

Role of Low Density Lipoprotein Cholesterol in Progression of Diabetic Retinopathy

RAHUL¹, NARSINGH VERMA², MAYANK AGARWAL³, PRAVESH KUMAR⁴, SANDEEP SAXENA⁵, ANUJ MAHESHWARI⁶

ABSTRACT

Introduction: Various cross-sectional studies indicate that dyslipidemia and increased activity of enzyme Aldose Reductase (ALDR-2) are associated with the establishment of Diabetic Retinopathy (DR). However, it remains unclear that among dyslipidemia and ALDR-2 activity, which is more important for the progression of DR from nonproliferative to the proliferative stage.

Aim: The present study was designed to explore the role of dyslipidemia and ALDR-2 activity in the progression of DR.

Materials and Methods: Two hundred subjects were involved in this cross-sectional study, 150 subjects had Type 2 diabetes while 50 had No Diabetes and No Retinopathy (NDNR) and acted as controls. On the basis of fundus examination, diabetic subjects were further divided equally into those having: No Retinopathy (DNR), Non-Proliferative DR (NPDR), and Proliferative DR (PDR). Fasting lipid profile, ALDR-2 level,

fasting and postprandial blood sugar were measured using a standard protocol. Data were analysed using both descriptive and inferential statistics using unpaired t-test.

Results: ALDR-2 level was significantly ($p\text{-value}=0.0001$) higher in NPDR than DNR. Low-Density Lipoprotein Cholesterol (LDL-C) was significantly ($p\text{-value}=0.0235$) higher in PDR than NPDR. Fasting Blood Sugar (FBS), Postprandial Blood Sugar (PPS), and LDL-C were significantly higher in DNR than NDNR while high-density lipoprotein cholesterol was significantly lower in DNR than NDNR group.

Conclusion: Persistent hyperglycaemia causes an increased ALDR-2 activity that has a significant role in the establishment of DR. However, dyslipidemia is more important as a risk factor for progression of NPDR to PDR. Hypolipidemic drug as an adjunctive therapy could prevent the progression of DR from NPDR to PDR.

Keywords: Aldose reductase, Diabetes mellitus, Diabetic retinopathy, Dyslipidemia

INTRODUCTION

The increased activity of enzyme ALDR-2 has been recognised as a major risk factor for the development of microvascular complications in diabetic patients [1-3] that comprises neuropathy, nephropathy and retinopathy. Among these microvascular complications, DR is different in having two-staged disease progression. DR initially develops as NPDR due to the increased activity of ALDR-2. If NPDR remains uncontrolled, it later progresses to PDR that has a multifactorial causation resulting in active angiogenesis in the retina [4].

Proliferative diabetic retinopathy is a serious diabetic microvascular complication that affects the quality of life by contributing to the visual loss in diabetic patients. The pathological importance of PDR lies in the fact that it presents as a sudden diminution or complete loss of vision; whereas, the development and manifestation of the other two microvascular complications involves a relatively long time course [5]. Neovascularization in PDR results in the formation of delicate capillaries over the retina, which ruptures easily by a sudden change in the blood pressure, leading to the haziness of vitreous humour due to haemorrhage and may cause immediate blindness.

Although, the age of the patient [6], the duration of diabetes, higher glycosylated haemoglobin levels, the presence of proteinuria and hypertension have been established as risk factors for DR [7], the current knowledge about other causes like deranged lipid profile remains unsatisfactory. Polyol pathway initiation is the foundation unit for production of sorbitol, fructose and other harmful by products in PDR that leads to various alterations in the intracellular and the extracellular environment of cells [8]. These changes include: modification of intracellular tonicity, due to the generation of

'Advanced Glycation End Products' (AGEs); and exposure of cells to oxidative stress presumably through the production of oxidant species coupled with declining antioxidant defence mechanism [9]. Apart from polyol pathway, other pathways like Protein Kinase C (PKC) activation, glucose autoxidation, prostanoid synthesis, protein glycation and hexosamine pathway might play a minor role in the establishment of DR [10,11]. These pathways ultimately result in the degeneration of pericyte cells by apoptosis that leads to the damage of blood-retinal barrier causing haemorrhage along with the leakage of lipid and protein macromolecules present in plasma [12]. In the highly oxidative environmental condition of the retina, these lipids and modified protein will cause accumulation of 'advanced lipoxidation end products' [13]. Thus, DR progresses to PDR, manifesting the signs of the disease.

