

The Effect of Darapladib Therapy for the Expression of Lp-PLA₂ in Dyslipidemia and Type 2 Diabetes Mellitus Atherosclerosis Model

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ABSTRACT

Atherosclerosis is the main cause of mortality and morbidity globally. *Lipoprotein-associated phospholipase A₂* (Lp-PLA₂) activity is suspected to have a significant role in atherosclerosis. 50 Sprague-Dawley Rats were divided into five groups: normal, dyslipidemia, Type 2 diabetes mellitus (T2DM), dyslipidemia with darapladib administration and T2DM with darapladib administration. These groups were divided into two serial times: 8 and 16 weeks. mRNA Lp-PLA₂ was measured from blood and aortic tissue extraction. Aortic tissue Lp-PLA₂ was measured by immunofluorescence. Lp-PLA₂ expression in aortic tissue was consistently increased in dyslipidemia and T2DM. The expression of Lp-PLA₂ enzymatic was significantly suppressed ($p < 0.05$) with the administration of darapladib especially in 8 weeks groups in both dyslipidemia and T2DM. The administration of darapladib in dyslipidemia and T2DM didn't significantly suppress the expression of mRNA Lp-PLA₂ in blood and aortic tissue. The failure of genetic expression suppression of Lp-PLA₂ was found in both 8 weeks and 16 weeks groups. The expression of Lp-PLA₂ protein also showed an inclined difference between dyslipidemia and T2DM. These results showed that administration of darapladib significantly decreased Lp-PLA₂ protein but prone to increase the expression of mRNA Lp-PLA₂ in blood and aortic tissue in dyslipidemia and T2DM model.

INTRODUCTION

Cardiovascular disease (CVD) has become a serious global health problem and the leading cause of death in both modern and developing countries (Zhang *et al.*, 2008). Atherosclerosis is the main cause of mortality and morbidity globally. It is predicted that prevalence of CVDs will increase quickly in developing countries (Wihastuti *et al.*, 2014, 2015). Atherosclerotic plaque formation is a long-term process and often without clinical symptoms. Atherosclerosis development is highly associated with lipid metabolic disorders and inflammation. Oxidized LDL (OxLDL) has been widely demonstrated to have an important role in vascular inflammation (Zhang *et al.*, 2013) Ox-LDL is capable of inducing monocyte adhesion to endothelial cells (Heriansyah *et al.*, 2015). oxLDL initiates chemoattractant release, which in this

way attracts macrophage migration into the lesion, and triggers the formation of reactive oxygen species (ROS), leading to oxidative stress, and the death of vascular smooth muscle cells (Heriansyah *et al.*, 2017). *Lipoprotein-associated phospholipase A₂* (Lp-PLA₂) activity that highly correlated with oxLDL is suspected to have a significant part in atherosclerosis development and also contributes to plaque destabilization process in different pathways. Several epidemiology studies with 80.000 subjects have clearly shown that the higher level of Lp-PLA₂ in the circulation is associated with higher risk of cardiovascular event, either in primary prevention and in secondary prevention cohort (Vepa *et al.*, 1999; Unno *et al.*, 2000; Von der Thusen, 2003; Wæhre *et al.*, 2004; Tsimikas *et al.*, 2009; van Dijk *et al.*, 2013). The physiology and potential function of Lp-PLA₂ in atherosclerosis pathogenesis is still controversial and far from achieving an understanding. Several pieces of evidence from previous experiments showed that there are two parts of the Lp-PLA₂ enzyme, which are opposites of one another, as an anti-inflammatory agent and pro-inflammatory agent. As an anti-inflammatory agent, the Lp-PLA₂ enzyme has the ability to

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hydrolyze mediators that have a potential role in atherogenesis (such as oxLDL and *platelet activating factor* (PAF)). In the same time, hydrolyzed products of this molecule which are mediated by Lp-PLA₂ stimulate atherosclerosis lesion either in the human and animal trial. Analysis using *reverse transcriptase-polymerase reaction* (RT-PCR) showed an increased expression of mRNA Lp-PLA₂ in human atherosclerotic lesions (Tsimihodimos *et al.*, 2002; Wæhre *et al.*, 2004).

