 3% (v/v) H2O2/PBS. Serial slides were incubated in normal rabbit serum (Dakopatts) [1:50 in 1% (w/v) BSA/PBS], incubated with primary antibodies (in BSA/PBS), incubated with rabbit-anti-mouse-HRP (Dakopatts) (1:25 in BSA/PBS), incubated in 0.5 mg/ml 3,3′-diaminobenzidine tetrahydrochloride (DAB, Sigma, St. Louis, MO, USA) in 0.01% (v/v) H2O2/PBS and counterstained with hematoxylin.

Three investigators (P.A.J.K., N.H. and I.K.) independently scored all slides. The total number of blood vessels was determined with CD31. The percentage of DPP4- and Tissue Factor-positive blood vessels was then determined and equated for slide areas.

Cell culture

Freshly isolated HUVECs were cultured in M199 culture medium containing heat-inactivated human serum (10% v/v), fetal calf serum (10% v/v), 2,500 Units heparin and 100 μg/ml endothelial cell growth factor (cells cultured in this medium served as controls). For metabolic inhibition, cells were incubated in 0.9 mM CaCl2/C1H2O, 20 mM 2-deoxy-D-glucose (Sigma, St. Louis, MO, USA), 106 mM NaCl, 5 mM NaCN, 4.4 mM NaHCO3, 3.8 mM KCl and 1 mM MgCl2/H2O in H2O (pH 6.6). Apocynin (100 μM), diprotin A (10 mM), polymyxin B sulfate (2 μg/ml) and TNFα (10 μg/ml) (all Sigma) were incubated for 6 h.

Western blotting

After culturing in culture medium or metabolic inhibition buffer, HUVECs were lysed in buffer containing 0.01% NP-40, 2.5 mM Tris–HCl (pH 8.0), 150 μM NaCl, and 5 mM EDTA in H2O. Protein concentration was determined using the BCA protein assay kit (Pierce, Rockford, IL, USA). Then reducing (β-mercaptoethanol-containing) sample loading buffer [0.25 M Tris (pH 6.8), sodium dodecyl sulfate (SDS), glycerol, 2-mercaptoethanol, bromophenol blue] was added and the samples were mixed and heated at 95°C for 5 min. Proteins were separated via electrophoresis and blotted onto nitrocellulose membranes, incubated with primary antibodies, incubated with rabbit-anti-mouse-HRP (1:1,000) and then visualized by Enhanced ChemiLuminescence (ECL; Amersham, Buckinghamshire, UK).

Digital imaging microscopy

HUVECs were grown in 4-well chamber slides (Nalge Nunc International, Naperville, IL, USA). After treatment, the cells were fixed in 4% formaldehyde, permeabilized with acetone–methanol (70–30%), incubated with primary
antibodies, incubated with donkey-anti-mouse-Alexa 488 (1:40) (DPP4, Tissue Factor) and/or donkey-anti-rabbit-Alexa 647 (1:40) (nitrotyrosine) and incubated with wheat germ agglutinin (WGA), (1:40) (Invitrogen) to visualize cellular membranes (all in 1:2,000 v/v PBS/Tween20). Slides were covered in mounting medium containing DAPI (Vector Laboratories Inc, Burlingame, CA, USA) to visualize nuclei and analyzed with a 3I Marianas™ digital imaging microscopy workstation (Carl Zeiss, Sliedrecht, Netherlands). To determine cell viability, cells were stained with trypan blue and then quantified. In the digital imaging microscopy experiments on average 70 cells were analyzed per condition per donor.

Platelet isolation

Whole blood, obtained from healthy individuals, was centrifuged at 150×g. The supernatant was 1:1 diluted in Ca²⁺-free Krebs–Ringer buffer (4 mM KCl, 107 mM NaCl, 20 mM NaHCO₃, 2 mM Na₂SO₄, 19 mM tri-sodium
citrate, 0.5% (w/v) glucose in H2O (pH 5.0) and centrifuged at 500×g. The obtained non-stimulated platelets were labeled with calcein (Invitrogen) diluted 2:1 (w/v) in DMSO and used in a final concentration of 300,000/µl in Ca2+-free Krebs–Ringer flow buffer (pH 6.1).

