The Implication of Cigarette Smoking and Cessation on Macrophage Cholesterol Efflux in Coronary Artery Disease Patients

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Running title: Effect of smoking cessation on macrophage cholesterol efflux

Abbreviations: CAD: Coronary Artery Disease; LOX-1: Lectin-like oxidized low-density lipoprotein receptor-1; SR-B1: scavenger receptor B1; ABCG1: ATP-binding cassette transporters G1; apoA-1: apolipoprotein A-1; CO: carbon monoxide; α-7 nAChR: nicotinic acetylcholine receptor α-7 subunit; PBMCs: peripheral blood monocyte cells; THP-1: Human acute monocytic leukemia cell line; CORM-2: Tricarbonyldichlororuthenium (II) dimmer.
Abstract:

Background: We investigated ATP-binding cassette transporters A1/G1 expression and function in mediating cholesterol efflux by examining the macrophages of cigarette-smoking patients with coronary artery disease (CAD) before and after smoking abstinence.

Methods and Results: Peripheral blood monocyte cells were collected from non-smokers (n=17) non-CAD (NCAD) smokers (n=35), and CAD smokers (n=32) before and after 3 months’ smoking cessation. We found that the ABCA1 expression level was lower in macrophages from both NCAD and CAD smokers than for non-smokers at baseline. The ABCA1 function of mediating cholesterol efflux was reduced in NCAD and CAD smokers as compared with non-smokers. After 3 months’ smoking cessation, ABCA1 expression and function were improved in CAD smokers. However, ABCG1 expression and function did not change after smoking cessation. Furthermore, ABCA1 expressions were inhibited by tar in THP-1-derived macrophages through the inhibition of Liver X receptors. Nicotine and carbon monoxide did not inhibit ABCA1 expression.

Conclusion: Our results indicate that chronic cigarette smoking impaired ABCA1-mediated cholesterol efflux in macrophages, and tobacco abstinence reversed the function and expression of ABCA1, especially in CAD patients. It was tobacco tar, rather than nicotine or carbon monoxide, that played a major role in the tobacco-induced disturbance of cellular cholesterol homeostasis.

Key words: cigarette smoking; coronary artery disease; macrophages; cholesterol efflux; ABCA1

Introduction:

Cigarette smoking is an independent risk factor of atherosclerosis (1, 2). In atherosclerotic lesions, the primary cell type is foam cells, which are macrophages overloaded with cholesterol ester. Macrophage over-expression of scavenger receptors, such as CD36, lectin-like oxidized low-density lipoprotein receptor-1 (LOX-1), and scavenger receptor A (SR-A), can internalize
modified lipoprotein during differentiation under inflammation and oxidative stress, leading to the overload of cholesterol ester (3, 4). Meanwhile, cholesterol efflux of macrophages can be mediated by different pathways including ATP-binding cassette transporters, scavenger receptor B1 (SR-B1), and aqueous diffusion (5). ATP-binding cassette transporters A1 and G1 (ABCA1/G1) play pivotal roles and have been shown additive activities in cholesterol efflux from macrophages in vivo (6). The imbalance between influx and efflux of cholesterol turns macrophages into lipid overloaded foam cells.

Both ABCA1 and ABCG1 are well studied as mediators that regulate cholesterol homeostasis. ABCA1 deficiency was identified as Tangier’s disease, which is characterized by low level of high density lipoprotein cholesterol (HDL-C) level in patient’s serum (7). Many studies indicated that ABCA1 is a key player in modulating cholesterol efflux to apolipoprotein A-1 (apoA-1). The dysfunction of this membrane transporter is associated with cellular cholesterol overload and premature atherosclerotic diseases (8–10). Although ABCG1 is ubiquitously expressed in macrophages, data from human and animal models give contradictory evidence as to whether it is a protector of atherosclerosis (11–13).

Nicotine, tobacco tar, and carbon monoxide (CO) are major alkaloids in cigarette smog, and have been considered, at least partially, responsible for the deleterious effects of tobacco because they are all toxic components of cigarettes. Although nicotine, with the stimulation of the nicotinic acetylcholine receptor α-7 subunit (α-7 nAchR), has now become the focus of attention because of its anti-inflammatory effect (14), other studies have demonstrated that it induces oxidative stress, which may play an important role in the development of atherosclerotic disease (15). CO has been well studied in apoptosis and proliferation (16, 17), and there is little evidence for its association with lipid metabolism.

Although cigarette smoking promotes inflammation, thrombosis, and oxidative stress, the underlying mechanisms involved in the pathophysiology of peripheral blood monocyte cell (PBMCs) dysfunction in response to cigarette smoking remain to be understood. Data on which
toxic components of tobacco contribute to macrophage cholesterol accumulation are not available. In the present study, we focused on the effects of cigarette smoking and smoking abstinence on ABCA1/G1 expression and function in macrophages from chronic smokers and patients with coronary artery disease (CAD). We also observed the effect of different tobacco components on ABCA1/G1 expression, as well as the related signal transduction pathway.