In a recent study, Lp-PLA₂ selective inhibitors [darapladib (DP)] have become a potential therapeutical candidate to decrease cardiovascular accident (Wæhre *et al.*, 2004). Several epidemiology studies with CAD patients as a subject showed that

DP has the ability to suppress Lp-PLA₂ activity and inflammatory markers (Tsimihodimos *et al.*, 2002; Tsimikas *et al.*, 2009; Tousoulis *et al.*, 2013; Tian, 2014; White, 2014; Taruya *et al.*, 2015). A clinical trial using animal treated by DP also succeed to suppress Lp-PLA₂ activity and advanced stage atheroma plaque size (Szasz *et al.*, 2013). However, two large clinical trial in stable CAD patients (STABILITY) and acute coronary syndrome patients (SOLID-TIMI 52) reported that administration of DP did not lower major cardiovascular accident, and even in fact increase the risk of Acute Myocardial Infarction (AMI) (Tsimikas *et al.*, 2009).

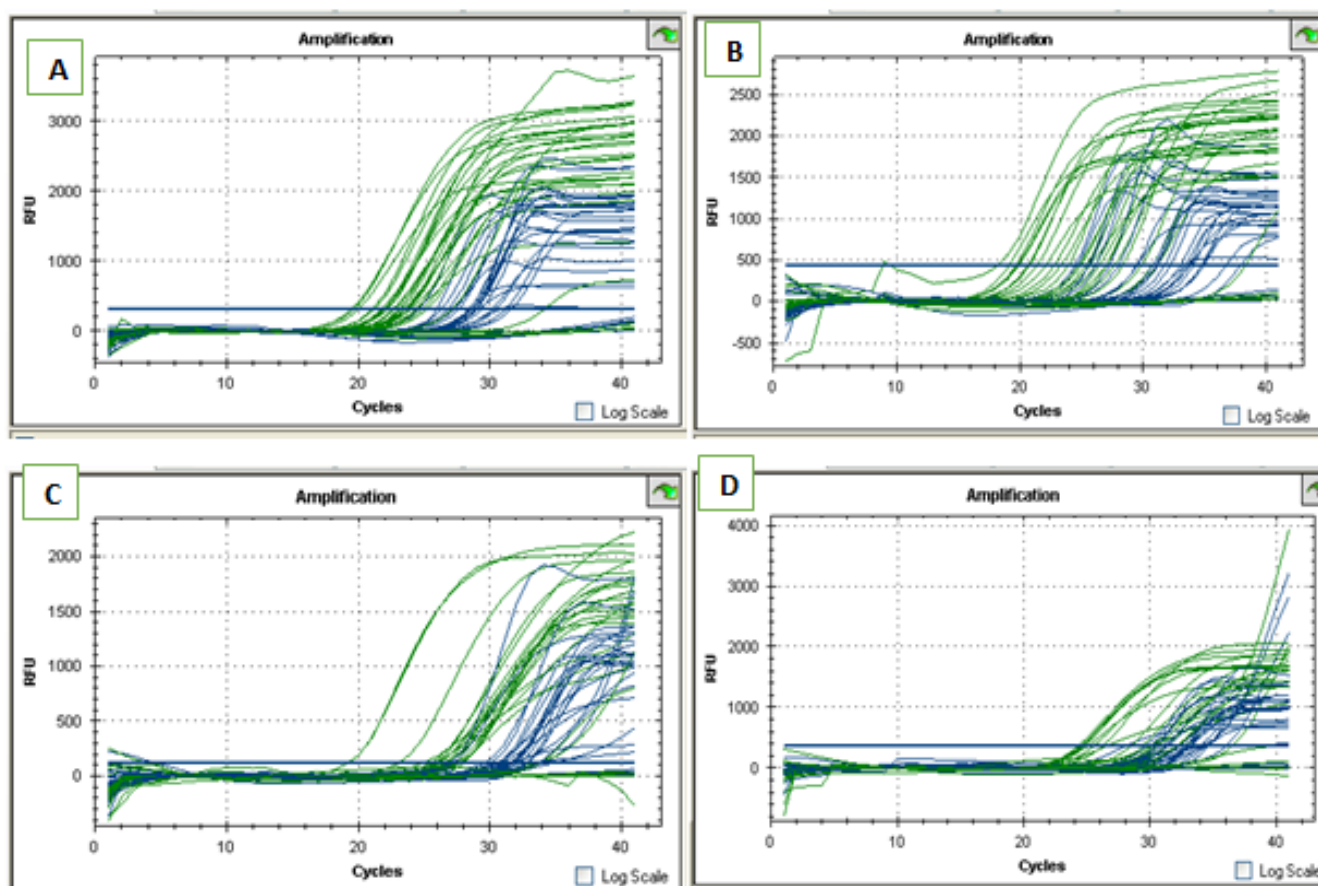


Fig. 1: Lp-PLA₂ and GAPDH Amplification; (a) Aorta Tissue Sample 8 Weeks, (b) Aorta Tissue Sample 16 weeks, (c) Blood 8 weeks, (d) Blood 16 weeks; (Green: GAPDH, Blue: Lp-PLA₂).