Several studies have been done to correlate the increased activity of ALDR-2 with various traditional and non-traditional lipid markers for the establishment of DR [14]. However, to the best our knowledge, till date none of the studies have clarified the role of dyslipidemia in the progression of DR. We hypothesised that both, dyslipidemia and increased ALDR-2 activity are responsible for the progression of NPDR to PDR. The aim of the current study was to determine whether dyslipidemia or increased ALDR-2 activity plays a major role in the progression of DR from NPDR to PDR.

MATERIALS AND METHODS

The present cross-sectional was conducted during 2016-2017 that involved 200 subjects, of which, 50 subjects were serving as healthy controls (NDNR) while rest 150 had Type 2 DM. Subjects

having Type 2 DM were further divided into three equal groups after ophthalmoscopy examination into those having diabetes with no retinopathy (DNR, n=50), NPDR (n=50) and PDR (n=50). Group and subgroup wise distribution is shown in [Table/Fig-1].

Group	Sub group	Description	No. of patients
Study group (Type 2 DM)	1	Diabetes mellitus with Non-Proliferative Diabetic Retinopathy (NPDR)	50
	2	Diabetes mellitus with Proliferative Diabetic Retinopathy (PDR)	50
	3	Diabetics mellitus with No Retinopathy (DNR)	50
Control group	4	No Diabetes and No Retinopathy (NDNR)	50

[Table/Fig-1]: Group and subgroup wise distribution of subjects.

According to WHO (2015 data) prevalence of diabetes in India was 8.7% [15]. From the above data p-value for diabetics is 0.087, at 5% margin of error and 95% confidence interval, the total sample size calculated was 122. However, we included 150 diabetic patients and 50 normal subjects in our study.

The present study was performed at the Retina Clinic of Ophthalmology Department in collaboration with the Physiology Department, King George's Medical University, Lucknow; after obtaining ethical clearance from the Institutional Ethical Committee. Subjects were recruited after a thorough history and blood sugar test. Before inclusion of diabetic patients in the study, they had their fundus examined by both direct and indirect ophthalmoscopy for the demarcation and categorization of DR according to (ETDRS) (Early Treatment Diabetic Retinopathy Study) classification [16].

Exclusion criteria were the diabetics on insulin or statin treatment and those who were anaemic or had any liver disease. A complete record of each participant, about their age, gender, clinical symptoms, duration of diabetes, medication and the socioeconomic background was noted using a well-designed questionnaire. All individuals gave the written informed consent before involvement in the study as subjects.

A fasting blood sample was drawn in the morning at 8-9 AM from the subjects into the Ethylene Diamine Tetra Acetic acid (EDTA), sodium fluoride (NaF), and plain vial, for ALDR-2, glucose and lipid profile estimation respectively. All the tests were executed on the same day of sampling in the clinical laboratory of Biochemistry Department, KGMU, Lucknow.

A 10% erythrocyte suspension was made by adding 50 mM sodium phosphate buffer, pH 7.4, containing 150 mM NaCl. The cells in the suspension were lysed by repeated freezing and thawing (three cycles) and then centrifuged to remove any insoluble material. ALDR-2 activity was measured spectrophotometrically using an appropriately diluted haemolysate according to a previously described method [17] using a SpectraMax spectrophotometer (Molecular Devices, Sunnyvale, CA).

