

# Glycoxidized Low-Density Lipoprotein Enhances Monocyte Chemoattractant Protein-1 mRNA Expression in Human Umbilical Vein Endothelial Cells: Relation to Lysophosphatidylcholine Contents and Inhibition by Nitric Oxide Donor

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Low-density lipoprotein (LDL) may undergo more glycation or oxidation in patients with diabetes mellitus than in nondiabetic subjects. We investigated whether glycoxidized LDL (goLDL) induces monocyte chemoattractant protein-1 (MCP-1) mRNA expression through activation of nuclear factor-kappaB (NFκB), and determined the effect of nitric oxide (NO) on MCP-1 mRNA expression in human umbilical vein endothelial cells (HUVEC). Oxidized (oxLDL) or goLDL enhanced MCP-1 mRNA expression in HUVEC, and preincubation with NOR3, a NO donor, abrogated such stimulation. goLDL increased NFκB-DNA binding activity in HUVEC and this effect was also suppressed by NOR3. We measured lysophosphatidylcholine (lyso-PC) contents in modified LDL using electrospray ionization liquid chromatography/mass spectrometry (LC/MS) to identify its molecular species. MCP-1 mRNA expression and NFκB activation correlated significantly with palmitoyl- and stearoyl-lyso-PC contents in LDL. Our results suggest that LDL modified by glycation and oxidation may contribute to the development of accelerated atherosclerosis in the presence of diabetes, a process that may be prevented by increased vascular NO availability.

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IT IS WELL ESTABLISHED that oxidative modification of low-density lipoprotein (LDL) plays a causal role in human atherogenesis. The development of atherosclerosis is accelerated in patients with diabetes mellitus, and LDL in a diabetic state is susceptible to glycation<sup>1,2</sup> and oxidation.<sup>3,4</sup> Recent immunohistochemical studies of human atherosclerotic lesions demonstrated that advanced glycation end products (AGE) and oxidized LDL (oxLDL) are colocalized in the cytoplasm of macrophage-derived foam cells.<sup>5</sup> Therefore, it is conceivable that glycoxidized LDL (goLDL) may contribute to the development and progression of atherosclerosis in diabetes mellitus.

LDL oxidized by enzymes or copper ion induces the expression of chemokines, including monocyte chemoattractant protein-1 (MCP-1), in various cells.<sup>6-8</sup> MCP-1 has been implicated as a pathogenetic factor in the infiltration of monocytes into the subendothelial space,<sup>9,10</sup> which is a crucial initial step in atherogenesis. Monocyte penetration through the vascular endothelium is thought to occur in response to a gradient of MCP-1 released from activated endothelial cells.<sup>11</sup> Although oxLDL upregulates the gene expression of MCP-1 by activating the transcription factor nuclear factor-kappaB (NFκB) in endothelial cells,<sup>12</sup> whether goLDL also activates MCP-1 expression similar to oxLDL has not yet been determined.

Nitric oxide (NO), a potent vasodilator released by endothelial cells, is thought to play a key role in preventing the development of atherosclerosis by inhibiting smooth muscle cell proliferation,<sup>13</sup> endothelin generation,<sup>14</sup> and leukocyte adhesion to the endothelium.<sup>15</sup> NO suppresses MCP-1 expression induced by tumor necrosis factor-alpha (TNFα) in human endothelial cells<sup>16</sup> and by lipopolysaccharide in rabbit smooth muscle cells<sup>7</sup> by inhibiting NFκB activation. The present study was designed to investigate whether goLDL induces MCP-1 mRNA expression through NFκB activation, and to determine any protective effect for NO on MCP-1 expression in human umbilical vein endothelial cells (HUVEC). In addition, we measured 2 molecular species of lysophosphatidylcholine (lyso-PC) contents (16:0, 18:0), the main constituent of oxidation, in modified LDL and investigated the relationship be-

tween lyso-PC contents and MCP-1 mRNA expression or NFκB activation in HUVEC.

## MATERIALS AND METHODS

### Cell Culture

HUVEC were isolated from umbilical cord veins using 0.25% trypsin (Difco Laboratories, Detroit, MI) according to the method of Jaffe et al<sup>17</sup> with minor modifications. Cells were grown in M199 medium supplemented with 10% fetal calf serum (FCS) (Gibco BRL, Life Technologies, Rockville, MD), 100 μg/mL heparin (Sigma Chemical Co, St Louis, MO), 20 μg/mL endothelial cell growth supplement (Upstate Biotechnology, New York, NY), 0.33 mg/mL piperacillin sodium (Sankyo Co, Tokyo, Japan) in a humidified atmosphere containing 5% CO<sub>2</sub>. The cells were used prior to the 6th passage.

