Clinical and Experimental Immunology ORIGINAL ARTICLE

doi: 10.1111/cei.13315

Circulating lectin pathway proteins do not predict short-term cardiac outcomes after myocardial infarction

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Summary

Despite improvements in treatment, coronary artery disease is still responsible for one-third of all deaths globally, due predominantly to myocardial infarction (MI) and stroke. There is an important potential in developing new strategies for treatment of patients with these conditions. Inflammation, and in particular the actions of the complement system, has emerged as part of the pathogenesis in reperfusion injury in patients with MI. To further qualify this, we examined the association between the plasma levels of lectin pathway proteins and myocardial end-points, left ventricular ejection fraction (LVEF) and infarct size in a cohort of patients with ST-elevation myocardial infarction (STEMI). A blood sample was drawn the day after percutaneous coronary intervention from 73 patients with STEMI. The primary end-points, LVEF and infarct size, were measured with magnetic resonance imaging 6-9 days after the infarct. Complement pattern-recognition molecules of the lectin pathway (mannanbinding lectin, H-ficolin, L-ficolin and M-ficolin) were analysed along with soluble membrane attack complex (sMAC) and C-reactive protein (CRP) in plasma with immunofluorometric assays <50%. CRP correlated negatively with LVEF, regression coefficient = -0.17 (P = 0.01). None of the lectin pathway proteins correlated to LVEF or infarct size, nor did soluble membrane attack complex (sMAC). There were no differences in plasma levels of these complement proteins when comparing patients with ejection fraction <50% to patients with ejection fraction <50%. Pattern-recognition molecules of the lectin pathway and sMAC do not predict shortterm cardiac outcomes after MI.

Keywords: complement system, ischaemia/reperfusion injury, lectin pathway, myocardial infarction

Introduction

Coronary artery disease (CAD) is one of the leading causes of mortality and morbidity worldwide. Although there has been an improvement in prognosis, there is still a residual risk which demonstrates the scientific and therapeutic challenge we are facing in the search for improved treatment of CAD-related diseases such as myocardial infarction (MI) [1,2]. The pathogenesis of acute MI is a complex multi-factorial process, in which inflammation is an essential causative factor [3,4]. Inflammation normally arises in response to tissue damage caused by pathogens or irritants. The primary role of inflammation is defence, but paradoxically inflammation is suggested to worsening the outcome of MI during reperfusion. A possible explanation is that dying cells present patterns of neoepitopes, which is recognized by pattern-recognition molecules and thus activate the complement system during reperfusion. Recently, the association between the lectin pathway and ischaemia/reperfusion (I/R) injury has generated scientific attention in a comprehensive review and calls for further investigation in clinical cohorts [5].

The lectin pathway is one of three pathways capable of activating the complement system, which integrates the innate and the adaptive immune systems. The lectin pathway is activated by five soluble pattern-recognition molecules: mannan-binding lectin (MBL), H-ficolin, L-ficolin, M-ficolin and collectin-LK [6–8], followed by activation of the MBL-associated serine proteases (MASPs), and ultimately formation of the membrane attack complex (MAC). Collard *et al.* was the first to investigate the association between MBL and I/R injury in an animal model. They observed MBL deposition in ischaemic rat heart and furthermore illustrated that MBL was responsible for iC3b deposition after oxidative stress in an *in-vitro* study [9]. The research performed in murine I/R models is persuasive, and recently Clark *et al.* showed that mice treated with anti-MASP-2 antibody have smaller infarcts than their controls [10].

In clinical research, the association between the lectin pathway and myocardial ischaemia is controversial. On one hand, MBL is shown to be protective in the development of coronary artery disease, but on the other hand it is suggested to be harmful [11,12]. L-ficolin and combinations of L-ficolin, MBL and MAp44 are shown to be associated with left ventricular dilatation after MI [13]. We showed that proteins involved in regulation of the lectin pathway, MAp44, MASP-1 and MASP-3, are found at higher concentrations in patients with acute MI compared to healthy controls, but these proteins had no association with infarct size [14]. In contrast, two small clinical studies reported lower MASP-2 level in patients with acute MI compared to healthy controls [15,16].

In this study, we aimed to investigate if markers of structural and functional cardiac damage [infarct size and left ventricular ejection fraction (LVEF)] in patients with STEMI were associated with plasma concentration of lectin pathway components. We therefore examined pattern-recognition molecules which are central for initiation of the lectin pathway (MBL, M-ficolin, L-ficolin, H-Ficolin) and the soluble membrane attack complex (sMAC) as an end-point of complements activation.