METHODS

Study group

This study used 50 4-weeks male Sprague-Dawley and weighted around 150-200 grams. Samples were obtained from Bogor Agricultural University, Bogor, Indonesia. These rats were divided into five groups; normal group (N); Dyslipidemia group which fed with High Fat Diet (DL); Type 2 DM Model group (T2DM) which fed with High Fat Diet and injected with Streptozotocin (STZ) intraperitoneal low dose 35 mg/KgBW, Dyslipidemia with Darapladib administration group (DLDP) and

type 2 DM model with Darapladib administration group (DMDP). Each group was divided into two serial times (8 weeks and 16 weeks) and consist of 5 rats. Darapladib was obtained from Glaxo Smith Kline. Samples were given Darapladib orally 20mg/KgBW once a day according to the time-serial groups given. Normal rats' food contained 3.43 kcal/g total energy calories (67% carbohydrate, 21% protein and 12% fat), while the HFD contained 5.29 kcal/g total calorie energy (58% fat, 17% carbohydrate and 25% protein). 30 grams of food were given for each rat every day. Parameters measurement were done at the Central Laboratory of Biological Sciences, Brawijaya University.

Blood glucose level measurement

Mouse blood glucose level was measured using enzymatic method after fasting for 16 hours, after 4 weeks of HFD administration in T2DM mouse group before low dose STZ injection, 7 days after STZ injection, and before dissection. A blood sample was collected from the top of mouse tail (vena lateralis). Blood glucose level measurement using glucometer (GlucoDrCo.Ltd Korea). Using eligible measurement scale mg/dl.

Plasma insulin level measurement and insulin resistance

Insulin level was measured in T2DM groups before 30 mg/kgBW of STZ injection to confirm insulin resistance. With *enzyme-linked immunosorbent assay* (ELISA) method. Plasma insulin level normal value is <2500 pg/mL. Insulin resistance (IR) in the mouse was measured with HOMA-IR (*Homeostatic Model Assessment-Insulin Resistance*) formula as described below:

$$[\text{HOMA} - \text{IR}] = \frac{\text{FBS} \times \text{FINS}}{14,1}$$

Information:

HOMA-IR: *Homeostatic Model Assessment-Insulin Resistance*

FBG: *Fasting Blood Glucose* (mmol/mL)

FINS: *Fasting blood insulin* ($\mu\text{U/mL}$)

Insulin resistance is determined if HOMA-IR value >1.716.

Measurement of mRNA expression *lipoprotein-associated phospholipase A₂* of aortic tissue and blood

Messenger Ribonucleic acid (mRNA) from blood (whole blood sample) and aortic tissue extraction were prepared for quantitative examination with *polymerase chain reaction* (PCR) using primer and probe from *Applied Biosystems 7300 Real-Time PCR system*, Foster City, California (Goncalves *et al.*, 2012; van Dijk *et al.*, 2013).