Materials and methods

This was a randomized, prospective, and parallel controlled study. All the subjects, including 17 non-smokers (non-smokers), 35 non-CAD smokers (NCAD smokers), and 32 CAD smokers (CAD smokers), were screened at Peking Union Medical College Hospital (PUMCH) (Beijing, China). Smoking subjects eligible for this study were individuals aged 40–80 years who had smoked at least 10 cigarettes per day for 10 years. Smoking subjects were randomized in a 1:1 ratio to either a smoking cessation subgroup or a continued smoking subgroup. In the NCAD smoker group, 17 of 35 smoking subjects were randomized into a smoking abstinence group (NCAD-abs) and 18 into a continuing smoking group (NCAD-smo) as a paralleled control. Sixteen of 32 CAD smokers were randomized into a smoking abstinence group (CAD-abs) and 16 into a smoking group (CAD-smo). Smokers placed into smoking abstinence groups were asked to stop smoking for the entire study period of 3 months. Cotinine concentration in urine and carbon monoxide (CO) of expiration were tested to insure compliance with the study requirements (18). Seventeen non-smoking subjects were enrolled as a control group (Figure 1). There were 14 non-CAD smokers and 13 CAD smokers who completed the 90 days of smoking cessation. Four CAD smokers and 5 non-CAD smokers in smoking cessation groups withdrew during follow up. All the CAD smokers had documented CAD, including individuals who survived myocardial infarction (MI) or experienced angina pectoris and coronary stenosis ≥ 75% in a major epicardial artery by coronary angiography. The exclusion criteria included acute coronary syndrome, diabetes, taken statin within two weeks preceding blood sampling, serum triglyceride ≥ 400mg/dl, ischemic cerebrovascular or cardiac episodes within 3 months preceding
the randomization, and hypertension. Fifty milliliters of blood was collected. All participants showed normal renal function, liver function, and an acceptable lipid profile.

**Ethics statement**

The study was registered in the Chinese clinical trial registry (Registration number: ChiCTR-RCH-10000748). Ethical approval for the study was obtained from the Human Ethics Committee of PUMCH. Informed consents were obtained from each study subject. The study protocol conformed to the ethical guidelines of the Declaration of Helsinki.

**Reagents**

The Ficoll-Paque premium was purchased from Haoyang (Tianjing, China), the human macrophage colony stimulation factor (M-CSF) was from PeproTech, Inc., and 22-[N-nitrobenz-2-oxa-1, 3-diazol-4-yl-amino]-23, 24-bisnor-5-cholen-3-ol (NBD-cholesterol) was obtained from Molecular Probes (Eugene, OR, USA). Antibodies against ABCA1, ABCG1, and GAPDH were from Novus Biologicals (ABCAM, CO, USA). The human monocytes line THP-1 was obtained from the cell bank of the Basic Research Institute of the Chinese Academy of Medical Sciences (Peking, China). Phorbol myristate acetate (PMA), purified human high density lipoprotein (HDL), and apolipoprotein A-1 (apoA-1) were purchased from Sigma (Sigma-Aldrich, St. Louis, MO, USA). The fluorescence cholesterol was diluted in ethanol, and the final concentration of ethanol was 0.5%. Cigarette tar, a generous gift from the China National Tobacco Corporation, was dissolved in ethanol. Hydrogen tartrate salt of nicotine, which was also obtained from Sigma, was dissolved in normal saline (0.9% NaCl) to obtain the stock concentration, and the pH of the nicotine solution was adjusted to 7.4 by NaOH (1M). Tricarbonyldichlororuthenium (II) dimmer (CORM-2) released CO was used to mimic carbon monoxide function (19). CORM-2, α- bungarotoxin, and TO901317 were purchased from Sigma.

**Biochemical analysis**

Serum biochemical tests, including plasma lipid profile (total cholesterol, triglyceride, LDL-C, HDL-C, apoA-1, and apoB), liver function (ALT, AST, Bil, A/G, AKP, GGT), kidney function (Cr,
BUN), electrolytes (K, Na, Cl) (Olympus AU-5400), complete blood count (Sysmex XE-5000-2),
and urinalysis (SIEMENS Clinitek Atlas), were conducted in the clinical laboratory of PUMCH.

Isolation of human monocytes and cell culture

Isolation of peripheral blood monocytes (PBMCs) was performed by gradient centrifugation layering of 50 ml heparinized (10 U/ml) blood. The viability and purity of monocytes were determined by flow cytometric analysis (Accuri Cytometers, BD, USA) (CD14 staining), confirming that at least 85% purity was achieved. Monocytes were first cultured in a serum-free RPMI 1640 medium containing 25 mmol/L Hepes and 10 ng/mL Human M-CSF for 4–6 hours and then cultured in RPMI1640 containing 10% autologous serum and M-CSF for 7 days at 37°C in humidified incubators with 5% CO₂. CD68 staining was positive in the PBMC-derived macrophages.