Methods

Study population

This study includes STEMI patients originally enrolled into a prospective randomized study at Karolinska University Hospital, Sweden and admitted to the coronary care unit for primary percutaneous coronary intervention (PCI) [17]. In brief, 89 patients met the inclusion criteria; chest pain >30 min and ≤ 6 h, ST elevation ≥ 0.1 mV (≥ 0.2 mV in V1–V3) in two contiguous electrocardiogram (ECG) leads or left bundle branch block, and a thrombolysis in myocardial infarction (TIMI) grade 0 flow in the infarct-related artery. The patients were randomly allocated to either primary PCI (n = 45) or primary PCI

and postconditioning (n=44). In the postconditioning group, an angioplasty balloon was positioned upstream for the primary stent at a pressure of 2–4 atm within the first minute of reflow. The balloon was reinflated four times for 1 min, each separated by 1 min of reflow. Seven patients randomized to the primary PCI group and seven patients from the postconditioning group were excluded because of unwillingness, poor image quality, sudden death (n=1), missing plasma samples or the protocol was not followed.

The study thus included 73 patients in total; 36 patients were allocated to the control group and 37 patients were allocated to the postconditioning group. The original study by Sörensson *et al.* [17], investigating the effect of postconditioning after PCI in STEMI, showed no difference in LVEF or infarct size between the two randomized groups. Based on this, and the fact that there were no differences in the complement proteins or C-reactive protein between the control group and the postconditioning group (see Table 1 in the Result section), we pooled the two groups into one group of STEMI patients.

The original randomized study was approved by the local ethics committee at the Karolinska Institutet and conducted in accordance with the Declaration of Helsinki. Written informed consent was given by all patients included [17].

Ethylenediamine tetra-acetic acid (EDTA)-plasma was drawn the day after reperfusion and stored at -80° C for later analysis.

Clinical end-points

Infarct size and LVEF were measured on days 6–9 after the infarct with magnetic resonance imaging (MRI), as previously described in the original study [17]. In brief, the MRI was performed in the supine position with an

Table 1. Mean plasma concentration of the complement proteins are divided into two groups: percutaneus coronary intervention (PCI) and PCI with postconditioning.

	'	PCI + postconditioning	5
	PCI (mean \pm s.e.)	$(mean \pm s.e.)$	P-value
MBL (μg/l)	1847 ± 287	1855 ± 260	0.82*
M-ficolin (μg/l)	4166 ± 207	4700 ± 177	0.06
L-ficolin (µg/l)	3294 ± 159	3293 ± 160	0.99
H-ficolin (μg/l)	$22\ 584 \pm 648$	$23\ 090 \pm 748$	0.61
sMAC (μg/l)	136 ± 6	158 ± 15	0.69*
CRP (µg/l)	17 ± 3	22 ± 3	0.17

MBL = mannan-binding lectin; sMAC = soluble membrane attack complex; CRP = C-reactive protein. *Student's t-test was used to compare and test the levels between the groups. Mannan-binding lectin (MBL) and soluble membrane attack complex (sMAC) were tested with Wilcoxon's rank sum test.

eight-channel cardiac coil. Gadolinium contrast (0·2 mmol/kg) was administrated prior to scanning the patient. The imaging protocol included Scout images, localization of the short axis and then covering the whole left ventricle with retrospectively gated cine steady-state free precession images. Delayed-enhancement images were acquired 15–20 min after injection of the contrast. The images were analysed with segmentation software (Segment version 1.8 R0857). Infarct size is expressed as the percentage of area at risk determined with bi-plane left ventriculography before primary PCI at admission day.

Analysis of plasma proteins

MBL concentration was measured using our in-house time-resolved immunofluorometric assay (TRIFMA) assay based on ligand recognition, as described elsewhere [18]. The lower level of detection was 5 μ g/l, and the intraand interassay coefficients of variance (CV) were both below 10%.

M-ficolin [19] and H-ficolin [20] were estimated with an in-house TRIFMA. The interassay coefficient of variance was 13% for M-ficolin and 16% for H-ficolin. Intraassay imprecision was below 10% in both assays. L-ficolin was measured using an enzyme-linked immunosorbent assay (ELISA) from Hycult Biotech (Uden, the Netherlands), according to the manufacturer's instructions. The intra- and interassay CVs are below 6 and 1%, respectively. Plasma sMAC was determined by an in-house TRIFMA, described in detail elsewhere [21]. The limit of detection for the sMAC assay was 1 μg/l. The intra- and interassay CVs were below 5 and 12%, respectively.