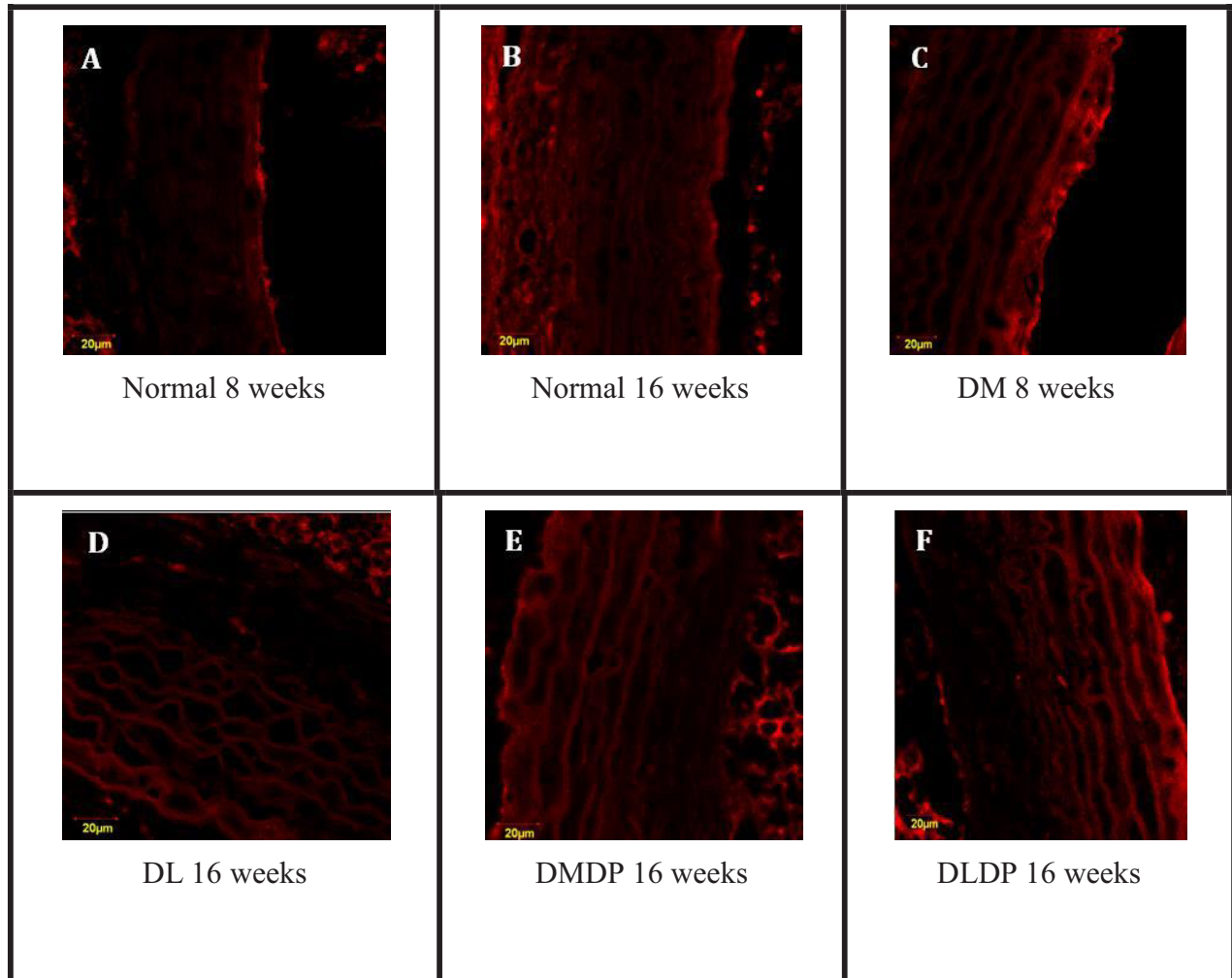


Fig. 2: Aorta with Rhodamin Secondary Antibody on N8 Groups, N16, DM8, DL16, DMDP8, and DLDP 16. (A) Normal mouse groups of 8 weeks (N8); (B) Normal Mouse Groups of 16 weeks (N16); (C) DMT2 Mouse Groups of 8 weeks (DM8); (D) Dyslipidemia Mouse Group of 16 weeks (DL16); (E) DMT2 Mouse groups given DP administration 30 mg/kg body weight for 16 weeks (DMDP16); (F) Dyslipidemia mouse groups given DP administration 30 mg/Kg body weight for 16 weeks (DLDP16).

Aortic tissue that has been separated from animal carcass and was stored in a cooler (-70°C) until used. mRNA from early-stage atherosclerosis plaque that was stored in an isolation was using *Fast Track mRNA isolation kit* (Invitrogen) and *cDNA synthesis kit* (Invitrogen) following a factory published procedure. Meanwhile, extracted RNA from blood was isolated using *Trizol reagent* (GIBCO BRL) and reverse transcriptase from DNase I (GIPCO BRL)-treated total RNA using Superscript II and *random hexamer primer* (GIPCO BRL) following factory published procedure (Hatoum *et al.*, 2010).

cDNA samples (approximately 5 μL each sample) were analyzed to see Lp-PLA₂ expression and Glyceraldehyde 3-phosphate dehydrogenase (GAPDH) *housekeeping* genes using real-time quantitative reverse transcriptase-polymerase chain reaction (RT-PCR) with fluorescent TaqMan 5' nuclease assay. Taqman oligonucleotide primers and probes examination were designed using *Primer Express software version 1.0* (PE Biosystems). Each TaqMan hydrolyzed probe consist of fluorescent reporter dye 6-carboxyfluorescein (FAM) that have a covalent association with 5' end of the oligonucleotide I and also the quencher dye 6-carboxytetramethylrhodamine (TAMRA), which is attached on the 3' end using a group linker (PE Biosystems).

