

IMPACT OF ALTERED EXPRESSION OF MATRIX METALLOPROTEINASES (MMPs) ON THE DEVELOPMENT AND SEVERITY OF DIABETIC RETINOPATHY.

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INTRODUCTION: The primary cause of visual loss in the country like India are cataract, glaucoma, malnutrition, age related macular degeneration and diabetic retinopathy (DR)¹. DR is the most common microvascular complication of both type 1 and type 2 diabetes mellitus (DM) and is the most frequent single cause of newly reported cases of blindness among the adults in the age group of 20 – 75 years ². Visual loss due to diabetes either develops from increased permeability of retinal vessels (diabetic macular edema) or by proliferation of new retinal vessels ².

Several angiogenic factors such as VEGF, PDGF, and TNF- α exhibit their potential signature on the development and progression of DR which ultimately results in the neovascularization process³. So far only VEGF has been largely implicated in deposition of extracellular matrix (an essential step in the new vessel formation) and play the crucial role by stimulating angiogenesis and inhibiting the endothelial function in retinal ischemic conditions.³ Among the various processes that regulate angiogenesis, the generation of proteolytic activity is thought to be pivotal in the regulation of cell migration and capillary tube formation. Pericellular proteolysis and capillary-like tubule formation by endothelial cells are achieved by cell-bound urokinase-type plasminogen activator (u-PA) and plasmin as well as matrix metalloproteinases (MMPs) ⁴. Membrane-type MMPs (MT - MMPs) have been suggested to play a key role in angiogenesis, in addition to the gelatinases MMP-2 and -9 ⁵. As a family of zinc-binding, calcium-dependent endopeptidases, matrix metalloproteinases (MMPs) expressed in various tissues and cells are responsible for the dissolution of extracellular matrix ⁶. MMPs play a major role in normal development, reproduction, and tissue remodeling⁷. In addition, these molecules are involved in a number of pathogenic processes including cancer, arthritis and other angiogenic disorder.⁸ Recently there have been a number of important discoveries on the role of MMPs in the eye. MMP activity is modulated through interaction with the MMP inhibitors, i.e., tissue inhibitors of metalloproteinases (TIMPs).⁷ MMP activities are regulated by transcriptional control, by proenzyme activation. MMPs are

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involved in the degradation of a variety of extracellular matrix (ECM) molecules, and ECM homeostasis.

Considering the finding that DR is initially caused by breakdown of inner blood-retinal barrier and later by fibrovascular proliferation; it is possible that MMP expression might change in the development and progression of DR and the extracellular matrix degradation would be also accompanied accordingly. To understand the pathogenic mechanism of the multifactorial disease like DR, we investigate the role of gelatinized MMPs from clinical isolates.

METHODOLOGY: SELECTION OF PATIENTS:

INCLUSION CRITERIA: The study included 373 subjects suffering from DR (n=253, 60, 60 subjects suffering from proliferative DR, severe non proliferative DR and mild non proliferative DR, respectively) as a long term complication of type2 DM and 240 age, sex, nutrition and glycemic level matched type 2 diabetic controls (duration of diabetes 17 ± 5 years) without retinopathy and 100 non diabetic healthy control. DR cases were recruited from the retina clinic of the Regional Institute of Ophthalmology, Kolkata, India and the controls were recruited at the diabetic clinic of the Institute of Post-graduate Medical Education and Research, Kolkata; India. All the study subjects belonged to the same community (same ethnic group) -Bengali Hindu, living in geographical proximity and are assumed to be pan-mixing. Institutional ethical clearance and written informed consent from each subject was obtained according to the Declaration of Helsinki.

Diagnosis of DM was made according to WHO criteria (WHO, 2006). Diagnosis of PDR was done by well qualified retina specialist through dilated fundus examination with slit lamp biomicroscopy by +90D and 3 mirror lens, seven field digital fundus photography with fluorescence angiography. Grading or scale of severity of retinopathy was based on modified early treatment diabetic retinopathy study (ETDRS). Among those the PDR (n=48) subject who were in the proliferative stage; had undergone pars plana vitrectomy and undiluted vitreous were collected from these patients for selective protein study.

30 subjects who underwent vitrectomy for idiopathic macular hole were included as control for vitreous study. This disorder is caused by vitreo-macular traction occurring prior to posterior vitreous detachment and there are no signs of ischemia, proliferation or inflammation.

EXCLUSION CRITERIA: People with coronary artery diseases (CAD), hypertension, peripheral vascular diseases, CAD, history of any thrombotic event, acute infection, patients having any other ocular disorder like glaucoma, branch retinal venous occlusion and Eales' disease were excluded from the study. In order to exclude overt diabetic nephropathy patients, subjects with micro albumin-creatinine ratio > 30 mg/gm and urinary micro albumin level > 300 mg /day were excluded from the study (Alvin, 2005).