Analysis of ABCA1 and ABCG1 protein expression

Western blot analysis was performed to analyze ABCA1 and ABCG1 protein expression. Cells were lysed in a lysis buffer containing Tris (50 mmol/L), NaCl (100 mmol/L), NAF (50 mmol/L), EDTA (1 mmol/L), 0.1% sodium dodecyl sulfate (SDS), 0.5% deoxycholic acid sodium salt, 1% Triton X-100, and protease inhibitor. Cell lysate was kept on ice for 30 min and then centrifuged at 12,000 g for 10 min at 4°C. The protein level was measured using a BCA kit. A total of 80 μg of protein per sample was electrophoresed on 4% to 12% gradient PAGE gel and electroblotted on polyvinylidene difluoride membranes. After blocking the blots for 3 hours at room temperature with TBS-T containing 5% bovine serum albumin and 0.5% Tween 20, membranes were incubated with anti-ABCA1/G1 antibody (1:1000 dilutions) for 24 hours. Immunoreactivity was detected by HRP-conjugated goat anti-mouse secondary antibody. Protein bands were detected with an ECL detection kit (Pierce Rockford, IL, USA). Densitometry analysis of the immunoreactive band was performed using AlphaEaseFC (Alpha Innotech, Santa Clara, CA, USA).

Analysis of ABCA1 and ABCG1 mRNA expression
A quantitative real-time polymerase chain reaction (RT-PCR) was performed to evaluate mRNA expression. RNA was extracted from cells by a TRI reagent (Sigma). Two µg of RNA was used for complementary DNA (cDNA) synthesis with oligo (dT) and SuperScriptIII reverse transcriptase (Invitrogen). For investigation of ABCA1/G1 mRNA, cDNA was synthesized using the Reverse Transcription System for First Strand cDNA Synthesis kit (Promega, Shanghai, China). The thermocycler protocol for the RT phase was 1 cycle at 20°C for 10 min, 1 cycle at 42°C for 60 min, and 1 cycle at 95°C for 5 min. RT-PCR was performed with specific primers for human ABCA1 (forward: 5'-TACAGCCAGAAAGACACCAG-3'; reverse: 5'-CACAGTAGACTTTTGGGAGAG-3'), ABCG1 (forward: 5'- GCAGTTACTCTGCAGAGATG -3'; reverse: 5'- CGGAAATTCCTTTCAGGAGG -3'). Human GAPDH primers (forward: 5'-ATG GAT GAT GAT ATT GCAG C-3'; reverse: 5'- CTCG GAA GCA TTT GCG GTG GAC G-3') were used for the internal standard. PCR was performed in an ABI 7500 thermocycler (Opticon, USA), and the sample was normalized to GAPDH using the \( 2^{-\Delta\Delta Ct} \) (cycle threshold) method.

**Cholesterol efflux assay**

NBD-cholesterol, a fluorescence analog of cholesterol, was used to evaluate intercellular cholesterol efflux (20, 21). ApoA-1 and HDL were used as the acceptors to evaluate ABCA1- and ABCG1-mediated cholesterol efflux, respectively. Macrophages were incubated in the medium with 5µM NBD cholesterol for 4 hours, at which point the supernatant was discarded. The cells were washed 3 times with PBS buffer and then incubated for 4 hours with medium containing apoA-1 (50µg/ml) or HDL (50µg/ml). The medium was collected and centrifuged at 12000g for 10 min to remove cell debris, and cells were lysed in 0.1% Triton X-100 for at least 1 hour after washing with cold PBS. The fluorescence intensity (FI) of the medium and cell lysate were measured in a black polystyrene 96-well assay plate (Costar, Corning Inc., USA) using a multilabel counter (PerkinElmer, Waltham, MA) at 469 nm wavelengths for excitation and 537 nm for emission. The calculations were performed as follows: the percentage of cholesterol efflux
= FI (efflux medium) / FI (efflux medium) + FI (cell lysate), and changes of cholesterol efflux = rate of cholesterol efflux at endpoint — rate of cholesterol efflux at baseline.