PCR (5' > 3' nuclease assay) were performed in *microamp optical 96-well reaction plates* with ABI PRISM 7300 *sequence detection system* for thermal cycle and *real time fluorescence* measurement (PE Biosystems). Each 25 μL reaction consist of $1 \times 1 \times$ TaqMan Universal PCR Master Mix (10 mmol/L Tris HCL (pH 8.3) 50 mmol/L KCL, 10 mmol/L EDTA, 60 nmol/L passive reference dye 1 (6-carboxy-X-rhodamine), 0.2 mmol/L dATP, 0.2 mmol/L dCTP, 0.2 mmol/L dGTP, 0.4 mmol/L dUTP, 5.5 mmol/L MgCl₂, 8% glycerol, 0.625 U AmpliTaq Gold DNA polymerase and 0.25 U AmpErase uracilN-glycosylase), 300 nmol/L forward primer, 300 nmol/L reverse primer, 100 nmol/L TaqMan Quantification probe and 5 μL template with 20 μL mineral oil (Promega).

Primary Sequence

Lp-PLA₂ F: 5'-CCACCCAAATTGC-ATGTGC-3'
 R: 5'-GCCAGTCAAAGGATAAACCACAG-3'
 GAPDH F: 5'-GCCAAGGTCATCCATGACAAC-3'
 R: 5'-GGGGCCATC-CACAGTCTTC-3'
 TaqMan Probe (FAM-5' > 3'-TAMRA)
 Lp-PLA₂: 5'-TTCTGCCTCTGCGGCTGCCTG-3'
 GAPDH: 5'-CTCATGACCACA-GTCCATGCCATCACT-3'

Certain circumstances for the reaction are described as follow: 50°C for 2 minutes, 95°C for 10 minutes, and continued for 40 cycles at 95°C for 15 seconds and 60°C for 1 minute.

Continued analyses were performed from external data of *sequence detector software* using *Microsoft Excel* software. Lp-PLA₂ expression quantitative value that obtained from *sequence detector* will be divided with each GAPDH sample quantitative value to achieve a normal value of Lp-PLA₂ for each sample. This value will be divided with the lowest value that is achieved to get the increased value of each sample. The obtained results were analyzed using $2^{-\Delta\Delta\text{Ct}}$ method (Hatoum *et al.*, 2010; Goncalves *et al.*, 2012).

Expression of Lp-PLA₂ enzyme measurement in form of protein in aortic tissue

Lp-PLA₂ in aortic tissue was measured by immunofluorescence that was previously fixed with PHEMO buffer (68 mM PIPES, 25 mM, HEPES, pH 6.9, 15 mM EGTA, 3 mM MgCl₂, 10% [v/v] dimethyl sulfoxide containing 3.7% formaldehyde and 0.05% glutaraldehyde) and were processed by immunofluorescence labelling with anti-rat antibody Lp-PLA₂ using rhodamine secondary antibody (BIOS Inc., Boston, MA, USA). This parameter was observed with confocal laser scanning microscopy (Olympus Corporation, Tokyo, Japan) and was quantitatively analyzed using Olympus FluoView software (version 1.7A; Olympus Corporation).

Ethics

We obtained ethical approval for the animal treatment and experimental processes in this study from the Animal Care and Use Committee Brawijaya University Number 229-KEP-UB.

RESULT

Lipid profile and fasting blood glucose level

Total cholesterol level and non-HDL cholesterol in 8 weeks normal groups are lower than experimental group. This result was also shown in 16 weeks normal groups, there is a lower trend compare to the experimental group. Total cholesterol and non-HDL cholesterol level in dyslipidemia and type 2 diabetes mellitus are decreased with darapladib administration. The decrease in total cholesterol level with darapladib administration on 16 weeks is lower than 8 weeks. Non-HDL cholesterol level is lower parallel with administration period. Treatments with darapladib administration for 8 weeks and 16 weeks have total cholesterol and non-HDL cholesterol level near normal group level.

HDL level on 8 weeks normal group is higher compared with experimental groups. Moreover, 16 weeks normal group tends to increase compared with the experimental group. Dyslipidemia and type II DM are affecting the increment of HDL level in experimental groups with darapladib administration. Darapladib administration for 8 weeks and 16 weeks have HDL level near normal group level.

Fasting blood glucose level on 8 weeks normal groups is lower than experimental group. Moreover, 16 weeks normal groups tend to be lower than the experimental groups. Mouse with dyslipidemia and type II DM have a decrease fasting blood glucose level in the experimental group with darapladib administration. The experimental group with darapladib administration for 8 weeks and 16 weeks have fasting blood glucose level near normal groups level.

Insulin resistance measurement using HOMA-IR formula is conducted after 4 weeks in either standard feeding or HFD feeding. Interpretation result from HOMA-IR calculation on the mouse as the experimental animal model is if the result > 1.710 then insulin resistance is diagnosed with sensitivity 83.87% and specificity 80.56% (95% *confidence interval*). HFD feeding mouse groups are experiencing insulin resistance compared with standardized feeding groups (normal diet) that does not experience

insulin resistance. According to the calculation of groups with diabetes mellitus experience insulin resistance before DO administration. Insulin resistance is an indication for T2DM. This show that T2DM model in these trials are what we are expected. All of the results above can be seen in Table 1.