SAMPLE COLLECTION AND PREPARATION FOR ANALYSIS: 10 ml of whole blood from DR patients and controls were collected by venipuncture from peripheral veins. 7ml of blood was taken in a EDTA containing tube for PBMC layer separation. Rest of 3ml was collected in clot vial and allowed to clot. After clot retraction, serum was separated by centrifugation at 3,000 g at 4°C for 20 minutes stored at -80°C until further use.

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Vitreous samples were collected from the study subjects and control patients who underwent three-port pars plana vitrectomy. After the construction of the ports, vitreous cutter was introduced in mid vitreous, before turning the infusion fluid 200µl of undiluted vitreous gel was excised and aspirated into the hand held sterile syringe attached to the suction port of the vitrectomy probe, using manual suction with high cutting rate. The vitreous biopsy samples thus obtained were immediately put in ice, centrifuged at 10,000g for 15 minutes at 4°C. After centrifugation supernatants were divided in two equal aliquot and stored at -20°C for immediate use and at -80°C for future use. Simultaneously venous blood was collected from those patients at the time of vitrectomy and processed as stated earlier.

The Demographic and clinical characterization of the study subjects were presented in table 1(DR and DC) and table2 (DC and HC) There was no significant difference in age, sex, blood pressure, and nutritional status between the study groups. The glycemic control were matched among the diabetic subjects.

TOTAL PROTEIN ESTIMATION FROM CLINICAL ISOLATES: Total protein from clinical isolates (serum and vitreous fluid) as well as cell culture supernatant and cell lysates were estimated by Bicinchoninic acid (BCA) assay (Smith et al., 1987)⁹ as per requirement. Protein (mg/ml) levels were estimated by using BCA reagent (Sigma) containing 4% copper sulphate (CuSO₄). Total concentration of gelatinized MMPs i.e. MMP2 and MMP9 in vitreous and serum of study subjects were measured using a commercial ELISA kit (R&D Systems).

CONCENTRATION OF GELATINIZED MMPS (MMP2 AND MMP9) IN VITREOUS FLUID OF PDR SUBJECTS AND THE SERUM LEVEL OF THE SAME CYTOKINE IN DIFFERENT STAGES OF DR.

Gelatinized vitreous MMPs concentration (MMP2 and MMP9) was found significantly higher than that of controls [260.18±97.69ng/ml (PDR) vs. 86.79±39.73 ng/ml (Control); p<0.0001 for MMP2 and 52.34±26.14 ng/ml (PDR) vs.26.21±8.87 ng/ml (Control); p= 0.0002 for MMP9.Table3

The serum level of MMP2 and MMP9 were found to be significantly elevated in mild NPDR(MMP2: 167.36 ± 42.4 ng/ml,MMP9: 732.62 ± 256.51) subject compared to control (DC) [MMP2: 99.44 ± 27.58 pg/ml MMP9: 611.11 ± 60.26]; p<0.0001 for MMP2 and p<0.0001 for MMP9 .Further the concentration of MMP2 and MMP9 significantly upregulated in severe NPDR subject(MMP2:188.13 ± 44.29 ng/ml.MMP9: 902.25 ± 114.28) compared to mild NPDR ;p<0.0118 for MMP2 and p<0.0001 for MMP9.The MMP2 level insignificantly elevated in PDR subject (198.85 ± 48.47 ng/ml) compared to severe NPDR p= 0.3447and MMP9 level reach its most activation sate on PDR subject (1265.4 ± 156.04ng/ml) vs. severe NPDR; p<0.0001]. Further it was also noted that the serum MMP2 and MMP9 concentration significantly elevated in DC compared to Healthy control (Table 4).

DISCUSSION: PDR is a common complication of type2 (and type1) DM, characterised by periretinal neo-vascularization and development of epiretinal fibrovascular tissue^{10,11}.Early features of DR include selective loss of intramural pericytes from retinal capillaries which leads to damage of the inner blood-retinal barrier (Aiello et al.,1995. This process is followed by neo-vascularization, involving the production of angiogenic factors as well as synthesis of extracellular matrix (ECM) necessary for anchorage of migrating endothelium and other cells such as retinal pigment