### Isolation and Modification of LDL

LDL was isolated from human plasma using density-gradient ultracentrifugation.<sup>18</sup> In brief, venous blood samples were collected into tubes containing EDTA from consented healthy volunteers after an overnight fast. After centrifugation, the plasma was adjusted to a density of 1.21 g/mL by adding KBr. In centrifuge tubes, a discontinuous density gradient was created by overlaying 2.8 mL plasma solution with 6.6 mL HEPES buffer solution (0.02 mol/L HEPES, 0.4 mol/L NaCl, and 0.4 mol/L NaOH, pH 7.4, *d* = 1.007 g/mL). The tubes were ultracentrifuged at 65,000 rpm for 3 hours at 15°C. The LDL fraction was aspirated and dialyzed by 0.15 mol/L NaCl, pH 7.4. Thereafter, the LDL solution was sterilized through a filter with a pore size of 0.45 μm

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Submitted August 27, 2001; accepted March 4, 2002.

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0026-0495/02/5109-0009\$35.00/0

doi:10.1053/meta.2002.34703

(Millipore, Bedford, MA). EDTA at a final concentration of 0.26 mmol/L was added to an aliquot of LDL solution, which was stored at 4°C under N<sub>2</sub> gas, as a native LDL (nLDL) preparation until final dialysis.

Glycated LDL (gLDL) was obtained by incubating an aliquot of the LDL solution with 200 mmol/L glucose at 37°C for 3 days. oxLDL was prepared by incubation with 5  $\mu$ mol/L CuSO<sub>4</sub> at 37°C for 12 hours after dialysis with 0.15 mol/L NaCl, pH 7.4. goLDL was prepared by incubating gLDL with 5  $\mu$ mol/L CuSO<sub>4</sub> at 37°C for 12 hours. Thereafter, all LDL preparations were dialyzed with 0.15 mol/L NaCl and 0.26 mmol/L EDTA, pH 7.4, and then stored at 4°C under N<sub>2</sub> gas. The protein concentration of each LDL preparation was determined by the Coomassie brilliant blue method using bovine serum albumin as a standard (Nacalai Tesque, Kyoto, Japan).

#### *Electrophoresis of Lipoprotein and Measurement of Lipid Peroxidation*

Electrophoresis of lipoprotein was carried out to ascertain the purity of the LDL fraction and to measure the electrophoretic mobility using a commercial kit (Titan Gel Lipoproteins; Helena Laboratories, Saitama, Japan). In brief, 1  $\mu$ L of nLDL and human serum were loaded on an agarose gel sheet and electrophoresed in barbital buffer at a voltage of 100 V for 25 minutes. The gel sheet was dried and stained with 0.04% Fat Red 7B in methanol.

Lipid peroxidation was estimated by thiobarbituric acid-reactive substances (TBARS) method of Yagi<sup>19</sup> with minor modifications. In brief, 200  $\mu$ g of protein of each modified LDL was mixed with 10% phosphotungstic acid and was centrifuged. The sediment was mixed with 0.67% thiobarbituric acid solution and then heated at 95°C for 60 minutes. After cooling, n-butanol was added and centrifuged. The absorbance of the n-butanol layer was read at 532 nm by spectrophotometer (U-1100, Hitachi, Tokyo, Japan). Serial dilutions of 1,1,3,3-tetramethoxypropane (Sigma), which yields malondialdehyde (MDA), was used to construct the standard curve. TBARS was expressed as nanomoles of MDA per milligram LDL protein.

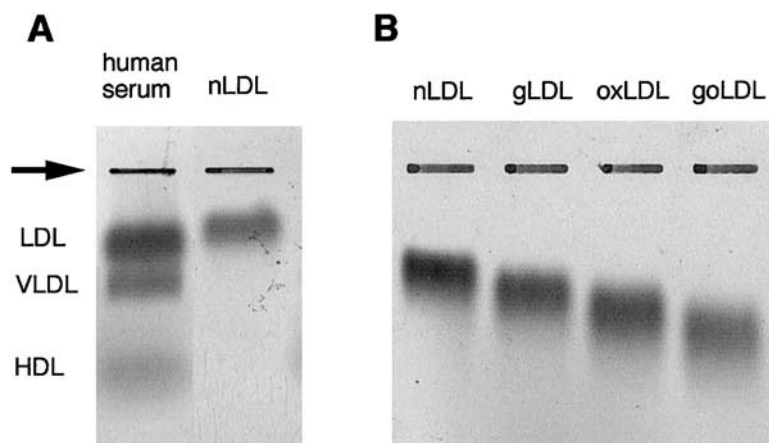
#### *Measurement of lyso-PC in Modified LDL by Electrospray Ionization Liquid Chromatography/Mass Spectrometry*