Table 1: Lipid Profile and Fasting Blood Glucose Level.

Variable	Groups	8 weeks	16 weeks
		Mean \pm SD	Mean \pm SD
Total Cholesterol	Normal	72.80 \pm 4.05 ^b	56.56 \pm 5.43 ^a
	DL	109.56 \pm 12.25 ^{cd}	100.57 \pm 22.64 ^c
	DLDP	79.57 \pm 10.54 ^b	96.96 \pm 18.97 ^c
	DM	123.00 \pm 2.86 ^d	111.72 \pm 7.30 ^{cd}
	DMDP	97.96 \pm 1.70 ^c	98.85 \pm 3.25 ^c
HDL	Normal	34.74 \pm 8.31 ^f	35.77 \pm 1.68 ^f
	DL	19.16 \pm 0.30 ^{de}	17.55 \pm 0.85 ^{cde}
	DLDP	20.62 \pm 0.20 ^c	13.00 \pm 1.82 ^b
	DM	4.96 \pm 0.41 ^a	13.96 \pm 0.87 ^{bc}
	DMDP	15.94 \pm 1.21 ^{bed}	20.79 \pm 2.76 ^c
Non HDL	Normal	49.83 \pm 5.06 ^b	19.24 \pm 3.67 ^a
	DL	88.20 \pm 3.08 ^{de}	98.14 \pm 11.43 ^c
	DLDP	60.34 \pm 2.64 ^c	96.51 \pm 17.06 ^{de}
	DM	95.53 \pm 8.66 ^{de}	88.25 \pm 6.23 ^{de}
	DMDP	85.92 \pm 6.84 ^d	61.52 \pm 6.03 ^c
Fasting Blood Glucose	Normal	91.60 \pm 7.16 ^{ab}	79.60 \pm 14.64 ^a
	DL	114.60 \pm 7.73 ^{bc}	118.60 \pm 15.65 ^{bcd}
	DLDP	110.20 \pm 6.18 ^{abc}	99.60 \pm 19.03 ^{abc}
	DM	128.00 \pm 15.02 ^{cd}	147.80 \pm 58.23 ^d
	DMDP	103.60 \pm 13.72 ^{abc}	101.80 \pm 19.07 ^{abc}

Analysis of two mean different groups with *Duncan Multiple Range Test* (DMRT), there is a similar letter notation, it means that the difference between two mean values is not significant. * $\alpha = 0.05$.

Aortic tissue and blood mRNA Lp-PLA₂ and aortic tissue protein Lp-PLA₂

In 8 weeks group, mRNA Lp-PLA₂ blood expression in experimental groups are always lower compare with normal groups. However, mRNA Lp-PLA₂ blood expression in the experimental group are always higher compare with the normal group in 16 weeks groups.

mRNA Lp-PLA₂ tissue expression in experimental groups tends to be lower in 8 weeks normal group. A different result is found in 16 weeks normal groups, mRNA Lp-PLA₂ tissue expression in the experimental group always higher than normal groups.

In 8 weeks normal groups, Lp-PLA₂ protein level is lower than the experimental group. 16 Weeks normal groups also tend to be lower than experimental group. Descriptively, in dyslipidemia and type II DM mouse, 8 weeks of Lp-PLA₂ protein level tissue were decreased with darapladib administration, and in 16 weeks diabetes mellitus group were experiencing a higher level. 16 weeks of Lp-PLA₂ protein level tissue were lower if we compared it with 8 weeks group. Darapladib administration in 8

and 16 weeks diabetes mellitus groups has Lp-PLA₂ protein level near normal level.

DISCUSSION

The newest information from this study is that we able to achieve a more complete and integrated data for mRNA Lp-PLA₂ aortic tissue and blood expression pattern in two risk factors model of cardiovascular disease, which are dyslipidemia and type II diabetes mellitus by following chronicity theory of atherosclerotic process using time series experimental design (animal model). The animal model used in this experiment is using the combination of low doses of STZ and high-fat diet could induce metabolic syndrome mimicking human criteria (Rohman *et al.*, 2017). This experimental has Lp-PLA₂ activity data in aortic tissue with protein/enzyme form. Which is far more different with a previous study that its goal is to only identify Lp-PLA₂ expression's degree only on one side (either only on aortic tissue or blood) and also without time series, and in turn Lp-PLA₂ expression pattern data cannot be achieved.