Parameters	Control	Study Group			Unpaired t-test (p-value)		
	NDNR (N=50)	DNR (N=50)	NPDR (N=50)	PDR (N=50)	NDNR vs DNR	DNR vs NPDR	NPDR vs PDR
Age (years)	55.84±5.62	56.02±7.93	56.92±7.22	57.50±6.95	0.8961	0.5543	0.6833
FBS (mg/dL)	97.00±26.46	132.68±51.13	137.28±56.17	141.73±38.54	0.001*	0.6694	0.47
PPS (mg/dL)	134.64±71.53	210.08±88.96	192.32±79.18	224.65±58.03	0.001*	0.2943	0.48
TG (mg/dL)	153.04±61.37	161.08±47.78	168.35±92.22	190.36±80.57	0.4665	0.6217	0.2080
TC (mg/dL)	148.80±23.77	155.38±38.39	168.36±41.70	168.40±41.08	0.3053	0.1011	0.1182
HDL-C (mg/dL)	51.24±11.87	44.60±16.28	43.42±10.62	42.40±10.64	0.0218*	0.4685	0.6390
LDL-C (mg/dL)	63.92±19.52	78.77±29.61	84.60±40.66	101.60±32.82	0.0038*	0.4143	0.0235*
VLDL (mg/dL)	31.60±12.58	31.93±8.72	35.50±17.5	38.40±16.38	0.8791	0.2015	0.3914
ALDR-2 (unit/gm of Hb)	3.56±1.68	4.24±2.11	7.47±2.02	8.22±2.19	0.0777	0.0001*	0.20

[Table/Fig-2]: Showing comparison for variable on the basis of mean, number of subjects (n) and standard deviation between NDNR, DNR, NPDR and PDR subgroups.

*significant, NDNR; No diabetes and no retinopathy, DNR; Diabetic but no retinopathy, NPDR; Non-proliferative diabetic retinopathy, PDR; Proliferative diabetic retinopathy, FBS; Fasting blood sugar, PPS; Post prandial sugar, TG; Triglyceride, TC; Total cholesterol, HDL-C; High density lipoprotein cholesterol, LDL-C; Low density lipoprotein cholesterol, VLDL; Very low density lipoprotein, ALDR; Aldose reductase.

A total of 5 parameters: Triglyceride (TG), Total Cholesterol (TC), High Density Lipoprotein (HDL), LDL and Very Low Density Lipoprotein (VLDL) were assessed by auto-analyser.

Glucose was estimated in plasma by the Glucose Oxidase-Peroxidase (GOD-POD) method using a kit. 10 microlitres of serum or standard (100 mg/dL glucose) was added to a reagent (100 mM phosphate buffer, pH 7.5 with 5 mM phenol, 10 U/mL glucose oxidase, 1 U/mL peroxidase and 0.4 mM 4-amino antipyrine) and incubated for 5 minutes at 37°C. Absorbance was measured at 505 nm by an auto-analyser.

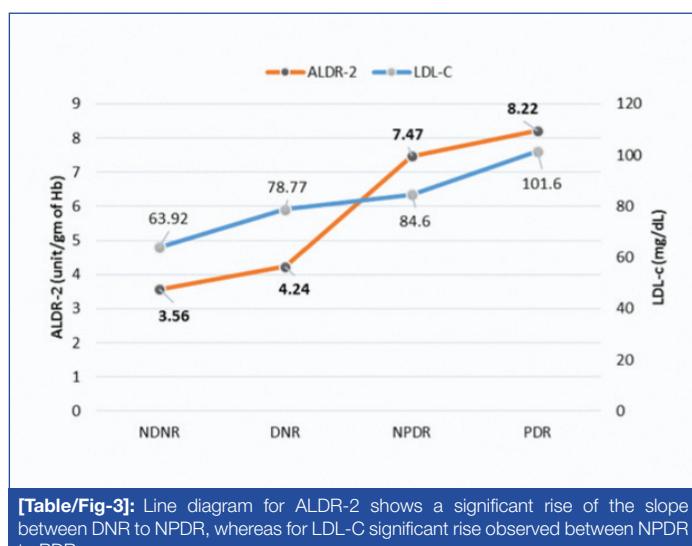
STATISTICAL ANALYSIS

The results are presented in Mean±SD. The Chi-square test was applied to compare the age between groups and subgroups. The unpaired t-test was applied to compare the continuous study parameters between two groups and subgroups. The p-value<0.05 was considered significant. All the analysis were carried out using IBM SPSS Statistics for Windows, version 22.0 (Armonk, NY: IBM Corp).

RESULTS

Outcomes of current study based on mean age, fasting blood glucose level, ALDR-2 enzyme level, and various traditional lipid markers with respect to NDNR, DNR, NPDR and PDR subgroups are summarised in [Table/Fig-2].