The content of lyso-PC in modified LDL was measured by liquid chromatography/mass spectrometry (LC/MS).<sup>20</sup> First, the modified LDL was concentrated and dialyzed with 0.05 mol/L HEPES buffer containing 1 mol/L NaCl and 1 mol/L NaOH, pH 7.4, by centrifuging in an Ultrafree-15 centrifugal filter (Millipore). After determination of the protein concentrations of each LDL, lipids were extracted from 50  $\mu$ L of LDL supplemented with 500 ng of 1,2-Dimyristoyl-sn-Glycero-3-

Phosphatidylcholine (DMPC; Avanti Polar-Lipids, Alabaster, AL) as an internal standard, according to the method of Bligh and Dyer.<sup>21</sup> Lipid extracts were evaporated under N<sub>2</sub> gas, and the residue was dissolved in 100  $\mu$ L chloroform. Phospholipids were separated from the extracted lipids by the method of Kaluzny et al<sup>22</sup> using aminopropyl solid-phase extraction chromatography (BAKERBOND spe Columns; J.T. Baker Inc, Phillipsburg, NJ). The separated phospholipid fractions were evaporated under N<sub>2</sub> gas and resuspended in 200  $\mu$ L methanol. Phospholipids were immediately introduced into the electrospray mass spectrometer (LCQ; ThermoQuest, Tokyo, Japan) via high-performance liquid chromatography (HPLC; LC-10, Shimadzu, Kyoto, Japan). Reverse-phase high-performance liquid chromatography (HPLC) was performed by injecting 20  $\mu$ L of isolated phospholipids in methanol into an STR-ODS analytical microcolumn (150 mm  $\times$  2.0 mm, 5  $\mu$ mol/L; Shimadzu) at a flow rate of 0.3 mL/min and eluting with a mobile solvent of methanol/acetonitrile/deionized water (84:14:2, vol/vol/vol). The mass spectrometer was operated in positive mode employing the "full-scan" function set from m/z 100 to 1,000. Quantitative analysis of phospholipids was performed essentially as described by Han et al,<sup>23</sup> using DMPC as an internal standard. Phospholipids were quantitated based on their ion intensity relative to the internal standard. LC/MS chromatogram of LDL showed the peaks of palmitoyl-lyso-PC, stearoyl-lyso-PC, and DMPC (used as an internal standard), at retention times of 2.03, 2.50, and 5.54 minutes, respectively. The electrospray ionization mass spectrum of LDL at retention time 2.03 minutes showed 4 masses for palmitoyl-lyso-PC: m/z 496.5, 497.5, 518.4, and 519.4, which resembled the masses of the standard product of palmitoyl-lyso-PC. The concentration of palmitoyl-lyso-PC was calculated by integrating the relative intensities of the 4 masses for 10 minutes after application in comparison with the standard DMPC. The concentration of stearoyl-lyso-PC in each LDL was calculated similarly by integrating the relative intensities of stearoyl-lyso-PC's 4 masses, m/z 524.6, 525.5, 546.5, and 547.6. The coefficient of variation for electrospray LC/MS assay was 4.8% (n = 12) with 5 ng/mL DMPC as the internal control. Standard curve experiments showed that the electrospray LC/MS method was linear over a wide range (0.1 to 5.0 ng/ $\mu$ L of palmitoyl- or stearoyl-lyso-PC).

#### *Measurement of MCP-1 mRNA Expression*

Cells were incubated with or without 100  $\mu$ g/mL modified LDL in M199 containing 2% FCS for 4 hours. MCP-1 mRNA content was then measured by Northern blot analysis. In brief, total RNA was isolated from HUVEC by using Isogen (Nippon Gene, Tokyo, Japan). Total RNA (15  $\mu$ g) was separated by electrophoresis and transferred to Hybond-N+ nylon membranes (Amersham International, Bucking-



**Fig 1.** (A) Agarose gel electrophoresis of LDL. Human serum or nLDL was applied to agarose gel electrophoresis, and stained with Fat Red 7B. Arrow indicates the site of application. (B) Electrophoretic mobility of nLDL, gLDL, oxLDL, and goLDL.

**Table 1. Parameters of Oxidation in LDL Modified In Vitro**

	nLDL	gLDL	oxLDL	goLDL
Electrophoretic mobility (mm)	7.58 ± 0.45	8.56 ± 0.48	9.56 ± 0.36*	11.08 ± 0.54†§¶
TBARS (nmol MDA/mg protein)	1.26 ± 0.08	1.34 ± 0.10	7.73 ± 0.94†§	7.90 ± 0.82†§
Palmitoyl-lyso-PC (μg/mg protein)	3.32 ± 0.30	7.62 ± 0.34*	14.69 ± 1.77†‡	19.81 ± 2.46†§¶
Stearoyl-lyso-PC (μg/mg protein)	2.61 ± 0.34	5.18 ± 0.48*	9.71 ± 0.84†‡	11.85 ± 1.58†§

NOTE. Values are means ± SEM of 5 measurements for electrophoretic mobility and TBARS or 6 measurements for lyso-PC.