Perfusion of HUVECs with platelets

HUVECs were cultured in fibronectin-coated µ-Slides VI0.4 (Ibidi GmbH, Martinsried, Germany) until uniform confluent monolayers were obtained. Prior to perfusion cells were incubated with either culture medium (controls) or with culture medium supplemented with Diprotin A (10 mM) or with TNFα (10 µg/ml) for 6 h. The slides were connected to two ANTEC syringe pumps [ANTEC (leyden), Zouterwoude, The Netherlands] via a flow kit (Ibidi) to generate a constant flow rate with a shear stress of 0.35 dyne/cm². The slides were mounted on a Zeiss Axiovert 200m microscope (Zeiss, Göttlingen, Germany). The HUVECs were perfused with calcein-labeled platelets for 3 min.

Statistical analysis

Paired or non-paired t tests, Mann–Whitney analysis, Chi-square test + ordinal Gamma test, Kruskal–Wallis test + Dunns multiple comparison tests or one-way ANOVA + Bonferroni’s multiple comparison tests were used when appropriate. In figures, values are given as means ± standard error. A p value (two-sided) of less than 0.05 was considered to represent a significant difference.

Results

DPP4 in the human myocardium

In human hearts, DPP4 was found on the endothelium in larger intramyocardial arteries (Fig. 1a) as well as in venules and capillaries (Fig. 1b). Cardiomyocytes were negative for DPP4 in all patients. DPP4 was found in the hearts of control patients as well as in the infarcted myocardium of AMI patients (Fig. 1c, grey bars). However, the percentage of DPP4-positive blood vessels was significantly lower in infarctions of all durations compared with controls, except for early-phase infarctions (PMN phase p < 0.001, chronic phase p < 0.01, early- and PMN-phase reinfarctions p < 0.05). In early-phase infarctions, the percentage of DPP4-positive blood vessels was markedly lower than in controls, but not significantly. DPP4 expression coincided with enzymatic activity in all infarct durations as well as in controls (Fig. 1c, black bars). Pearson correlation analysis revealed that in the controls and all phases of infarction patients’ age did not correlate with the DPP4 expression score: control (r² = 0.05167), early phase (r² = 0.01368), PMN phase (r² = 0.1468), chronic phase (r² = 0.001577), early-phase reinfarction (r² = 0.05768) and chronic-phase reinfarction (r² = 0.02594) (p > 0.05 in all groups). Notably, in chronic-phase infarctions DPP4 positivity was also observed in blood vessels, lymphocytes and granulation tissue (not shown).

In non-infarcted myocardium from AMI patients, no reduction in the percentage of DPP4-positive blood vessels was found compared to controls (Fig. 1d).

Tissue Factor in the human myocardium

As DPP4 may act as an anti-coagulant enzyme, the expression of pro-coagulant Tissue Factor was analyzed in control hearts and in PMN-phase infarctions and PMN-phase reinfarctions.

In control hearts, low levels of Tissue Factor were found in the sub-endothelial layer of a small number of intramyocardial blood vessels (Fig. 2a). Cardiomyocytes were negative for Tissue Factor. In the infarcted myocardium, the percentage of Tissue Factor-positive intramyocardial blood vessels significantly increased (PMN-phase infarctions p < 0.01, PMN-phase reinfarctions p < 0.001; Fig. 2a, b). Tissue Factor was found in the endothelial- and sub-endothelial layer of blood vessels (Fig. 2c, d) as well as on damaged cardiomyocytes that also stained positive for complement (factor C3d, not shown). As macrophages can also express Tissue Factor [13], we examined putative co-localization between macrophage marker CD68 and Tissue Factor. On Tissue Factor-positive endothelial cells no CD68-positive macrophages were found (not shown).