Levels of C-reactive protein (CRP) were quantified by commercially available monoclonal antibodies (Biotechne, R&D Systems, Minneapolis, MN, USA) using an in-house assay for high sensitivity CRP. Briefly, wells were coated with 0·1 μg anti-CRP antibody (MAB17071; R&D Systems) in 100 µl phosphate-buffered saline (PBS) overnight at 4°C. Residual protein-binding sites were blocked with PBS, 1% Tween 20 (Tw) for 1 h at room temperature and washed in PBS containing 0.05% Tween 20 (PBS/ Tw). Recombinant human CRP (NIBSC 85/506) was used as standard and samples were diluted 200- or 1000-fold in assay buffer (PBS, 1% Tw, 0.1% human serum albumin and 25 µM EDTA) and incubated overnight at 4°C. Bound CRP was determined by incubation with 25 ng biotinylated anti-CRP antibody (BAM17072; R&D Systems) in 100 µl assay buffer for 2 h at room temperature. The wells were washed and subsequently incubated with 10 ng Eu³⁺labelled streptavidin (Perkin Elmer, Waltham, MA, USA) in 100 µl assay buffer for 1 h at room temperature. After washing, bound Eu3+ was detected by the addition of 200 µl of enhancement solution (Perkin Elmer), 5 min of vigorous shaking and reading the time-resolved fluorescence on a DELFIA fluorometer (Victor³; Perkin Elmer). The limit of detection was $0.05~\mu g/l$ and the intra- and interassay CV were below 5 and 6%, respectively.

Statistical analysis

Assumptions of normally distributed data were checked with q-q plots and histograms. If data were not normally distributed, log-transformation was performed before statistical analysis. Results are reported as mean ± standard deviation (s.d.) or median (25th percentile; 75th percentile), as appropriate. Pearson's correlation was used to test the relationship of plasma levels of sMAC, ficolins and CRP with LVEF and infarct size. For the association between MBL and the cardiac outcomes we used the Spearman's rank correlation test due to the non-normal distributed MBL plasma levels. A *t*-test was used to test the difference between two groups, but due to unequal variance between groups when measuring H-ficolin and sMAC, we used Wilcoxon's rank-sum test to test our hypothesis.

The statistical analysis was performed by STATA version 15.0 for Mac OS, and figures are made in GraphPad Prism version 8.0.1.

Results

Levels of plasma CRP did not vary between the two groups defined in the original study: $22.7 \mu g/l$ [95% confidence interval (CI) = 16.9, 28.5] for the PCI and postconditioning group *versus* $17.4 \mu g/l$ (95% CI = 12.1, 22.6) in the PCI group, P = 0.17, nor did the two randomized groups differ when looking at the lectin pathway proteins; see Table 1.

Plasma CRP correlated negatively with LVEF, with a regression coefficient = -0.17, P = 0.01. This was expected, and indicates that higher CRP correspond to worsening and lowering of LVEF; see Fig. 1.

We found no association between LVEF or infarct size and plasma levels of the pattern-recognition molecules from the lectin pathway or sMAC (Table 2). When stratifying the patients into the original randomized groups the cardiac outcomes remained without an association with complement factors (results not depicted).

In order to investigate if the lectin pathway alter cardiac function after MI, we divided patients into two groups; low ejection fraction (<50%) and normal ejection fraction ($\ge50\%$) in accordance with the American Heart Association [22]. We found no differences in plasma levels of sMAC, MBL, M-, L- and H-ficolins or CRP when comparing patients with EF <50% (n=40) with patients with EF $\ge50\%$ (n=33); see Fig. 2 for details.

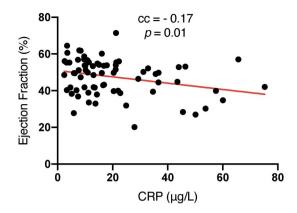


Fig. 1. Left ventricular ejection fraction plotted with C-reactive protein (CRP) on the *x*-axis and the fitted linear model in red.

Finally, we examined if the patients deficient in MBL (<100 ng/ml) differed in LVEF or infarct size from the patient with sufficient MBL. We found no differences between these two groups.

