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**Serum from patients with** 

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**atherosclerosis resulted in** 

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**macrophage cholesterol accumulation**

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# **Abstract**

Aims: The aim of our study was to analyze the effect of serum derived from patients with chest pain, with or without significant atherosclerosis, on J774A.1 macrophages cholesterol metabolism.

**Methods:** Thirty-nine patients with chest pain underwent CT Coronary Angiography (CTCA) to assess atherosclerosis. They were divided into three groups (n=13 for each group), according to extent of Coronary Artery Disease (CAD): No disease (NCAD), Non-significant (NSCAD), or significant CAD (SCAD).

**Results:** Serum total cholesterol level was similar in all three groups, whereas the SCAD group had the lowest serum HDL level and highest serum triglyceride levels, compared with the other groups.

The patient's serum (30μl) was incubated with J774 macrophages for 18h. Cellular cholesterol mass was found to be significantly higher by 44 percent in SCAD patients compared to NCAD patients, and by 23 percent compared to NSCAD patients.

In parallel, we observed a significant enhanced cholesterol biosynthesis rate by 53 percent in macrophages treated with serum from the SCAD patients, as compared to NCAD patients, or by 17 percent when compared to the NSCAD patients.

In accordance with the above results, HMGCR (the rate limiting enzyme in cholesterol biosynthesis) expression was significantly upregulated in macrophages treated with serum from SCAD patients, in comparison to NCAD or NSCAD patients.

**Conclusion:** These results clearly demonstrate high macrophage atherogenicity for serum harvested from patients with significant atherosclerosis.

# **Abbreviations**

CTCA: CT coronary angiography; SCAD: Significant Coronary Artery Disease; NSCAD: Non-Significant Coronary Artery Disease; HMGCR: Hydroxy Methyl Glutaryl Coenzyme Reductase; CRP: C Reactive Protein; LDL: Low Density Lipoprotein; HDL: High Density Lipoprotein; IMT: Intima Media Thickness; PBS: Phosphate Buffer Saline; DMEM:

Dulbecco's Modified Eagle's Medium; PCEC: Percentage of Cholesterol Efflux Capacity; BMI: Body Mass Index

## **Introduction**

Atherosclerosis is a progressive disease and the leading cause of morbidity and mortality worldwide [1]. A crucial early step in atherosclerosis development is infiltration of blood monocytes

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from the circulation into the arterial wall [2], where they differentiate into macrophages. These intimal macrophages can take up oxidatively modified lipoproteins at enhanced rate, leading to lipid accumulation and foam cells formation, the hallmark of early atherogenesis [2,3]. Cholesterol accumulation in macrophages can also result from increased cholesterol biosynthesis, and/or decreased cholesterol efflux from the cells [3]. Experimental data support a role of monocytes in acute coronary syndromes and outcome post-infarction[4]. Indeed, macrophages, C-Reactive Protein (CRP), and complement and oxidized Low-Density Lipoprotein (LDL) were colocalized in plaque tissue from patients with unstable coronary artery disease[5]. This may increase vulnerability to plaque rupture and thrombosis.

Plasma levels of oxidized LDL are higher in patients with unstable angina [6], suggesting that antioxidants may have beneficial effects on macrophages [7] and atherosclerotic lesions. Indeed, we have shown that pomegranate juice consumption by patients with carotid artery stenosis reduced their Intima-Media Thickness (IMT), blood pressure, and LDL oxidation [8].

Observational studies have shown that LDL and High-Density Lipoprotein (HDL) cholesterol have opposing associations with risk of myocardial infarction, with LDL cholesterol being positively associated, and HDL cholesterol inversely associated [9-11].

Benjamin F Voight, et al. [12], showed that some genetic mechanisms that raise plasma HDL cholesterol do not seem to lower risk of myocardial infarction. It was previously shown that blood serum of patients with angiographically assessed coronary atherosclerosis can stimulate the accumulation of intracellular lipids (mostly free and esterified cholesterol) in primary culture of intimal smooth muscle cells of human aorta [13,14]. Thus, blood atherogenicity could be a target for antiatherosclerotic therapy [15].