As Tissue Factor is a coagulation initiator we analyzed the presence of thrombi in the intramyocardial (micro)vasculature. We detected CD31-positive thrombi in the lumen of intramyocardial blood vessels in only 5 out of 32 analyzed patients (not shown). In these five patients, the percentage of intramyocardial blood vessels positive for Tissue Factor was higher than in AMI patients without detectable thrombi, although not significant (p = 0.06).

DPP4 and Tissue Factor expression in endothelial cells

DPP4 expression was decreased in those parts of the heart that were exposed to ischemia. To investigate whether ischemia by itself can induce the loss of endothelial DPP4, its expression was studied in metabolically inhibited HUVECs via digital imaging microscopy (Fig. 3a–c).

In control cells (Fig. 3aI), DPP4 was abundantly present on the cell surface as well as in the cytoplasm. DPP4 expression was reduced significantly after 3 (p < 0.001) and 6 h (p < 0.001) of metabolic inhibition (Fig. 3aII–III,
These results were verified by Western blot analysis (Fig. 3c). Notably, after 3 and 6 h of metabolic inhibition cell viability was still more than 95%, as determined by trypan blue staining (not shown).

In contrast, Tissue Factor was induced reaching statistical significance after 6 h of metabolic inhibition ($p < 0.001$; Fig. 3dI–III, e).

As exogenous reactive oxygen species (ROS), which are produced also during ischemia, can be involved in endothelial Tissue Factor expression [30], involvement of ROS in ischemia-induced Tissue Factor and/or reduced DPP4 expressions in HUVECs was analyzed via digital imaging fluorescence microscopy. Nuclear ROS significantly increased after 6 h of metabolic inhibition ($p < 0.001$; Fig. 4a). The antioxidant and inhibitor of NADPH oxidase-derived ROS apocynin significantly counteracted this increase ($p < 0.001$; Fig. 4a) as well as the effects of metabolic inhibition on the expressions of both DPP4 and Tissue Factor ($p < 0.001$; Fig. 4b, c). These results suggest that the effects induced by metabolic inhibition on DPP4 and Tissue Factor expression involve NADPH oxidase-derived ROS.

In addition, the putative effects of DPP4 activity inhibition by diprotin A on nuclear ROS and on Tissue Factor expression were analyzed in HUVECs via digital imaging microscopy. In contrast to metabolic inhibition Diprotin A induced a significant decrease in nuclear ROS ($p < 0.001$; Fig. 4d). Remarkably, diprotin A did induce a significant increase in endothelial Tissue Factor expression ($p < 0.0001$; Fig. 4e), which was verified via Western blot analysis (Fig. 4f). To confirm that the effects of diprotin A on nuclear ROS and Tissue Factor expression were not due to endotoxin contamination, the effects of diprotin A on nuclear ROS and Tissue Factor expression were analyzed in the presence of the endotoxin antagonist polymyxin B. Polymyxin B did not affect the diprotin A-mediated effects (3 independent experiments; not shown).

Inhibition of DPP4 activity induces adherence of platelets to endothelial cells

Aggregation of platelets to endothelial cells is a sign of thrombogenicity. Here we tested whether inhibition of DPP4 activity by diprotin A in HUVECs leads to increased adherence of non-stimulated platelets under flow conditions. Platelet adhesion-inducing TNFα was used as a positive control [2].
In untreated HUVECs, very few platelets adhered, mainly as solitary cells (upward arrows; Fig. 5a). Unbound platelets were recognized as unfocussed blurred stripes as they passed (dotted arrows). TNFα induced a marked increase in adhering platelets (Fig. 5b). These platelets adhered both individually (upward arrows) and as strings of platelets (downward arrows) of various length, sometimes up to 20 or more platelets per string. These strings were formed within seconds. Aggregation of platelets was not observed. Similarly, diprotin A induced a marked increase in adhering platelets compared to untreated cells (Fig. 5c). Again, platelets adhered both individually (upward arrows) and as strings of platelets (downward arrows), but did not aggregate. Interestingly, the adherence of platelets was in majority temporary in these conditions. Adhered platelets sometimes released after seconds, but most platelets adhered for longer and released after minutes. These results thus indicate that loss of DPP4 activity in endothelial cells can induce a prothrombotic status.
Discussion