THP-1 culture and treatment

THP-1 monocyte was used to evaluate the effects of different concentrations of nicotine, cigarette tar, and CO on ABCA1 expression in vitro. Cells were cultured in RPMI 1640 medium supplemented with 20% heat-inactivated fetal bovine serum in 5% CO2 at 37°C and were differentiated into macrophages by incubation with 100 ng/mL PMA for 72 hours. Cigarette tar stock dissolved in ethanol was diluted by RPMI 1640 to 0.1 g/L. Hydrogen tartrate salt of nicotine was dissolved in basic RPMI 1640 to obtain the required concentrations (ranging from 10^-7M to 10mM); CORM-2 (from 10^-5M to 100mM) was then diluted by RPMI 1640. Macrophages were cultured in the medium containing different concentrations of tar, nicotine, and CO for 72 hours. Liver X receptors (LXR) agonist TO901317 was used to detect whether LXR was involved in ABCA1 expression mediated by cigarette tar. Macrophages were preincubated with 1 µM TO901317 for 2 hours, and then co-incubated with cigarette tar for another 72 hours. Antagonist of α-7 nAchR, α-bungarotoxin (BTX) was used to investigate whether α-7 nAchR was involved in ABCA1 expression mediated by nicotine. Macrophages were preincubated with 10^-5M BTX for 2 hours, and then stimulated by nicotine for another 72 hours. Total protein and mRNA were harvested for ABCA1 detection, and a 3-(4, 5)-dimethylthiazoliumromide (MTT) assay was carried out to evaluate the survival rate of cells.

Statistical analysis

Data is represented as mean ± SEM. Data were analyzed using the SPSS Statistical Analysis System V-16.0 (SPSS, Chicago, Illinois). One-way ANOVA and paired T test were used. Wilcoxon signed-rank tests were used for ordinal variables and p<0.05 was considered to be statistically significant. Data was presented by GraphPad Prism 5 (GraphPad Software, San Diego,
RESULTS

1. Study Population

Seventy-five subjects (17 non-smokers, 30 NCAD smokers, and 28 CAD smokers) completed the study. Subjects who had been randomized into the smoking cessation group and who failed to stop smoking for 90 days, which was assessed by the cotinine in urine and CO in expiration (supplementary table 1), were excluded from the study.

Baseline plasma apoA-1 and HDL-cholesterol (HDL-C) levels were significantly lower in both NCAD smokers ($p=0.002$ and $p<0.001$, respectively) and CAD smokers ($p=0.019$ and $p=0.004$, respectively) than in non-smokers. With respect to apoA-1 or HDL-C, there was no statistical disparity between CAD and NCAD smokers. Triglyceride and apoB levels were higher in NCAD smokers than in non-smokers ($p=0.003$ and $p=0.038$, respectively). Other parameters, including age, BMI, blood pressure, plasma TC, LDL-cholesterol, and glucose levels, made no difference among the study groups at baseline (table 1).

After 3 months’ smoking cessation, plasma apoA-1 levels increased more significantly in CAD-abs (0.32±0.30 g/L) than in CAD-smo (0.02±0.29 g/L, $p=0.032$), but did not change in NCAD-abs. The change of plasma HDL-C levels in CAD-abs (0.04±0.18 mmol/L) was comparable to the change in CAD-smo (0.13±0.13 mmol/L) ($p=0.203$). There were no significant differences in other biochemical parameters between the study groups (table 2).

2. ABCA1 expression and its function among study groups at baseline

At baseline, ABCA1 protein expression was significantly lower in macrophages from both CAD smokers and NCAD smokers, as compared with the expression from non-smokers ($p=0.001$ and $p=0.033$, respectively). Compared with NCAD smokers, ABCA1 expression was further inhibited in macrophages from CAD smokers ($p=0.037$) (fig. 2A and 2B). ABCA1 mRNA expression in macrophages from CAD smokers was significantly reduced compared with non-smokers ($p<0.001$), while paradoxically, the expression was augmented in macrophages from NCAD
ABCA1-mediated cholesterol efflux to apoA-1 was attenuated in macrophages from CAD smokers (8.15±1.50%) compared with non-smokers (15.30±5.70%, p<0.001). It was also diminished in NCAD smokers (9.97±2.00%) compared with non-smokers (15.30±5.70, p<0.001). ABCA1-mediated cholesterol efflux in CAD smokers was also lower than in NCAD smokers (p=0.017) (fig. 1D).

3. Changes of ABCA1 expression and function in macrophages after smoking cessation from CAD smokers

Both protein and mRNA expression of ABCA1 in macrophages from CAD-abs subjects were increased after 3 months’ smoking cessation (figs. 3A, 3C and 3D) (all with significance levels of p<0.05 compared with baseline). There was no significant change in ABCA1 expression in macrophages from CAD-smo subjects (figs. 3B, 3C and 3D). Compared with baseline, the end of 3 months, ABCA1-mediated cholesterol efflux to apoA-1 was significantly increased in CAD-abs (8.14%±1.61 vs. 11.47%±3.61, p=0.004) but not in CAD-smo (8.16%±1.55 vs. 8.62%±1.49, p=0.473) (fig. 3E). The changes of ABCA1-mediated cholesterol efflux between CAD-abs and CAD-smo after a 3-month follow up were still significantly different (3.33%±3.72 vs. 0.64%±1.97, p=0.036) (fig. 3F).

ABCG1 mRNA expression was up-regulated dramatically (p<0.001) in macrophages from CAD-abs after 3 months’ tobacco abstinence, while ABCG1 protein expression was not increased and the function was not improved (supplementary fig. 2).