Discussion

In this study we show that the plasma level of CRP measured the day after PCI in STEMI patients is negatively associated with LVEF (Fig. 1). The present findings are in line with previous studies [23,24], which have shown that increasing CRP levels are correlated with decreased left ventricular function; however, when stratifying the patients into LVEF above and below 50%, the disadvantages with having a high CRP level disappear in our study (Fig. 2).

Although inflammation is known to play an important role after MI, the exact inflammatory mechanism is not clear. Several studies using animal models support a role for the lectin pathway in the severity of MI after reperfusion [9,10,25], but only a limited number of human clinical studies have investigated this. Even though we found an association between inflammation (CRP) and LVEF, the pattern-recognition molecules of the lectin pathway and

sMAC, as a measure of downstream complement activity, was not associated with LVEF or infarct size following MI. However, our study cannot rule out local production of complement proteins and/or local capture of the lectins to neoepitopes in the heart. In theory, this may contribute to local tissue damage and will not necessarily be reflected in the circulation. The pathogenesis of apoptosis of cardiomyocytes following MI is highly complex, and is probably the net result of insufficient reperfusion, inflammatory response, reduced clearance and complement activation [26]. As an example, pentraxins (e.g. CRP, PTX3) are also described as pattern-recognition molecules and are reported to be capable of modulating the complement system both via the classical and lectin pathways, and may thus be involved in the inflammatory processes [27,28]. Due to the pleiotropic effects of PTX3, it is still debated if PTX3 is cardioprotective or a negative prognostic factor in cardiovascular diseases [29].

A potential association between lectin pathway proteins and LVEF or infarct size could, in theory, be obliterated by selection bias. As an example, the inclusion criteria require a TIMI grade at 0, which is defined as 'no perfusion' in the coronary artery involved. The STEMI patients are therefore a very homogeneous group, which narrows the variability of the two outcomes, LVEF and infarct size.

By determining sMAC levels, we assume that this reflects the activation of the entire complement activation. However, we do not know if MAC is deposited locally in the heart and not reflected by the soluble form.

No pharmaceutical drug targeting the complement system is approved in MI treatment. Panagiotou et al. [5] have reviewed clinical trials using the complement inhibitor C1 esterase inhibitor (C1INH) in human myocardial I/R injury [30,31]. C1INH is an obvious first choice in the search for treatment against IR injuries, as it has an additional effect by preventing cells to alter into procoagulatory and anti-fibrolytic states [32]. The most recent clinical study, a randomized clinical trial with 78 STEMI patients, showed a significantly better

Table 2. Linear regression, correlation analyses.

	LVEF		Infarct size	
	Regression coefficient (95% CI)	P-value	Regression coefficient (95% CI)	P-value
MBL (μg/l)	Spearman	P = 0.95	Spearman	P = 0.71
M-ficolin (µg/l)	-1.88 (-10.8, 7.07)	P = 0.67	-0.24 (-15.41, 14.92)	P = 0.97
L-ficolin (µg/l)	0.00 (-0.00, 0.00)	P = 0.16	-0.00 (-0.00, 0.00)	P = 0.74
H-ficolin (μg/l)	0.00 (-0.00, 0.00)	P = 0.42	-0.00 (-0.00, 0.00)	P = 0.94
sMAC (μg/l)	0.00 (-0.02, 0.02)	P = 0.83	0.02 (-0.02, 0.05)	P = 0.30
CRP (µg/l)	-0.17(-0.31, -0.04)	P = 0.01	0.14 (-0.10, 0.38)	P = 0.25

Pearson's correlation was applied to check if complement proteins correlated with left ventricular ejection fraction (LVEF) or infarct size. The regression coefficient was determined and the corresponding 95% confidence interval (CI). Spearman's rank correlation test is marked with 'Spearman'.