Currently, DPP4 receives an attention as a target for anti-hyperglycemic therapy in type 2 diabetes patients and cardioprotective properties have been ascribed to GLP-1 analogues and to DPP4 inhibition in AMI. However, DPP4 can act as a proteolytic enzyme, a receptor and a costimulatory protein and has been shown to be involved in a wide range of different cellular functions [10]. We now show that enzymatically active DPP4 is expressed by the endothelium of the intramyocardial (micro)vasculature in humans. Interestingly, this DPP4 expression and activity decreased within the infarction area of AMI patients, coinciding with increased microvascular Tissue Factor expression. In HUVECs, metabolic inhibition induced ROS-dependent loss of DPP4 and concomitant Tissue Factor upregulation. And inhibition of DPP4 activity induced adhesion of platelets. Taken together, our data...
suggest a direct relation between DPP4 activity and thrombogenic status of endothelial cells.

The post-infarct decrease of intramyocardial microvascular DPP4 expression may be a direct result of the ischemia endured or due to enhanced availability of inflammatory cytokines such as TNFα. Involvement of ischemia is supported by our finding that persisting metabolic inhibition markedly reduced DPP4 expression in still viable human endothelial cells. This seems at variance with observations in a rat model of hindlimb skeletal muscle ischemia, in which control muscles did not contain DPP4 mRNA at all, and DPP4 mRNA was induced by ischemia [37]. However, neither protein levels nor cellular origin of DPP4 were determined. Furthermore, we observed an increase of DPP4 in blood vessels within granulation tissue in chronic-phase infarctions. Therefore, these differences

![Image](A control) (B TNFα) (C diprotin A)
may be related to differences in expression or timing between muscle cells and endothelial cells. Inflammatory cytokines such as TNF-α and IL-6 were shown to be involved in impaired coronary microvascular perfusion [32, 33, 38, 45, 51]. As TNF-α can both reduce DPP4 [48] and induce Tissue Factor [11, 20], we cannot exclude that such inflammatory cytokine also may contribute to the effects observed in infarcted hearts [33]. However, it does not contribute to the observations in metabolically inhibited cultured endothelial cells.

Contrary to DPP4, Tissue Factor was upregulated in the infarcted heart, both in the intramyocardial microvasculature as well as in damaged cardiomyocytes. This increase may also result from the ischemia endured, as demonstrated in endothelial cells in vitro. Tissue Factor expression was shown earlier in human and animal hearts and was implicated in mediating reperfusion injury, as Tissue Factor inhibition resulted in decreased inflammation and infarct size in rabbit and mouse AMI models [12, 14, 22]. Tissue Factor, a member of the cytokine receptor superfamily, initiates blood coagulation through the binding of FVII/FVIIa [41]. Additionally, the Tissue Factor/FVIIa complex or Tissue Factor alone can activate different (pro-inflammatory) cellular signaling pathways [17, 41]. Tissue Factor inhibition was shown by two independent groups to reduce myocardial infarct size in rabbits, although one group suggested that this was due not through inhibition of thrombotic events but through inhibition of inflammation [12], while the other concluded that it reduced intramyocardial coagulation and reduced the no-reflow phenomenon [22].

Platelet adhesion resulting in microvascular plugging was shown to cause reduced perfusion and no-reflow after AMI [7, 46]. In our in vitro experiments, inhibition of DPP4 activity induced adherence of non-stimulated platelets to HUVECs under flow conditions. Adherence of non-stimulated platelets to cytokine- or microparticle-stimulated endothelial cells was shown before [1, 2, 16, 26, 49, 53]. It was shown that platelet adhesion can depend on Tissue Factor [49, 53], interactions between platelet F11 receptor and endothelial JAM [2] and on secreted von Willebrand factor (vWf) [1, 16, 26]. It was shown recently that platelet-derived growth factor CC (PDGF-CC), released from activated platelets induced Tissue Factor expression in human microvascular endothelial cells [19]. Although in our platelet adhesion experiments we used non-stimulated platelets we cannot exclude such a role for PDGF-CC. In our experiments, platelets adhered individually and in strings. These strings of platelets indicate a role for vWF, as it was shown earlier that platelets bind endothelial cells through vWF in such fashion [1, 16, 26].