4. Changes of ABCA1 expression and function in macrophages from NCAD smokers after smoking cessation

There was no change in ABCA1 protein and mRNA expression in both NCAD-abs (figs. 4A and 4C) and NCAD-smo (figs. 4B, 4C and 4D). Compared with baseline, ABCA1-mediated cholesterol efflux to apoA-1 was improved in macrophages from NCAD-abs (9.33%±2.01 vs.
11.10%±1.96, p=0.002), but not from NCAD-smo (10.61%±1.84 vs. 11.53%±2.78, p=0.137) after follow up at 3 months (fig. 4E). However, the absolute changes of cholesterol efflux did not reach statistical significance between the two study groups (1.77%±1.57 vs. 0.92%±1.98, p=0.255) (fig. 4F). There appeared to be no differences in ABCG1 expression and function from NCAD smokers after tobacco abstinence (supplementary fig. 3).

5. Inhibition of ABCA1 expression from cigarette tar in THP-1-derived macrophages

ABCA1 protein and mRNA expression was enhanced by incubation with 100ng/ml nicotine (figs. 5A and 5B), but inhibited by 0.1 g/L tar (figs. 5C and 5D). Liver X receptor (LXR) agonist TO901317 abolished the inhibitive effect of tar on protein and mRNA expression of ABCA1 (figs. 4C and 4D). Bungarotoxin (BTX), a selective α-7-n acetylcholine receptor (α-7-nAchR) antagonist, inhibited the up-regulation of ABCA1 induced by nicotine (figs. 5A and 5B). MTT assay was carried out to evaluate the toxic reaction and survival rate of cultured cells stimulated by these agents (supplementary fig. 4). The survival rate of cells was not affected by stimulation by tar and nicotine. ABCA1 expression was not affected by the administration of CORM-2 (supplementary fig. 5).

Discussion

The evidence linking cigarette smoking with cardiovascular disease are shown in many studies; however, the mechanisms involved in cigarette exposure associated cardiovascular dysfunction have been largely debated. Previous studies have proven that tobacco aggravates inflammation, thrombosis, oxidation of low-density lipoprotein, and oxidative stress reaction (22–24). Our study was focused on whether tobacco impacts cellular cholesterol metabolism from the lipid metabolic point of view. We found that ABCA1 protein expression and the function of mediated intercellular cholesterol efflux to be deficient in CAD smokers. This deficiency might play a detrimental role in cellular cholesterol metabolism and lead to intercellular cholesterol disposal, which may accelerate the formation of foam cells. Many studies have proven that defects in ABCA1 impaired apoA-1-mediated intercellular lipid efflux develop in early atherosclerosis...
Apart from its role in lipid metabolism, ABCA1 has also been implicated in engulfing apoptotic cells, inhibiting the release of inflammatory mediators, and binding to other lipoproteins (including apolipoprotein E), thus playing various roles in its anti-atherogenic effect (28, 29). Deficiency in ABCA1 protein expression in smokers may attenuate the role of anti-inflammation and anti-oxidation, resulting in atherosclerotic aggravation.

As to the results of this study, it was interesting that ABCA1 mRNA was up-regulated in NCAD smokers, which is inconsistent with the results from ABCA1 proteins. Sticozzi et al. also found that ABCA1 mRNA expression was up-regulated in keratinocyte exposure to air filled with cigarette smoking (30). However, as opposed to our study, Sticozzi’s conclusion was based on cultured cell lines’ *in vitro* exposure for only 50 min; there was 92% gas, which was not hydrosoluble, and 8% particulates in the air filled with cigarette smoke, including nicotine and tar (31). Thus, Sticozzi’s study was a more passive smoking model, while our data from subjects who had been active long-time smokers was a direct assessment of the exposure–response relationship between cigarette smoking and macrophage ABCA1 expression and function.

The plasma HDL-C level was inversely correlated with the risk of clinical events resulting from atherosclerosis. In our study, the HDL-C level was significantly low in all smokers, which is compatible with other observations that smokers have a low concentration of plasma HDL-C (-5.7%) and apoA-1 (-4.2%) (32). The mechanisms involved in diminishing HDL-C have not yet been clarified. We found that plasma apoA-1 was also reduced in all smokers. ABCA1-dependent lipid transfer to apoA-1 is the rate-limiting step in the biogenesis of nascent HDL, and it also plays a key role in the cardioprotective function of HDL (33). It has been proven that removing ABCA1 in macrophages does not affect the circulation of HDL cholesterol (34). ABCA1 in the liver and intestines is essential for maintaining plasma HDL cholesterol levels (35). Thus, the deficiency of ABCA1 protein expression and dysfunction in macrophages probably reflect what is occurring in liver ABCA1 expression, which results in low plasma HDL cholesterol in smokers. Moreover, after 3 months’ smoking cessation, ABCA1-mediated cholesterol efflux to apoA-1 was
dramatically elevated and was also accompanied by up-regulation of ABCA1 protein expression in macrophages from CAD patients. Although increased plasma HDL-C levels did not reach statistical significance, apoA-1 improvement was significant in the CAD smoking cessation subgroup. This fact re-emphasizes the importance of the therapeutic strategy for CAD smokers in quitting tobacco in addition to standard pharmacotherapy treatments.