Table 2: Lp-PLA₂ Expression.

Variable	Groups	8 weeks	16 weeks
		Mean \pm SD	Mean \pm SD
mRNA Lp-PLA ₂ Aortic tissue	Normal	1/1.00 \pm 0.00	1/1.00 \pm 0.00
	DL	1/1.13 \pm 0.00	1/0.62 \pm 0.00
	DLDP	1/0.37 \pm 0.00	1/0.48 \pm 0.00
	DM	1/1.08 \pm 0.00	1/0.72 \pm 0.00
	DMDP	1/2.00 \pm 0.00	1/0.48 \pm 0.00
mRNA Lp-PLA ₂ Blood	Normal	1/1.00 \pm 0.00	1/1.00 \pm 0.00
	DL	1/0.37 \pm 0.00	1/2.27 \pm 0.00
	DLDP	1/0.17 \pm 0.00	1/5.12 \pm 0.00
	DM	1/0.59 \pm 0.00	1/6.69 \pm 0.00
	DMDP	1/0.42 \pm 0.00	1/1.83 \pm 0.00
Lp-PLA ₂ Protein Aortic tissue	Normal	788.96 \pm 49.73 ^{ab}	752.59 \pm 153.47 ^{ab}
	DL	1220.52 \pm 200.61 ^d	804.61 \pm 116.52 ^{ab}
	DLDP	1069.05 \pm 93.59 ^{cd}	599.63 \pm 32.30 ^a
	DM	925.13 \pm 405.28 ^{bc}	751.65 \pm 240.29 ^{ab}
	DMDP	820.86 \pm 152.10 ^{abc}	798.5 \pm 108.67 ^{ab}

Trials result of two mean value with *Duncan's Multiple Range Test* (DMRT), there are two similar notations, which means the difference between two mean value are not significant * $\alpha = 0.05$.

This study found that aortic tissue and plasma mRNA Lp-PLA₂ expression profile is different between dyslipidemia and type II diabetes mellitus conditions. Lp-PLA₂ protein expression also showed an activity which tends to be different in such conditions. Much more surprising findings is aortic tissue mRNA Lp-PLA₂ expression pattern is similar with aortic tissue Lp-PLA₂ protein expression, which showed that Lp-PLA₂ activity either on both transcriptase level and on enzymatic activity in tissue are much more dominant on the early phase of atherogenesis (in this study shows a higher level of expression on 8 weeks, experimental groups). This expression pattern was seen on both dyslipidemia and type II diabetes mellitus condition. This study supports a

previous scientific proof (*in vitro* study) that found Lp-PLA₂ formation is dramatically increased in an arterial blood vessel at the time of monocyte differentiation into macrophage which occurs in a much early (Cao *et al.*, 1998; Ferguson *et al.*, 2012).

Contradictive expression patterns are found in blood mRNA Lp-PLA₂ expression pattern. mRNA Lp-PLA₂ expression pattern is significantly increased in 16 weeks experimental groups (both dyslipidemia and type II diabetes mellitus conditions). This findings support previous an *in vitro* study findings which found that the early phase of inflammation process (monocyte to macrophage pre-transformation phase), mRNA Lp-PLA₂ expression level are so low to be detected in the blood, mRNA Lp-PLA₂ expression will significantly increase in blood after macrophage maturation phase (longer observation period phase) (Ferguson *et al.*, 2012). Even though there is a possibility that Lp-PLA₂ is produced by macrophage locally in aortic tissue can also contribute to half of circulating Lp-PLA₂. It is not right to use the circulating Lp-PLA₂ level as an independent biomarker which represents Lp-PLA₂ action in atherosclerotic tissue. In other words, mRNA expression and Lp-PLA₂ protein activity in the blood is a poor predictor of PLA₂G7 action on the arterial plaque (Ferguson *et al.*, 2012). This result showed a supporting evidence for that *in vitro* study and support the pro-inflammatory pro-atherogenic role of Lp-PLA₂.

This study results showed that DP administration therapy tends to consistently decrease enzymatic activity of Lp-PLA₂ in aortic tissue either on both dyslipidemia or Type II diabetes mellitus. This finding is consistent with previous *in vivo* study result, that reported a significant decrease in enzymatic activity of Lp-PLA₂ on mouse serum (Wang *et al.*, 2011). Other *in vivo* study also reported that DP administration can decrease Lp-PLA₂ enzymatic activity in plasma and aortic blood vessel of animal's trial (Wilensky *et al.*, 2008). A different report is seen in a human study that showed DP therapy does not decrease Lp-PLA₂ plasmic activity significantly but change atheroma plaque characteristic from fragile plaque with wide necrotic lipid core into much more stable plaque with the smaller necrotic core (Serruys *et al.*, 2008).