Therefore, the reduced presence/activity of DPP4 and increased presence of Tissue Factor in the post-MI intramyocardial microvasculature suggests a shift towards a prothrombogenic status of the endothelium and an increased risk of intramyocardial thrombosis. However, in only 5 out of 32 analyzed AMI patients intramyocardial thrombi were found. As thrombolytic treatment is standard procedure for AMI patients, prospective thrombi may dissolve prior to analysis.

The ischemia-induced effects on DPP4 and Tissue Factor expressions were inhibited by the antioxidant and inhibitor of NADPH oxidase-derived ROS apocynin, suggesting ROS involvement in both events. Involvement of ROS in Tissue Factor expression was reported before. In primary endothelial cells exogenous superoxide induced concomitant NADPH oxidase activation and Tissue Factor expression [30] and LPS induced NOX1-mediated Tissue Factor expression [43], whereas PAR1/2-induced endothelial Tissue Factor expression was mediated solely via mitochondrial ROS [5]. Mitochondrial ROS are a potential candidate for involvement in ischemia-induced Tissue Factor expression and possibly ischemia-induced loss of DPP4.

In contrast to ischemia, DPP4 activity inhibition with diprotin A reduced nuclear ROS production. It was recently shown that GLP-1 protected endothelial cells against oxidative stress in vitro [44]. However, in the said study the effects of added GLP-1 on cultured endothelial cells were investigated and since there is no evidence available so far that endothelial cells can produce GLP-1 themselves, it is unlikely that GLP-1 was a factor in our in vitro experiments. Thus, diprotin A most likely reduced ROS independent of GLP-1. Remarkably, diprotin A simultaneously induced a significant increase in endothelial Tissue Factor expression. The fact that diprotin A, even though it lowered nuclear ROS, did induce Tissue Factor, strongly suggests that different cellular pathways are involved in Tissue Factor upregulation induced by ischemia on the one hand (which increased nuclear ROS) and by diprotin A on the other (which lowered nuclear ROS). Whether, and if so to what extent, these different pathways are relevant in the infarcted heart in vivo remains to be established.

Dysfunction of the intramyocardial microvasculature following AMI relates to left ventricular remodeling and cardiac events [6, 9, 18], may relate to no-reflow of the ischemic myocardium and is associated with larger infarct size and poor clinical outcome [29]. The AMI-induced loss of DPP4 and concomitant Tissue Factor upregulation we show here may contribute to this dysfunction. DPP4 activity was also implicated in revascularization of ischemic tissues via conversion of neuropeptide Y [37, 52]. Our observed DPP4 increase in chronic-phase infarctions in blood vessels within granulation tissue might support such a role. Alternatively, recent studies show evidence for a cardioprotective role for pharmacologic inhibition and of
genetic depletion of DPP4 in rat- and mouse-myocardial infarction models [27, 35, 47, 50]. These cardioprotective/infarct size limiting effects of DPP4 inhibition were largely attributed to the prevention of GLP-1 degradation. The cardioprotective effects of increased GLP-1 levels may involve improved glucose uptake, increased myocardial cyclic AMP levels and activation of cardioprotective kinases, such as PKA and AKT [27, 35, 47, 50]. However, Ban et al. [4] showed novel cardiac and vascular protective effects of truncated GLP-1(9–36), suggesting that DPP4 inhibition in patients at risk for AMI appears warranted.

In conclusion, we show that DPP4 activity is related to the thrombogenic status of endothelial cells in that lack of DPP4 activity induces expression of Tissue Factor and platelet adhesion, and that this relation may have implications regarding post-AMI dysfunction of the coronary microvasculature.

Acknowledgments Dr. Kholova was a PhD student of the Marie Curie Training Site of the European Community (No. HPMT-2000-114) at the ICaR-VU.

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