Why is ABCA1 function in subjects with CAD reversed, but not in NCAD subjects? We found that peroxisome proliferator-activated receptor-gamma (PPAR-γ) mRNA was up-regulated, while ABCA1 was decreased in CAD smokers compared with NCAD smoker at baseline (supplementary fig. 6A, all \( p<0.05 \)). PPAR-γ protein expression was obviously more enhanced in macrophages from CAD than from NCAD smokers (supplementary figs. 6B and 6C, \( p<0.01 \)). These results imply that the mechanism of ABCA1 reversion through smoking cessation may be different for CAD and NCAD smokers. Furthermore, we extended the smoking cessation time to 5 months (post-endpoint) and found that ABCA1 mRNA (supplementary fig. 7A) and protein expression (supplementary figs. 7B and 7C) were enhanced compared with baseline in macrophages from NCAD smokers (both \( p<0.001 \)). Changes in the ABCA1-mediated cholesterol efflux rate was increased significantly (\( p=0.005 \)), but there was no change in PPAR-γ expression (supplementary fig. 7D). We are not sure whether the activation of PPAR-γ is the reason for CAD or a compensatory reaction of CAD. However, Mehrabi et al. found that the expression of PPAR-γ differed according to organ (36). Thus, it is reasonable to suggest that PPAR-γ in macrophages is increased in certain pathological conditions, such as CAD. Although the role of PPAR-γ in the development of heart disease is controversial, the activation of PPAR-γ/LXR-α-induced ABCA1 transcription supports the hypothesis that is has an effect on its development (37, 38). It may be that activation of PPAR-γ leads to a fast reversal of ABCA1 expression and function.

ABCG1-mediated intercellular cholesterol efflux to HDL was shown to be coordinated with ABCA1 in removing excess cellular cholesterol, and it also affected inflammatory cellular
cytokine secretion by modulating cholesterol content in the plasma membrane and within intracellular compartments (39). We did not find any significant difference in ABCG1 expression and function among these groups before or after smoking cessation, which suggests that ABCG1 might not be critical in tobacco-induced changes of macrophage cholesterol metabolism. Because of the complexity of cigarettes, it is difficult to identify the components responsible for pathophysiologic actions and discern the relevant mechanisms. To understand the effects of tobacco ingredients on macrophage ABCA1 expression, we designed in vitro experiments with THP-1 cells. We found that cigarette tar inhibited ABCA1 expression in THP-1 macrophages, while the LXR agonist could abolish this inhibitive effect. This may explain why cigarette smoking reduces ABCA1 expression and function.

Studies in the last century have demonstrated that cigarette tar leads to DNA damage (40, 41). Recent research demonstrates that accumulation of tar-like substances in human neutrophils leads to atypical cell death, which shares features of apoptosis, autophagy, and necrosis (42). We found for the first time that tobacco tar inhibits ABCA1 expression, which may lead to impairment of cholesterol-mediated efflux in macrophages. Nicotine in cigarette smoke is probably the most frequently studied component, and the plasma levels of nicotine range from 10ng/ml to 40ng/ml after exposure to cigarettes (43). In our study, we found that nicotine, which was greater than or equal to the plasma nicotine concentration, increased the ABCA1 mRNA and protein expression in THP-1-derived macrophages. Nicotine may responsible for the results from Sticozzi, who stimulated keratinocytes with cigarette smoke and found up-regulation of ABCA1 expression (30). Although high doses of nicotine favor atherogenic changes in various models, the majority of evidence suggests that nicotine, at concentrations similar to a smoker’s blood level, has a minor effect on the initiation or propagation of atherosclerosis (44, 45). Some studies even show that nicotine plays an anti-inflammatory role that is mediated by the cholinergic pathway (46). Carbon monoxide (CO) is one of the more toxic agents in incompletely burned tobacco inhalation. There is sufficient evidence to suggested that CO affects cardiovascular, respiratory, and central
nervous systems (47, 48). In this study, however, CO did not interrupt the ABCA1 expression in THP-1 macrophages. There are thousands of components in cigarettes, and an abundance of work remains to be performed in exploring which components are responsible for the impaired cholesterol metabolism in macrophages. It was more effective to perform study in vivo than in vitro.

Our study investigated the relationships between cigarette smoking, ABCA1/G1 expression, and the function of cholesterol efflux in macrophages from humans. Cigarette smoking is associated with impaired intercellular cholesterol efflux resulting from the inhibition of ABCA1 expression and function, which is also associated with the reduction of plasma HDL-C levels. Tobacco tar, as a complex in tobacco, plays an important role in inhibiting ABCA1 expression. Our findings provide novel evidence that smoking cessation is an effective method for restoring ABCA1-mediated cholesterol efflux and raising HDL-C levels in CAD patients.