The current guidelines recommend CT Coronary Angiography (CTCA) for assessment of patients with stable angina pectoris and without changes in ECG or troponin[16]. The purpose of the present study was to examine, for the first time, the atherogenicity of serum from SCAD patients, as compared to NSCAD or to NCAD patients, by analyzing their serum effects on macrophage cholesterol metabolism.

## **Methods**

#### **Patients**

Thirty-nine patients with chest pain were admitted to the Internal Medicine Department at Rambam Health Care Campus (Haifa, Israel) for CTCA, in order to assess atherosclerosis in their coronary arteries. They were divided into three groups of 13 patients according to their results: No disease (NCAD), Non-significant (NSCAD) and significant CAD (SCAD). All patients gave written informed consent in accordance with the declaration of HELSINKI. The protocol was approved by the Rambam Health Care Helsinki Committee (protocol 0373-18- RMB).

#### **J774A.1 macrophages**

J774A.1 murine macrophage-like cell line was obtained from the American Type Culture Collection (ATCC, Rockville, MD, USA) and maintained in a humidified incubator (37 $\textdegree$ C, 5% CO2) in DMEM containing 1,000 U/L penicillin, 100 mg/L streptomycin, and 5% heat-inactivated FCS.

#### **Macrophage cholesterol mass**

The cells (1×10<sup>6</sup> cells) were incubated for 18h with 30µl of the patient's serum. After cell wash PBS (X2), cellular lipids were extracted with hexane:isopropanol (3:2, v:v). The hexane phase was evaporated under nitrogen. The amount of cellular cholesterol was determined spectrophotometrically using commercially available kit. The remaining cells in the plates were dissolved in 0.1 M NaOH, and an aliquot was taken measuring cellular protein by the Lowry assay [17,18]. This method uses the Folin-Ciocalteu Phenol reagent and albumin for the standart curve. Results were expressed as µgcholesterol/ mg cell protein.

#### **Macrophage cholesterol biosynthesis rate**

The cells  $(1\times10^6 \text{ cells})$  were incubated for 18h with 30µl of the patients' serum, then washed with PBS and further incubated for 3h at 37°C in DMEM supplemented with 3μCi/ mL [3 H]-labeled acetate (PerkinElmer). After incubation, the cells were washed twice in PBS. Cellular lipids were then extracted with hexane/isopropanol (3:2 v/v), the hexane phase evaporated under nitrogen, then lipids separated by thinlayer chromatography on silica gel plates using hexane/ether/ acetic acid (130:30:1.5 v/v). Cholesterol spots were visualized by iodine vapor using an appropriate standard for identification and [ ${}^{3}$ H] was counted by  $\beta$ -counter (Packard Tri Carb 2100TR; PerkinElmer).

The remaining cells in the plates were dissolved in 0.1 M NaOH, and an aliquot taken to measure cellular protein by Lowry assay [17]; [3 H] counts were normalized to cellular protein levels and expressed as cpm/mg cell protein.

#### **RNA extraction and real-time PCR**

Total RNA was extracted from J774A.1 macrophages with Master PureTM RNA purification kit. Thermo Scientific cDNA kit was used to extract 1 μg of cDNA. Using Absolute Blue qPCR ROX mix, products of the reverse transcription were subjected to quantitative PCR using TaqMan gene expression assays with a Rotor-Gene 6000 amplification detection system; 3-hydroxy-3-methylglutarylcoenzyme reductase (HMGCR) levels were normalized to GAPDH mRNA. Primers and probes were designed by Primer Design (South Hampton, UK).