Abbreviations: TBARS, thiobarbituric acid-reactive substances; MDA, malondialdehyde; lyso-PC, lysophosphatidylcholine.

\* $P < .05$ , † $P < .001$  v nLDL; ‡ $P < .01$ , § $P < .001$  v gLDL; ¶ $P < .05$  v oxLDL.

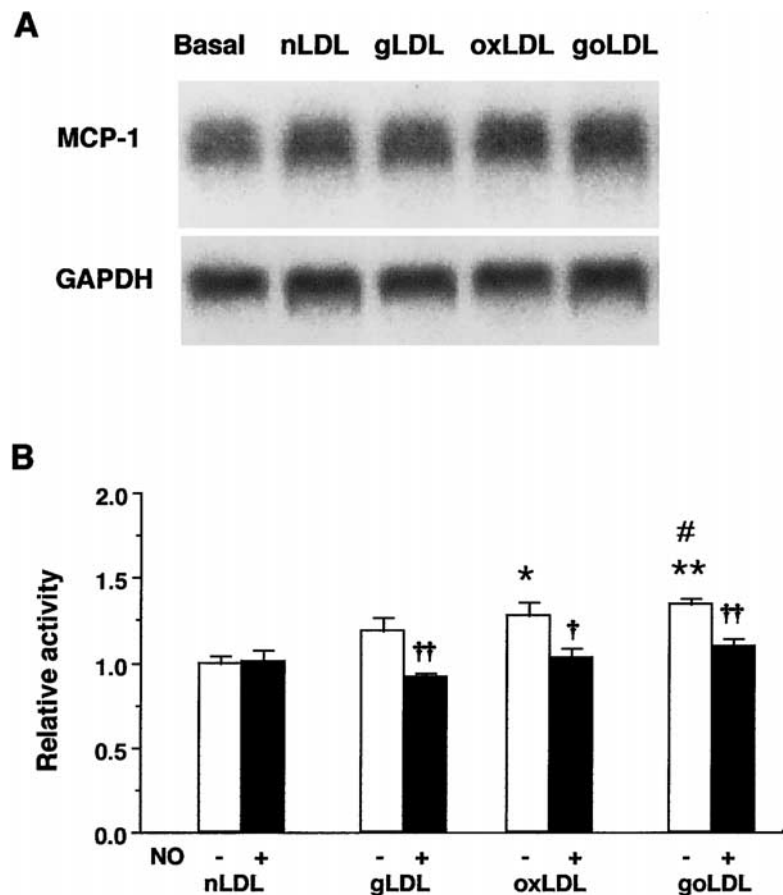
hamshire, England). Hybridization was performed for 1 hour at 65°C in QuikHyb hybridization solution (Stratagene, La Jolla, CA) with a human MCP-1 cDNA (R&D Systems, Minneapolis, MN) or GAPDH cDNA (Oncogene Research Products, Cambridge, MA), that had been labeled with ( $\gamma$ -<sup>32</sup>P) adenosine triphosphate (ATP) (Amersham) by T4 polynucleotide kinase (Promega, Madison, WI). Autoradiography and quantitative analysis were performed with a Bio-Imaging Analyzer (Fuji Photo Film Co, Kanagawa, Japan).

#### Measurement of NFκB Activity

HUVEC were incubated with or without 100 μg/mL of modified LDL in M199 medium containing 2% FCS for 2 hours, and nuclear extracts were then prepared by a modified method of Schreiber et al.<sup>24</sup> In brief, after HUVEC were stimulated with modified LDL, the cells were washed with phosphate-buffered saline (PBS) and suspended in 0.4 mL buffer. After incubation on ice for 15 minutes, 25 μL of 10%

Nonidet P-40 (Sigma) was added. The mixture was centrifuged and the nuclear pellet was suspended in 50 μL ice-cold buffer. The suspension was sonicated and shaken at 4°C for 15 minutes. The supernatant of nuclear extracts, spun at 13,000 × *g* at 4°C for 5 minutes, was stored at -80°C until use after determination of protein concentrations.