In DNR group fasting and postprandial blood sugar along with LDL-C and HDL-C were significantly deranged, as compared to NDNR group. However, ALDR-2 was non-significantly increased in DNR than NDNR. A significantly higher activity of ALDR-2 was observed for NPDR subjects as compared to DNR subjects (p-value <0.001) as evident in [Table/Fig-2].



[Table/Fig-3]: Line diagram for ALDR-2 shows a significant rise of the slope between DNR to NPDR, whereas for LDL-C significant rise observed between NPDR to PDR.

[Table/Fig-3] shows a higher value of LDL-C and ALDR-2 in NPDR and PDR group than DNR and NDNR group. However, PDR group showed a non-significant increase in ALDR-2 level while a significantly higher level of LDL-C as compared to NPDR group.

DISCUSSION

The present study conducted in diabetic patients to know the role of dyslipidemia in the progression of NPDR to PDR. The most likely reason for a significantly higher activity of ALDR-2 in NPDR than DNR subjects could be the activation of ALDR precursor gene in diabetics causing retinopathy [18]. However, the difference in ALDR-2 level in NPDR and PDR subjects was non-significant; thus, the ALDR-2 activity might not play a major role in the progression of NPDR to PDR. Further, out of the five traditional lipid markers in our study, LDL-C was significantly high in PDR than NPDR group signifying the role of dyslipidemia in the progression of DR.

Type 2 Diabetes occurs due to insulin resistance i.e., secretion of insulin is either normal or increased, however, this increased insulin level is unable to enter the cell through tyrosine kinase receptors present on cell membrane [19]. Most oral hypoglycaemic either stimulate the beta cells of the pancreas causing the enhanced release of insulin or increase the sensitivity of insulin receptors to overcome the insulin resistance. Over the years or decades, these pancreatic beta cells and receptor protein are exhausted and depleted, that cannot be resynthesized by medicines; thus, a diabetic individual could not revert to a normal physiological condition. These individuals will develop DR as a microvascular complication if they have a genetic susceptibility for ALDR precursor gene. Once DR is established, it will progress to NPDR after prolonged hyperglycaemic condition due to the activation and increased production of ALDR-2 enzyme [3]. NPDR progresses to PDR stage if proper treatment is not given to the patient.

The possible mechanism for progression of DR due to dyslipidemia could be the activation of PKC and AGE pathways [20]. Typically, PKC is involved in a number of physiological processes in our body, but its up regulation contributes to the pathogenesis of DR by: (i) differential synthesis of proteins and remodelling of Extracellular Matrix (ECM); and (ii) increasing angiogenic factors — both of which cause capillary occlusion and leukostasis due to endothelial dysfunction and altered retinal blood flow [21]. PKC family has 10 isoforms of which 1/2 isoform is strongly associated with the progression of DR [21]. De novo synthesis of Diacylglycerol (DAG) ensues as a by-product, when hyperglycaemia leads to an increase in glucose flux through the glycolysis pathway. Furthermore, the accumulated long-chain Fatty Acids (FAs) due to dyslipidemia instantaneously transforms into DAG. This disproportionate production of DAG activates PKC enzymes [22].

On the other hand, AGEs are constantly produced, non-enzymatically, from reducing sugars and lipoproteins, at a slower rate since the embryonic development that gets accumulated over time. Their production is markedly augmented in diabetes because of the sustained hyperglycaemia [23]. Carboxyethylpyrrole [24] and malondialdehyde [25] are the two types of AGEs product associated with the pathogenesis of DR. Their serum levels significantly correlate with the severity of the retinopathy. The interaction of AGEs with their specific cell surface G-Protein Coupled Receptors (GPCR) like RAGE, galectin-3, CD36, and the macrophage scavenger receptor has been linked with the advancement of DR [26]. An animal model study done on rats by Hammes HP et al., indicated that the accumulation of AGEs in the retinal capillaries and loss of pericytes cells occurred within 26 weeks of the diabetes establishment. Treatment with AGEs inhibitor significantly reduced AGEs accumulation, microaneurysms, acellular capillaries and pericyte cell loss in diabetic individuals [27].