#### **Human serum-mediated cholesterol efflux**

J774A.1 macrophages were pre-incubated with [3 H]-labeled cholesterol (2.5 mCi/mL) for 1h at 37°C, followed by cell wash with PBS and further incubation for 3h at 37°C with patients' serum (30μl). At the end of the incubation period, the media (500μl) were collected and [3 H]-labels were determined by liquid scintillation counting (LSC; b-counter, Packard Tri Carb

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2100TR, PerkinElmer, Waltham, MA). The remaining cells in each well were washed with PBS, solubilized overnight in 0.1 M NaOH, and the amount of [3 H]-labels in 500 μl of each cell lysate determined. Total cellular [3 H]-cholesterol was calculated as the sum of the radioactivity in the medium and radioactivity in the lysates. Human serum-mediated cholesterol efflux corresponded to the percentage of total [3H]cholesterol released into the medium.

## **Statistical analysis**

Each serum was analyzed in triplicate, and each experiment was separately performed three times. Statistical analyses used one-way analysis of variance followed by the Student-Newman -Keuls test for comparing differences among multiple groups. Results are expressed as mean ± SEM of 3 separate experiments.

#### **Results**

#### **The patient's serum biochemical parameters**

Information about the patients in each group included: age, gender, BMI, percentage HBA1C, CRP levels, number of patients with hypertension or diabetes, and those treated with statin (Table 1).

Serum total cholesterol and LDL-C levels were similar in all three groups. In contrast, the SCAD group had lower serum HDL levels compared to the NCAD and NSCAD groups (32% (p<0.02) vs. 31% (p<0.04), respectively) and increased levels of triglyceride (28% vs. 24%, p<0.24, respectively) (Figure 1).

## **The effect of the patient's serum on macrophage cholesterol content**

Cellular cholesterol mass was found to be significantly higher by 44% (p< 0.0003) or 23% (p< 0.008) in cells treated with serum of the SCAD patients, compared to cells treated with serum from NCAD or NSCAD patients, respectively (Figure 2A). In addition, cellular cholesterol mass was higher by 23% (p<0.01) in the NSCAD group compared to in the NCAD group (Figure 2A).

## **The effect of the patient's serum on macrophage cholesterol biosynthesis**

We observed a significant enhanced cholesterol biosynthesis rate of 53% (p<0.0006) and 33% (p<0.006), respectively, in cells treated with serum from the SCAD group, compared to the effect of serum from NCAD or NSCAD patients (Figure 2B). In addition, a non-significant higher cholesterol biosynthesis rate (14%) was noted in J774A.1 macrophages treated with serum from the NSCAD group compared to the NCAD group (Figure 2B).

## **The effect of the patient's serum on macrophage HMG-CR expression**

Furthermore, the enhanced macrophage cholesterol biosynthesis rate observed in cells treated with SCAD patients' serum was found to be associated with overexpression of

HMGCR mRNA levels by 27% (p<0.002) and 26% (p<0.003), respectively, in comparison to cells treated with serum from NCAD or NSCAD patients (Figure 2C). In addition, a higher  $(6%)$  but non-significant cholesterol biosynthesis rate was noted in cells treated with serum of the NSCAD compared to the NCAD group (Figure 2C).

#### **Table 1:** Clinical parameters of the NCAD, NSCAD and SCAD patients.



\*number of patients with hypertension medications; \*\*number of the patients with statins medications; BMI: Body Mass Index; CRP: C Reactive Protein



Figure 1: Serum lipid profile of patients from the NCAD, NSCAD or SCAD groups.



**Figure 2A:** Effect of the patient's serum from the 3 different CTCA groups on macrophage cholesterol mass. J774A.1 macrophages were incubated for 18h with 30μl of the patient's serum, followed by determination of cellular cholesterol levels, as described in the Methods section. Results are expressed as mean ±SEM (n=13, in each group). # p< 0.0003 SCAD vs. NCAD, \*\* p< 0.008 SCAD vs. NSCAD, \* p<0.017 NSCAD vs. NCAD.