Electrophoretic mobility shift assay (EMSA) was performed by incubating nuclear proteins (4 μg) in 10 μL of binding buffer (10 mmol/L Tris-HCl, pH 7.5, 50 mmol/L NaCl, 1 mmol/L MgCl<sub>2</sub>, 0.5 mmol/L EDTA, 0.5 mmol/L dithiothreitol (DTT), 4% glycerol, and 0.05 mg/ml poly(dI-dC) (Pharmacia Biotech, Piscataway, NJ) for 10 minutes at room temperature. Next, 1 μL of <sup>32</sup>P-labeled NFκB oligonucleotide probe (Promega) was added, and the reaction mixture was incubated for 20 minutes at room temperature. For competition assays, 100-fold concentrations of unlabeled oligonucleotides were added to the nuclear proteins. For supershift experiments, antibodies against p50 and p65 (Santa Cruz Biotechnologies, Santa Cruz, CA) were added to



**Fig 2. (A)** Northern blot analysis of MCP-1 mRNA induced by modified LDL in HUVEC. Cells were incubated with or without 100 μg/mL modified LDL for 4 hours. **(B)** Relative intensity of MCP-1 mRNA expression corrected by GAPDH when the expression in the basal state was considered to be 1.0. (□) Expression without NOR3, a NO donor; (■) expression with NOR3 preincubation. Each bar represents the mean ± SEM of 4 experiments. \* $P < .05$ , \*\* $P < .01$  v nLDL; # $P < .05$  v gLDL; † $P < .05$ , †† $P < .01$  v without NOR3.

the nuclear proteins and then incubated for 10 minutes at room temperature. All samples were loaded onto 6% polyacrylamide gel and electrophoresed. Gel contents were dried and autoradiographed by a Bio-Imaging Analyzer.

#### Effect of NO Donor on MCP-1 mRNA and NF $\kappa$ B

HUVEC were incubated with or without 100  $\mu$ mol/L of 4-ethyl-2-hydroxyimino-5-nitro-3-hexenamide (NOR3), NO donor (Dojindo Laboratories, Kumamoto, Japan) in M199 containing 2% FCS, 1 hour before incubation with the modified LDL. The expression of MCP-1 mRNA and NF $\kappa$ B activity were determined as described above.

#### Statistical Analysis

Data were presented as the mean  $\pm$  SEM. Statistical analysis was performed by 1-way analysis of variance (ANOVA) followed by Fisher's test to detect significant differences in multiple comparisons. Relationships between 2 continuous variables were examined by Pearson's correlation analysis. A *P* value of less than .05 was considered statistically significant.

### RESULTS

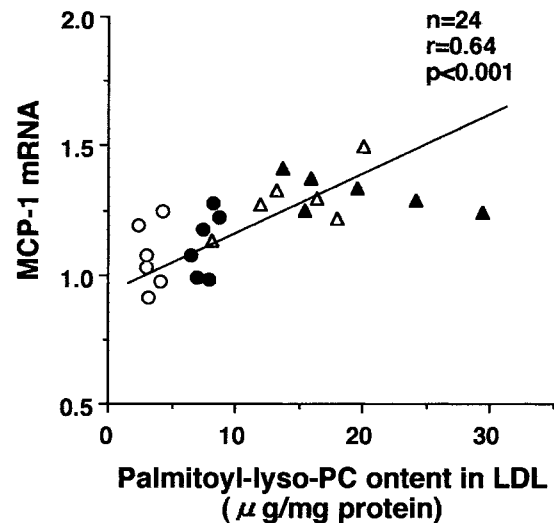
Isolated nLDL showed a single band located in the same position as LDL found in human serum in agarose gel electrophoresis stained by Fat Red 7B (Fig 1A), indicating no protein contaminations in nLDL (stained by Sudan Black B; data not shown). Figure 1B shows the electrophoretic mobility of nLDL, gLDL, oxLDL, and goLDL. The distance from the applied line to the band was significantly greater in oxLDL than in nLDL (Table 1). The mobility distance was significantly greater in goLDL than in nLDL, gLDL, and oxLDL, respectively.

Lipid peroxidation measured by TBARS was significantly higher in oxLDL and goLDL compared with nLDL and gLDL, respectively (Table 1). Palmitoyl- and stearoyl-lyso-PC contents expressed as micograms per milligram LDL protein were significantly higher in gLDL than in nLDL, respectively, and those in oxLDL and goLDL were significantly higher than in nLDL and gLDL, respectively. Palmitoyl-lyso-PC content was significantly higher in goLDL compared to oxLDL (Table 1).