epithelium (RPE), glial cells, and fibroblasts¹¹. Degradation of ECM proteins is exerted by MMPs, a family of zinc binding, calcium dependent enzymes¹². Our present study demonstrated that MMP2 and MMP9 markedly upregulated in the vitreous fluid in PDR subjects compared to control. Further this MMPs particularly MMP9 gradually elevated at regular manner in serum level during the severity of DR which may act as the marker for adverse prognosis of this microvascular complication of type2 DM. Serum MMP2 activated at MNPDR and SNPDR subjects and may established its pathogenic potentiality for the initiation of this retinal complication among the prolonged hyperglycemic subjects. Activation of MMPs both in vitreous and serum may play a crucial role in connective tissue remodelling and in the degradation of basement membrane and surrounding extracellular matrix during processes of angiogenesis. Extracellular proteinases (i.e., MMPs and urokinase plasminogen activator) may play an important and rate-limiting role in the process of neovascularization¹³ Expression of these proteinases is required for vascular endothelial cells to "break-through" their surrounding basement membrane and migrate through interstitial tissue toward the angiogenic stimulus¹⁴. Many angiogenic factors including VEGF, bFGF, TNF- α selectively stimulate the expression of various MMPs (MMP 1 to 3 and 9) in vascular endothelial cells¹⁵. Agents that are MMPs enzyme inhibitors (batimastat, marimastat) and agents that alter vascular endothelial cells MMPs expression currently are being evaluated as angiostaticagents¹⁶. It also appears that the MMP system may be involved in regulating angiogenesis in a much more complex manner (Joseph et al., 2008). In addition, MMPs may be responsible for the generation of natural angiostatic substances such as angiostatin, which inhibits vascular endothelial cell proliferation and is the proteolytic product of plasminogen¹⁶. A recent study adds additional complexity to the role of MMPs in angiogenesis (Joseph et al., 2008). Previous studies hypothesized that that PEX, a proteolytic fragment of MMP-2 containing a hemopexin-like domain, inhibits the functional association of MMP-2 with an integrin receptor (av/33) on the surface of vascular endothelial cells¹⁷. PEX binding effectively blocks the expression of collagenolytic activity on vascular endothelial cells and is thereby angiostatic¹⁸. PEX progressively accumulates in the retina during retinal vasculogenesis, suggesting that it may play a normal role in the regulation of blood vessel formation (Brooks et al., 1998; Rupp et al., 2008). Alteration pattern also seen in the vitreous fluid where MMP2, and 9 are significantly elevated in the vitreous fluid of PDR subjects. Previously, retinal pigment epithelial (RPE) cells in culture have been shown to secrete both MMPs and TIMPs¹⁹.

It is therefore suggested that systematic continuous activation of gelatinized MMPs in hyperglycemic subjects may exert its crucial role in connective tissue remodeling, degradation of basement membrane and surrounding extra cellular matrix during the process of progression of diabetic retinopathy leading to angiogenesis and fibro cellular proliferation.

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Table1. Clinical characteristics, nutritional and glycemc status among DR and DC

Parameter	DR	DC	p-Value
Age	53 ±16	54±12	0.4
Sex			
Male	203 (54.42%)	128(53.33%)	0.8
Female	170 (45.58%)	112 (46.67%)	
HBA ₁ C (%)	7.8 ± 1.2	7.6 ± 1.3	0.51
Urinary micro albumin (mg/day)	18.92 ± 5.8	18.1 ± 6.08	0.0942
Urinary micro albumin creatinine ratio (mg/gm)	18.23 ± 5.19	17.54 ± 5.64	0.121
Serum total protein (gm/dl))	7.18 ± 1.02	7.1 ± 0.88	0.3182
Duration of diabetes (years)	18 ± 6	17 ± 7	0.0599
Blood Systolic Pressure	132 ± 9	131 ± 8	0.1616
(mmHg) Diastolic	83 ± 6	82 ± 7	0.0597

Values are represented in mean ± SD. p- value Significant at (0.05) level.

Table2. Clinical characteristics, nutritional and glycemc status among DC and HC

Parameter	HC	DC	p-Value
Age	52 ±13	54±12	0.21
Sex			
Male	55 (55%)	128(53.33%)	0.82
Female	45 (45%)	112 (46.67%)	
HBA ₁ C (%)	-----	18.1 ± 6.08	-----
Urinary micro albumin (mg/day)			
Urinary micro albumin creatinine ratio (mg/gm)	-----	17.54 ± 5.64	-----
Serum total protein (gm/dl))	7.28 ± 1.3	7.1 ± 0.88	0.1395
Duration of diabetes (years)	-----	17 ± 7	-----

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Blood Systolic Pressure	130 ± 9	131 ± 8	0.3125
(mmHg) Diastolic	81 ± 8	82 ± 7	0.251

Values are represented in mean ± SD. p-value Significant at (0.05) level.

Table 3: Concentration of gelatinized MMPs (MMP2 and MMP9) in vitreous fluid of PDR subjects compare to control

	MMP2	MMP9
PDR	260.18±97.69ng/ml	52.34±26.14 ng/ml
Control	86.79±39.73 ng/ml	26.21±8.87 ng/ml
P-VALUE	p<0.0001	p= 0.0002

TABLE 4: Concentration of gelatinized MMPs (MMP2 and MMP9) in serum among different degrees of DR subjects

	MMP2	P-VALUE	MMP9	P=VALUE
HC	82.42±59.73ng/ml	<0.0001	473± 51.73ng/ml	<0.0001
DC	99.44± 27.58 ng/ml		732.62 ± 256.51ng/ml	
MNPDR	167.36 ± 42.ng/ml	<0.0001	611.11 ± 60.26ng/ml	<0.0001
SNPDR	188.13± 44.29.ng/ml	0.01	902.25 ± 114.28ng/ml	<0.0001
PDR	198.85 ± 48.47 ng/ml.ng/ml	0.03	1265.4 ± 156.04ng/ml	<0.0001