Acknowledgement

Part of this work was performed at the Institute of Basic Medical Sciences (IBMS) of the Chinese Academy of Medical Science. We would like to thank Dr. Wei Chen for helping with testing viability and purity of monocytes, and we would like to thank Dr. Shenghan Chen at the Cleveland Clinic Lerner Research Institute for polishing this manuscript. This work was supported by a grant from the Pfizer Inc. Corporation (WS554487).

Conflict of interest

The authors have declared no conflict of interest.

Reference List


Figure and legend

Figure 1. Study Flow Chart.

Figure 2. ABCA1 expression and its function among study groups at baseline. ABCA1 protein expression was suppressed in macrophages from NCAD and CAD smokers compared with that from non-smokers (each group n=6) (2A). Densitometry was shown as in 2B (p=0.033, p<0.001 vs. non-smoker). 2C showed ABCA1 mRNA was dramatically suppressed in CAD smokers, but obviously up-regulated in macrophages from NCAD smokers compared with that from non-smokers (p<0.001, each group n=10). 2D showed ApoA-1-mediated cholesterol efflux was decreased in macrophages from CAD smokers (p<0.001, n=26) and NCAD smokers (p<0.001, n=24) compared with that from non-smokers (n=10) at baseline. ApoA-1-mediated cholesterol efflux was significant lower from CAD smokers than that from NCAD smokers (p=0.017).

Figure 3. Changes of ABCA1 expression and function after smoking cessation in macrophages from CAD smokers. ABCA1 protein expression did not change in CAD-smo (p=0.156, n=6) (3A and 3C), but it was increased after smoking cessation for 3 months in CAD-abs (p=0.007, n=6) (3B and 3C). 3D showed ABCA1 mRNA expression was up-regulated in macrophages from CAD-abs (p=0.042, n=10). 3E showed that after 3 months' smoking cessation, ABCA1-mediated cholesterol efflux was increased significantly compared with baseline in CAD-abs (p=0.004, n=12) while no change in CAD-smo (p=0.473, n=10). 3F showed that after 3 months, ABCA1-mediated cholesterol efflux increased significantly in CAD-abs compared with that in CAD-smo (p=0.036).

Figure 4. Changes of ABCA1 expression and function in macrophages from NCAD smokers after smoking cessation. ABCA1 protein expression did not change in macrophages from both NCAD-smo and NCAD-abs (p=0.458, n=6) (4A, 4B and 3C). 4D showed ABCA1 mRNA expression did not change in both NCAD-smo and NCAD-abs (p=0.267 and 0.089, n=10). 4E
showed that after 3 months’ smoking cessation, ABCA1-mediated cholesterol efflux was increased significantly in NCAD-abs ($p=0.002$, $n=12$) while no change in NCAD-smo ($p=0.137$, $n=12$). 4F showed that the changes in ABCA1-mediated cholesterol efflux in NCAD-abs were not significantly different from that in NCAD-smo ($p=0.255$).

**Figure 5. Inhibition of ABCA1 expression from cigarette tar in THP-1-derived macrophages**

THP-1 cells were differentiated into macrophages. Macrophages were incubated in medium contained tar or nicotine for 72 hours respectively. 5A ABCA1 protein expression was stimulated by 100ng/ml nicotine, which could be reversed by BTX. 5C ABCA1 protein expression was inhibited by 0.1 g/L tar, which could be abolished by LXR agonist TO901317. 5B and 5D showed that ABCA1 mRNA was up-regulated by nicotine ($p=0.006$ vs. control) and down regulated by tar ($p=0.039$ vs. control). The effects of nicotine and tar on ABCA1 mRAN expression could be removed by BTX ($p<0.001$ vs. nico) and TO901317 ($p<0.001$ vs. tar), respectively.
Table 1. Laboratory profile of study population at baseline