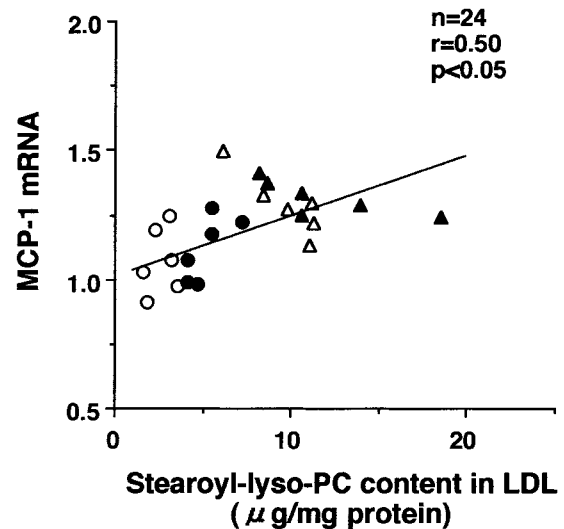
Expression of MCP-1 mRNA levels was evaluated in HUVEC incubated with nLDL or modified LDL for 4 hours using Northern blot analysis (Fig 2A). MCP-1 mRNA expression relative to GAPDH mRNA was significantly higher in oxLDL and goLDL compared with nLDL (Fig 2B). The addition of NOR3, a NO donor, significantly suppressed the expression of MCP-1 mRNA in HUVEC incubated with gLDL, oxLDL and goLDL (Fig 2B). As shown in Fig 3, MCP-1 mRNA expression correlated significantly with palmitoyl-lyso-PC (Fig 3A), as well as with stearoyl-lyso-PC contents in LDL (Fig 3B).

A weak NF $\kappa$ B-specific band was identified in the basal state by unlabeled NF $\kappa$ B oligonucleotide probe and by the supershift assay using specific antibodies against p50 and p65 (Fig 4A). We also determined NF $\kappa$ B-DNA binding activity in HUVEC following exposure to 100  $\mu$ g/mL of native or modified LDL for 2 hours (Fig 4B). The density of the NF $\kappa$ B-specific band relative to the basal state was significantly higher when the cells were incubated with goLDL compared with nLDL or gLDL (Fig 4C). NO donor significantly reduced the density of the NF $\kappa$ B-specific band of goLDL (Fig 4C). As shown in Fig

**A**



**B**



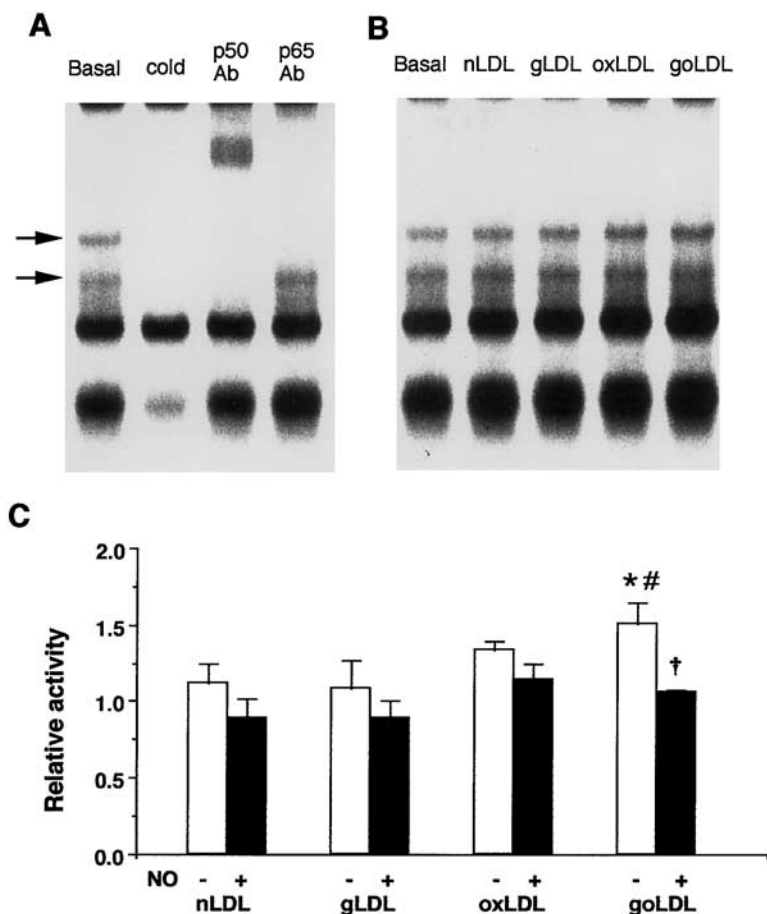
**Fig 3.** Correlation of MCP-1 mRNA expression with the contents of (A) palmitoyl-lyso-PC or (B) stearoyl-lyso-PC. (○) nLDL; (●) gLDL; (△) oxLDL; (▲) goLDL.

5, NF $\kappa$ B activity correlated significantly with palmitoyl-lyso-PC (Fig 5A), as well as with stearoyl-lyso-PC contents in LDL (Fig 5B).