Our results is in agreement with few previous studies [28-34] that indirectly correlated the significance of LDL-C, TC and TG in the

progression of DR. ETDRS Report 22 (1996) showed that the patients who had increase serum TC or higher serum LDL-C levels were more prone to have retinal hard exudate [28]. The Atherosclerosis Risk in Communities Study (2002) concluded that the presence of retinal hard exudates was associated with the plasma LDL-C and plasma lipids [29]. The Cardiovascular Health Study (2002) demonstrated that DR was associated with higher mean Systolic Blood Pressure (SBP), plasma TC, LDL-C and Cardiovascular Disease (CVD) presence; this study further specified that elevations in plasma LDL-C might contribute to the hard exudate deposition in the retina of diabetics [30].

A North Indian observational case-control study associated NPDR with Clinically Significant Macular Oedema (CSME) and dyslipidemia. On univariate analysis, retinal hard exudates were significantly associated with SBP, serum TC, serum LDL-C, and serum TG levels. However, on linear regression analysis, serum TC and serum LDL-C was established as an independent risk factor affecting the density of retinal hard exudate [31]. A prospective study of serum lipids and risk of diabetic macular oedema in Type 1 diabetes found that TC to HDL-C ratio and LDL-C separately were linked to an increased threat of CSME and retinal hard exudate; it was concluded that treatment with lipid-lowering agent among Type 1 diabetic subjects might decline the risk of the CSME [32]. SN-DREAMS Report Number 13 revealed that high serum LDL-C, non-HDL-C, and TC was related to CSME and non-CSME [33]. Another study underlined the association of dyslipidemia with macular oedema and retinal hard exudates in Type 2 diabetes patients. A statistically significant high levels of TG, TC, total lipids and cholesterol ester was found in the diabetic patients with manifested maculopathy as compared to those without diabetic maculopathy [34].

Some treatment trial studies also signify the role of dyslipidemia in the progression of DR. The Action to Control Cardiovascular Risk in Diabetes (ACCORD) study concluded that intensive combination treatment of dyslipidemia along with intensive glycaemic control reduced the rate of DR progression [35]. Gordon B et al., reported the effects of lipid lowering on DR and concluded that reduction in TC and LDL-C might have a beneficial effect on background retinopathy in diabetics [36]. Sen K et al., reported that an HMG-CoA reductase inhibitor significantly retarded the progression of retinopathy in diabetics with hypercholesterolemia [37].

LIMITATION

We estimated serum lipid level and correlated it with progression of DR, which is an indirect method. Study could have been more precise if the levels of various lipid markers would have been estimated in the vitreous humor that could be collected during vitrectomy indicated for patients of diabetic retinopathy.

CONCLUSION

Diabetic Retinopathy will occur in all genetically susceptible diabetics eventually. The persistent hyperglycaemia stimulates diverse molecular mechanisms, including increased ALDR-2 activity that results in the establishment of DR. However, the progression of DR is largely regulated by the mechanisms that are likely to be mediated by dyslipidemia rather than hyperglycaemia alone. Based on the results of our study, it can be concluded that the lipid lowering drugs as an adjunctive to the traditional diabetes therapy that involves rigorous blood sugar and blood pressure control, might retard the progression of DR.

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PARTICULARS OF CONTRIBUTORS:

- Assistant Professor, Department of Physiology, Hind Institute of Medical Sciences, Barabanki, Uttar Pradesh, India.
- Professor, Department of Physiology, King George's Medical University, Lucknow, Uttar Pradesh, India.
- Senior Resident, Department of Physiology, Hind Institute of Medical Sciences, Barabanki, Uttar Pradesh, India.
- Medical Officer, Employee State Insurance Corporation of India, Lucknow, Uttar Pradesh, India.
- Professor, Department of Ophthalmology, King George's Medical University, Lucknow, Uttar Pradesh, India.
- Professor, Department of Medicine, Babu Banarsidas Dental University, Lucknow, Uttar Pradesh, India.

NAME, ADDRESS, E-MAIL ID OF THE CORRESPONDING AUTHOR:

Dr. Mayank Agarwal,
Hind Institute of Medical Sciences, Safedabad, Barabanki, Uttar Pradesh, India.
E-mail: ma.gsvm@gmail.com

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