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J774A.1 macrophage HMGCR mRNA expression. Results are expressed as mean ±SEM (n=8, in each group) # p< 0.003 SCAD vs. NCAD, \*\* p< 0.002 SCAD vs. NSCAD. J774A.1 macrophages were incubated for 18h with 30μl of the patient's serum, followed by determination of HMGCR mRNA.

## **The effect of the patient's serum on serum-mediated cholesterol effl ux from macrophages**

It should be noted that there was no significant difference in the extent of serum-mediated cholesterol efflux rate from the macrophages, when comparing the three groups. Results were expressed as % serum-mediated cholesterol efflux vs. baseline (control cells incubated only with medium). Efflux rate in the SCAD group was 42±8%, compared to 40±9 and 41±7 percent, respectively, in the NSCAD and NCAD groups.

## **Discussion**

Anti- or pro-atherogenicity of human serum is determined by its ability to attenuate or induce lipid accumulation in cultured macrophages [19]. In the current study, we clearly demonstrated, for the first time, that serum from patients with significant coronary atherosclerosis (as indicated by CT

angiography), substantially increased J774A.1 macrophage cholesterol content.

Although serum cholesterol and LDL-C levels were similar in all three CTCA groups, incubation of macrophages with serum derived from atherogenic patients resulted in increased cellular cholesterol levels compared to less atherogenic patients, or to subjects with no atherosclerosis. This effect could be attributed to the significant increase in cholesterol biosynthesis rate, secondary to the observed up-regulation of HMGCR (rate limiting enzyme in cholesterol biosynthesis) expression [20]. Similarly, it was previously shown that serum taken from patients with angiographically coronary atherosclerosis induced cholesterol deposition in smooth muscle cells from human aortic intima [14]. In that study, cholesterol biosynthesis rate and HMGCR expression were not analyzed. However, the authors found in patient's blood an LDL fraction prone to multiple modifications, including desialylation and oxidation. This LDL fraction resulted in cholesterol accumulation in cultured intima smooth muscle cells [14].

The role of HDL cholesterol in CAD is well investigated. Epidemiological studies indicate an inverse relationship between HDL cholesterol levels and risk for CAD, independent of LDL levels [10-12]. On the other hand, recent data provide evidence that high HDL levels are not always protective [11]. HDL may provide cardiovascular protection by promoting reverse cholesterol transport from macrophages [21].

Although the SCAD patients had low HDL cholesterol levels, as compared to NSCAD or NCAD patients, the ability of their serum to induce cholesterol efflux from J774A.1 macrophage was similar, suggesting the importance of investigating the functionality of the HDL, and not just HDL cholesterol concentration as a risk factor for CAD. Indeed, it was previously demonstrated that HDL from CAD patients is dysfunctional and pro-oxidant/pro-inflammatory in macrophages [22] and triggers macrophage apoptosis [23].

Associations between the presence and the severity of CAD, and the Percentage of Cholesterol Efflux Capacity (PCEC) and HDL cholesterol level in patients who underwent CTCA were investigated [24]. While PCEC was not associated with the presence or severity of CAD, total PCEC and HDL-C in patients with CAD were significantly lower than in patients without CAD.

Findings from several large studies indicate that elevated levels of triglycerides (either fasting or nonfasting) or, more specifically, triglyceride-rich lipoproteins and their remnants, are independently associated with increased risk of CVD. Increased triglyceride levels were noted in the SCAD patients, compared to NSCAD or NCAD patients. It was previously shown that triglyceride-rich lipoproteins induced cholesterol accumulation in J774A.1 macrophages [25] and THP-1 macrophages [26]. Similarly, in our study, the increased triglyceride levels in the SCAD patients could contribute to cholesterol accumulation in macrophages.

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## **Conclusion**

The SCAD patients' serum atherogenicity can contribute to macrophage cholesterol accumulation, foam cell formation, accelerated plaque formation, and thus to a higher morbidity mortality percentage in these patients.

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