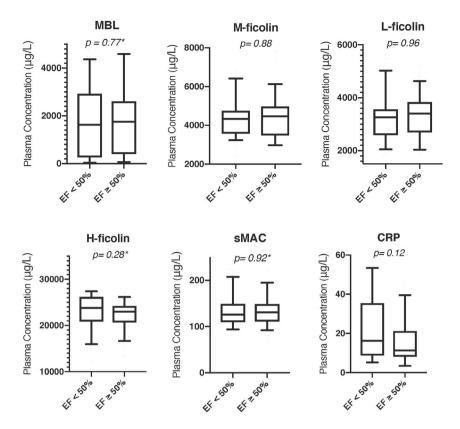


Fig. 2. Plasma concentration of mannan-binding lectin (MBL), M-ficolin, L-ficolin, H-ficolin, soluble membrane attack complex (sMAC) and C-reactive protein (CRP) given with 10th percentile, first quartile, median, third quartile and 90th percentile and divided into ejection fraction <50% and ejection fraction ≥50%.

myocardial contractility in the group receiving plasma derived C1INH compared to placebo [33]. Thus, pdC1INH is safe, with no adverse effects reported, and so far the clinical studies indicate that C1 inhibition is effective. These clinical trials have not been conducted with the state-of-the-art revascularization, such as stents and anti-thrombotic treatment, which is questioning if any additive effect of C1 inhibition remains when testing together with state-of-the-art treatment. In the same trial, C3a and C4a plasma concentrations are increasing during and after surgery (coronary artery bypass grafting). These concentrations are reaching a maximum concentration approximately 2 h after surgery. This suggests that the complement system is mainly activated during and immediately after reperfusion.

The association between MBL and cardiac I/R injury is contentious. Personen *et al.* found higher serum levels of MBL and lower serum concentration of C3 in a cohort of patients with unstable angina pectoris or acute MI when compared with healthy controls [11]. Our data confirmed this, with higher serum levels of MBL and lower sMAC in patients with STEMI as reported in healthy controls [21]. Plasma MBL levels could be persistently higher in patients with high MI risk or it might be an

acute immunological response due to cardiac ischaemia, thereby adversely causing enhanced thrombosis and complement-mediated IR injury. Other studies suggest MBL as being protective in the development of acute coronary syndrome, as it lowers the atherosclerosis by eliminating infection and clearing cellular debris and apoptotic cells. Saevarsdottir *et al.* found that high MBL levels were associated with decreased risk of MI [12], and Vengen *et al.* found that individuals with MBL-deficiency associated genotypes had twice the risk of developing a MI [34].

In the present study we did not focus on whether MBL or ficolin levels were higher or lower in patients with STEMI compared to healthy controls, but merely studied the impact of the lectin pathway on short-term cardiac outcomes, LVEF and infarct size after ischaemia. Our study shows no correlation between levels of the initiating molecules of the lectin pathway, nor the final complement activation end-product and the magnitude of the cardiac damage. This support our previous finding, that even though the lectin pathway components are elevated in patients with MI, they are not directly associated with the extent of cardiac damage [14].

The present findings must be carefully interpreted due to the cross-sectional design and small number of patients. In particular, it is unclear whether the plasma levels of complement factors are stable and represent the level prior to the MI or if local inflammation have caused acute alteration in the complement system during MI. Furthermore, peak CRP levels would have been more appropriate to examine due to liver production latency.

In conclusion, this study demonstrates that structural and functional damage following ischaemia are not associated with physiological variations in plasma concentration of MBL or ficolins. In the acute phase, the plasma levels of the pattern-recognition molecules of the lectin pathway (MBL and ficolins) or the end-product of total complement activation (sMAC) do not appear to be of predictable prognostic value of short-term cardiac outcomes after MI. Thus, it seems that once cardiac ischaemia has occurred, the patients with low plasma levels of either MBL or ficolins are not protected against the consequences of cardiac I/R compared to the patients with high plasma levels.

Future research must aim for randomized clinical trial using up-to-date revascularization and testing a specific lectin pathway inhibitor. Furthermore, to evaluate complement proteins in MI patients as a prognostic marker it is important to obtain bigger sample sizes and retrieve plasma samples during or immediately after the reperfusion.

Acknowledgements

The authors thank Lisbeth Jensen and Annette G. Hansen at the Department of Microbiology and Immunology, Aarhus University, and Zahra Partovi Nasr at the Medical Research Laboratory, Aarhus University, for laboratory assistance. This study was supported by Aarhus University.

Disclosures

The authors declare that there are no conflicts of interest concerning the publication of this paper.

Author contributions

P. S., L. M., T. K. H. and M.B. designed the study. The laboratory work was conducted by C. B. H. and the above-mentioned laboratory technicians with supervision from S. T. and M. B. C. B. H. conducted the analysis of raw data and performed the statistics with supervision from J. A. Ø. and M. B. All authors contributed to writing this article.

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