<table>
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<th>Characteristic</th>
<th>Non-smoker (n=17)</th>
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<td>57.68±8.12</td>
<td>0.382</td>
</tr>
<tr>
<td>BMI</td>
<td>24.70±2.93</td>
<td>25.81±2.52</td>
<td>25.38±3.26</td>
<td>0.213</td>
</tr>
<tr>
<td>ApoA-1 (g/L)</td>
<td>1.58±0.32</td>
<td>1.33±0.20</td>
<td>1.38±0.28</td>
<td>*0.002</td>
</tr>
<tr>
<td>ApoB (g/L)</td>
<td>0.88±0.14</td>
<td>0.99±0.19</td>
<td>0.81±0.16</td>
<td>*0.038</td>
</tr>
<tr>
<td>TC (mmol/L)</td>
<td>4.99±0.81</td>
<td>5.09±0.81</td>
<td>4.52±0.77</td>
<td>0.645</td>
</tr>
<tr>
<td>TG (mmol/L)</td>
<td>1.04±0.44</td>
<td>2.20±1.44</td>
<td>1.74±1.35</td>
<td>*0.003</td>
</tr>
<tr>
<td>HDL-C (mmol/L)</td>
<td>1.41±0.30</td>
<td>1.12±0.22</td>
<td>1.17±0.29</td>
<td>*&lt;0.001</td>
</tr>
<tr>
<td>LDL-C (mmol/L)</td>
<td>3.06±0.53</td>
<td>3.15±0.85</td>
<td>2.66±0.76</td>
<td>0.696</td>
</tr>
<tr>
<td>Glucose (mmol/L)</td>
<td>4.90±0.61</td>
<td>6.01±3.81</td>
<td>5.86±0.13</td>
<td>0.144</td>
</tr>
<tr>
<td>SBP (mmHg)</td>
<td>118±12</td>
<td>122±8</td>
<td>124±14</td>
<td>0.217</td>
</tr>
<tr>
<td>DBP (mmHg)</td>
<td>74±10</td>
<td>72±8</td>
<td>72±12</td>
<td>0.508</td>
</tr>
</tbody>
</table>

All data normally distributed variables are expressed as mean and standard deviation. 

Abbreviations: BMI, body mass index (kg/m2); TC, total cholesterol; TG, triglycerides; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol; SBP, systolic blood pressure; DBP, diastolic blood pressure

*p<0.05 obtained in the comparison between CAD or healthy smokers and non-smokers using ANOVA.
Table 2. Laboratory profile of study population after 3 month follow-up.

<table>
<thead>
<tr>
<th>Characteristic</th>
<th>NCAD smoker</th>
<th>CAD smoker</th>
<th>p value</th>
<th>p value</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>NCAD-smo (n=16)</td>
<td>NCAD-abs (n=14)</td>
<td></td>
<td>CAD-smo (n=15)</td>
</tr>
<tr>
<td>ApoA-1 (g/L)</td>
<td>0.09±0.10</td>
<td>0.14±0.25</td>
<td>0.486</td>
<td>0.02±0.29</td>
</tr>
<tr>
<td>ApoB (g/L)</td>
<td>-0.01±0.09</td>
<td>-0.03±0.33</td>
<td>0.891</td>
<td>-0.02±0.26</td>
</tr>
<tr>
<td>TC (mmol/L)</td>
<td>-0.10±0.36</td>
<td>-0.14±0.79</td>
<td>0.867</td>
<td>-0.05±0.17</td>
</tr>
<tr>
<td>TG (mmol/L)</td>
<td>-0.07±0.47</td>
<td>-0.05±0.36</td>
<td>0.925</td>
<td>-0.12±0.35</td>
</tr>
<tr>
<td>HDL-C (mmol/L)</td>
<td>0.10±0.09</td>
<td>0.20±0.26</td>
<td>0.184</td>
<td>0.04±0.18</td>
</tr>
<tr>
<td>LDL-C (mmol/L)</td>
<td>-0.10±0.46</td>
<td>-0.14±0.44</td>
<td>0.839</td>
<td>-0.04±0.64</td>
</tr>
</tbody>
</table>

*p<0.05 obtained in the comparison between cessation and continue using independent-samples T test.
Figure 1.

Study flow diagram

Subjects (N=84)

Non-smoker (n=17)

Non-CAD Smoker (N=35)

Smoking abstinence (N=17)
Smoking continue (N=18)

CAD smoker (N=32)

Smoking abstinence (N=16)
Smoking continue (N=16)

3 months follow up

NCAD-abs (n=14)
NCAD-smo (n=16)
CAD-abs (n=13)
CAD-smo (n=15)

N: number of subjects randomized into subgroup
n: number of subjects finished the study
Figure 2A  

**Figure 2B  ABM41 protein expression at baseline**

<table>
<thead>
<tr>
<th></th>
<th>non-smoker</th>
<th>NCAD smoker</th>
<th>CAD smoker</th>
</tr>
</thead>
<tbody>
<tr>
<td>ABCA1</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>GAPDH</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Fold ABCA1 protein (GAPDH)

- **p<0.001**
- **p<0.033**
- **p<0.037**

**Figure 2C  ABCA1 mRNA expression at baseline**

<table>
<thead>
<tr>
<th></th>
<th>non-smoker</th>
<th>NCAD smoker</th>
<th>CAD smoker</th>
</tr>
</thead>
<tbody>
<tr>
<td>ABCA1</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>GAPDH</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Relative expression of ABCA1 mRNA (GAPDH)

- **p<0.003**
- **p<0.049**
- **p<0.011**

**Figure 2D  Non-smoker, NCAD smoker and CAD smoker cholesterol efflux rate at baseline**

<table>
<thead>
<tr>
<th></th>
<th>non-smoker</th>
<th>NCAD smoker</th>
<th>CAD smoker</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cholesterol efflux to apop-1</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

- **p<0.001**
- **p<0.017**
- **p<0.001**