### DISCUSSION

The present study demonstrated that HUVEC incubated with oxLDL or goLDL enhanced MCP-1 mRNA expression, and preincubation with NO donor reversed the stimulation of MCP-1 mRNA expression. NF $\kappa$ B-DNA binding activity was increased in HUVEC incubated with goLDL and was also suppressed by NO donor. We measured lyso-PC by electro-





**Fig 4.** (A) NFκB-DNA binding activity was determined by an electrophoretic mobility shift assay. NFκB-specific bands (indicated by 2 arrows) were identified by competitive (lane 2) and supershift assays (lanes 3 and 4). (B) NFκB-DNA binding activity in HUVEC exposed to nLDL, gLDL, oxLDL, and goLDL. (C) NFκB-DNA binding activity relative to the basal state. (□) Activity without NOR3; (■) activity with NOR3 preincubation. Data are mean  $\pm$  SEM of 4 experiments. \* $P < .05$  v nLDL; # $P < .05$  v gLDL; † $P < .05$  v without NO.

spray to identify its molecular species. MCP-1 mRNA expression and NFκB activation correlated significantly with palmitoyl- and stearoyl-lyso-PC contents in LDL, respectively.

It has been reported that minimally modified LDL by oxidation induces MCP-1 mRNA expression in endothelial cells.<sup>6,25</sup> Minimally modified LDL has been defined as LDL that has mild lipid peroxidation and binds native LDL receptors but not scavenger receptors.<sup>6</sup> Recently, it was reported that highly oxidized LDL (TBARS 182 nmol/mg protein) upregulated lectin-like receptor for oxLDL (LOX-1) and activated MCP-1 mRNA.<sup>26</sup> Since it was reported that more oxidized goLDL than that in our study (TBARS 26.8 nmol/mg protein) was metabolized through native LDL receptors but not by scavenger receptors,<sup>27</sup> our goLDL seems to bind native LDL receptors and activate MCP-1 mRNA expression in HUVEC as minimally modified LDL. Since the effects of modified LDL and NO donor on HUVEC were investigated only at particular time, time studies will be required to clarify these effects in detail.

Bucala et al<sup>28</sup> demonstrated that incubation of LDL with 200 mmol/L glucose for 3 days resulted in the formation of readily measurable levels of AGE on both lipid and apoprotein. Other studies demonstrated that AGE induced the expression of heme oxygenase, tissue factor, and vascular cell adhesion molecule-1 (VCAM-1) genes by activating NFκB in cultured endothelial

cells.<sup>29-31</sup> gLDL mimics oxLDL in terms of accumulation of cholesteryl ester in macrophages<sup>32</sup> and the expression of adhesion molecules in endothelial cells.<sup>33</sup> The present study showed that lyso-PC contents were significantly higher in gLDL prepared by 3-day incubation under 200 mmol/L glucose than in nLDL, indicating that oxidation occurred in the glycation process in LDL. However, gLDL did not stimulate MCP-1 mRNA expression in HUVEC. The glucose autooxidation in our gLDL might be too mild to elicit biologic responses in HUVEC. On the other hand, gLDL seems to be oxidized more readily than nLDL, because palmitoyl-lyso-PC content was significantly higher in goLDL than in oxLDL. This is in line with the report of Sakurai et al<sup>34</sup> who showed that gLDL incubated with  $\text{Fe}^{3+}$  caused a higher level of lipid peroxidation than nLDL. Stimulation of MCP-1 mRNA expression by goLDL was significantly enhanced compared to nLDL and gLDL. These findings suggest that the co-occurrence of glycation and oxidation in LDL strongly promotes its atherogenicity.

Lyso-PC increased the DNA-binding activity of NFκB partly through a protein kinase C (PKC)-mediated pathway<sup>35</sup> and stimulated MCP-1 gene expression in HUVEC.<sup>36</sup> Lyso-PC increased vascular superoxide anion production via PKC activation in intact vessels,<sup>37</sup> and stimulated  $\text{O}_2^-$  production partly through a membrane-associated NADH-dependent  $\text{O}_2^-$  production system.<sup>38</sup> Superoxide is considered as a second messenger