Different with suppression of Lp-PLA₂ protein activity as mentioned above, DP administration therapy in this study is giving an interesting result in terms of mRNA Lp-PLA₂ expression in aortic tissue and blood. DP administration has the ability to decrease Lp-PLA₂ expression in blood, in type II Diabetes Mellitus conditions with 16 weeks observation period. Meanwhile, in dyslipidemia condition DP administration therapy tend to increased mRNA Lp-PLA₂ expression in blood. Different conditions are seen in tissue mRNA Lp-PLA₂ expression, decrease expression of mRNA Lp-PLA₂ only found in dyslipidemia conditions with 8 weeks observation period. In type II diabetes mellitus conditions with DP therapy, mRNA Lp-PLA₂ expression of aortic tissue increased significantly in 8 weeks observation period groups. This result phenomenon shows that both conditions (dyslipidemia and type II diabetes mellitus) are not a role for a different pattern on both groups. mRNA Lp-PLA₂ expression pattern in treatment group with DP administration in this study support previous *in vitro* result that showed with administration of Lp-PLA₂ inhibitor (SB435495) to oxLDL results in undetected enzymatic activity of Lp-PLA₂, however there is an increased of mRNA Lp-PLA₂ expression as an effect of oxPC that are hydrolysed in oxLDL which then cause an up-regulatory effect of oxLDL to stimulate

much mRNA Lp-PLA₂ expression (because of SB-oxLDL consist of much more oxPC as the result of less hydrolysed oxPC in those complex) (Wang *et al.*, 2010). Another study in an *in vivo* model (another animal model) also reported that with the administration of DP does not decrease mRNA Lp-PLA₂ expression significantly, however with the administration of RNA Lp-PLA₂ inhibitors (Lp-PLA₂ RNAi) using lentivirus were a success to significantly decreasing mRNA Lp-PLA₂ expression. Enzymatic activity of Lp-PLA₂ degree that was decreased in plasma is not different between DP groups and Lp-PLA₂ RNAi. Usable explanation to analyze this *in vivo* trial and previous two clinical trials is that DP works on such level as a reversible inhibitor between ox LDL with Lp-PLA₂ interactions, not on genetic transcription level. Inhibitions with DP can increase the number of mRNA Lp-PLA₂ expression level as a result of much more un-hydrolyzed oxPC. Our *in vivo* study result can also explain as an indirect answer to all three major clinical trials (STABILITY, SOLID-TIMI 52 and CIBIS 2) to observed DP effectivity in lowering inflammation response and cardiovascular morbidity and mortality. Inflammation response that involved Lp-PLA₂ is much more complex in nature and is not linear and not easy to predict.

CONCLUSION

Lp-PLA₂ protein expression in aortic tissue was consistently increased in both dyslipidemias and type 2 diabetes models. It was significantly seen in 8 weeks group compared to 16 weeks group. Similar phenomenon confirmed by the treatment using a selective blocker of Lp-PLA₂ (darapladib/DP). The expression of Lp-PLA₂ enzymatic was significantly suppressed ($p < 0.05$) with the administration of darapladib especially in 8 weeks groups in both dyslipidemia and type 2 diabetes models. This happened due to the high inflammatory response that appears at the beginning of the period that followed by high production of Lp-PLA₂ by its cellular source, namely macrophages. The administration of selective blockers Lp-PLA₂ (darapladib) *in vivo* models of dyslipidemia and type 2 diabetes in this study did not significantly suppress the relative expression of mRNA Lp-PLA₂ in both blood and aortic tissue. The failure in suppressing the genetic expression of Lp-PLA₂ was found in both 8 weeks and 16 weeks groups. The expression of Lp-PLA₂ protein also showed an inclined different in activity between dyslipidemia and type 2 diabetes. The surprising finding was the pattern of mRNA Lp-PLA₂ expression of aortic tissue was similar to the pattern of Lp-PLA₂ protein expression of aortic tissue which shows that the activity of Lp-PLA₂ in both transcription and protein level in tissues are more dominant in the phase before atherosclerosis process (in this study, it shows the level of expression was higher in the 8 weeks group). This expression pattern was shown in both dyslipidemias and type 2 diabetes models. This study supports the previous *in vitro* research that found that the formation of Lp-PLA₂ increased dramatically in arterial blood vessel walls tissue during the process of differentiation of monocytes into macrophages that occurs in an early phase.

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CONFLICTS OF INTEREST

None.

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