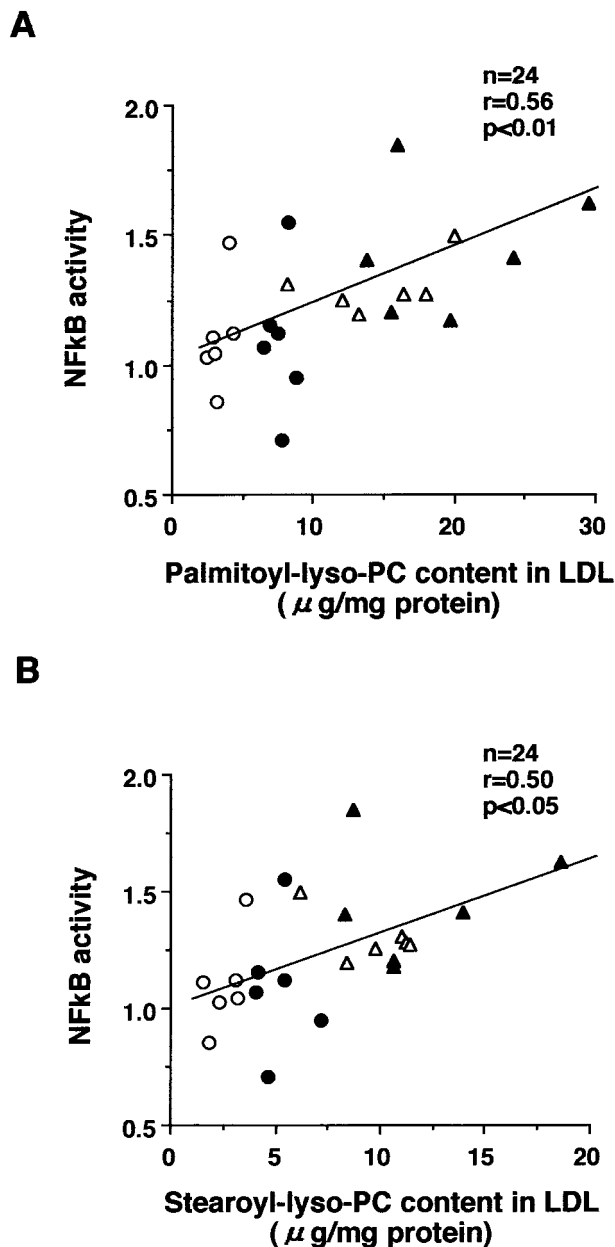


Fig 5. Correlation of NFκB-DNA binding activity with the contents of (A) palmitoyl-lyso-PC and (B) stearoyl-lyso-PC. (○) nLDL; (●) gLDL; (△) oxLDL; (▲) goLDL.

for activating the transcription factor NFκB, followed by the expression of MCP-1 mRNA in human smooth muscle cells<sup>39</sup> and in mouse mesangial cells.<sup>40</sup> In the present study, therefore,

increased lyso-PC content in goLDL may cause MCP-1 gene expression in response to NFκB activation probably due to the generation of superoxide in HUVEC. We have previously measured palmitoyl- and stearoyl-lyso-PC contents in LDL using HPLC, and showed that lyso-PC contents of both species were higher in diabetic patients than in nondiabetic control.<sup>41</sup> In addition, diabetic patients with macroangiopathy had higher levels of palmitoyl-lyso-PC than those without macroangiopathy, whereas stearoyl-lyso-PC levels did not differ between the 2 groups. In the present study, however, palmitoyl- and stearoyl-lyso-PC contents in LDL showed similar correlation with MCP-1 mRNA expression and NFκB activation in HUVEC. Since palmitoyl- and stearoyl-lyso-PC are known to exhibit certain biologic differences,<sup>42,43</sup> the differences between the 2 species of lyso-PC related to the atherogenic activities other than MCP-1 gene expression, remain to be clarified.

NO donors significantly attenuated lyso-PC-induced upregulation of p-selectin by suppressing PKC activation in cat endothelial cells.<sup>44</sup> In addition, MCP-1 expression and NFκB activation were reduced by NO donor in rabbit smooth muscle cells.<sup>7</sup> In the present study, we demonstrated that NO donor significantly suppressed MCP-1 mRNA expression and NFκB activation induced by goLDL in HUVEC. In this regard, Katsuyama et al<sup>45</sup> reported that NO donors inhibit the expression of inducible NO synthase (iNOS) and activation of NFκB induced by interleukin-1β in rat smooth muscle cells, by directly inhibiting the phosphorylation and subsequent degradation of IκBα, the cytoplasmic inhibitor of NFκB. In contrast, Spiecker et al<sup>46</sup> reported that NO donor decreased TNFα-induced VCAM-1 expression in human endothelial cells via inhibition of NFκB activation, by increasing the expression and nuclear translocation of IκBα, but not by preventing the phosphorylation and degradation of IκBα. The apparent discrepancy between these studies may be due to the different cell types, species, cytokines, and NO donors used in the experiments. Since it has been reported that vascular NO availability is reduced in diabetes mellitus,<sup>47-49</sup> goLDL may induce MCP-1 mRNA expression in endothelial cells under diabetic condition, leading to accelerated atherosclerosis.

In conclusion, goLDL stimulated HUVEC and upregulated MCP-1 mRNA expression at least in part due to an enhanced NFκB activity, which was completely inhibited by a NO donor. Palmitoyl- and stearoyl-lyso-PC in LDL correlated with NFκB activity and MCP-1 gene expression, respectively. The present study suggests that LDL modified by glycation and oxidation may contribute to the development of accelerated atherosclerosis under diabetic condition, a process that could be prevented by increased vascular NO availability.

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