Research paper

Early increase of specialized pro-resolving lipid mediators in patients with ST-elevation myocardial infarction

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Abstract

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Background: Termination of acute inflammation is an active process orchestrated by lipid mediators (LM) derived from polyunsaturated fatty acids, referred to as specialized pro-resolving mediators (SPM). These mediators also provide novel therapeutic opportunities for treating inflammatory disease. However, the regulation of these molecules following acute myocardial infarction (MI) remains of interest.

Methods: In this prospective observational study we aimed to profile plasma levels of SPMs in ST-elevation MI (STEMI) patients during the first week following MI. Plasma LM concentrations were measured in patients with STEMI (n = 15) at three time points and compared with stable coronary artery disease (CAD; n = 10) and healthy controls (n = 10).

Findings: Our main findings were: (i) Immediately after onset of MI and before peak troponin T levels, STEMI patients had markedly increased levels of SPMs as compared with healthy controls and stable CAD patients, with levels of these mediators declining during follow-up. (ii) The increase in SPMs primarily reflected an increase in docosapentaenoic acid- and docosahexaenoic acid-derived protectins. (iii) Several individual protectins were correlated with the rapid increase in neutrophil counts, but not with CRP. (iv) A shift in 5-LOX activity from the leukotriene B4 pathway to the pro-resolving RvTs was observed.

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1. Introduction

Cardiovascular disease (CVD) is the leading cause of death worldwide of which atherosclerotic disorders are the most important [1]. Atherosclerosis is now considered a chronic inflammatory disease, with an interaction between lipids and inflammation as the major characteristic. In clinical studies, increased levels of inflammatory markers are associated with accelerated disease and worsened prognosis following atherosclerotic complications like myocardial infarction (MI) [2,3]. MI leads to a sterile inflammatory reaction involving recruitment of inflammatory cells and activation of inflammatory pathways within the myocardium, responses that are vital for tissue repair [4]. However, if sterile inflammation is prolonged and not resolved, such inflammatory responses could aggravate tissue damage with unfavorable effects on the myocardium. Indeed, resolution of inflammation is a prerequisite for restoration of tissue integrity and function following MI [4,5].

Evidence before this study

MI leads to a sterile inflammatory reaction, a response vital for tissue repair. However, if the inflammation is prolonged and not resolved, it could aggravate tissue damage with unfavorable effects on the myocardium. Indeed, resolution of inflammation is a prerequisite for restoration of tissue integrity and function following MI. Specialized pro-resolving lipid mediators (SPMs) are key effectors of resolution of inflammation and are endogenously formed from n-3 polyunsaturated fatty acids (PUFAs), the balance of pro-inflammatory mediators and SPMs regulates the duration and strength of the inflammatory response. To this end, however, the role of SPMs and resolution in acute MI is unexplored in humans. We hypothesized that timely induction and resolution of inflammation is required for optimal MI healing and aimed to profile plasma levels of SPMs in STEMI patients during the first week following MI. The sources searched were all PubMed publications available and the search terms were myocardial infarction and resolution, with emphasis on ST-elevation myocardial infarctions.

Added value of this study

This is the first report on the regulation of SPMs during acute MI in humans. Our findings show that pro-resolving mechanisms are increased early during STEMI, indicating that the “inflammation breaks” are activated immediately after MI onset. Our novel findings, along with the existing evidence on resolution mechanisms in inflammatory disease, could represent the start of a new era in relation to targeting inflammation during MI, focusing not only on anti-inflammatory intervention, but also on enhancing the pro-resolving capacity.

Implications of the available evidence

Our data may lead to new targets for therapy in STEMI, not only focusing on how to down-regulate harmful inflammation, but also on how to enhance resolving and repair mechanisms. Future research should try to identify the most important pathways in these processes and their cellular targets, and if successful, these studies could lead to a new paradigm in the management of these patients.
developing signs of heart failure within the observation period, other known inflammatory comorbidities (e.g. autoimmune disease, infections, and malignancies), and patients using immune modulating drugs (e.g. steroids) and COX-inhibitors were excluded. Blood samples were also obtained from ten age- and sex-matched apparently healthy controls. The healthy volunteers were all apparently healthy based on clinical examination and history, mean age 61 years, and none used any medications. They all had high-sensitivity C-reactive protein (hsCRP) ≤2-5 mg/L and their clinical and biochemical characteristics are presented in Tables 1 and 2. One patient in each group used a daily n-3 PUFA supplement, in the STEMI group 240 mg n-3 PUFA daily, in the stable CAD group 2000 mg n-3 PUFA daily, and among healthy volunteers 1200 mg n-3 PUFA daily.

2.2. Blood sampling

Blood samples were collected from patients and controls at the time of inclusion (in STEMI patients, mean time from symptoms debut 2.5 h; Table 1). In blood samples from STEMI patients were also collected at day one and eight. In STEMI and stable CAD patients, peripheral arterial blood was drawn prior to procedural heparinization and catheterization on inclusion into endotoxin-free blood collection tubes with ethylenediaminetetraacetic acid (EDTA) as anticoagulant. In healthy volunteers and STEMI patients at day one and eight, peripheral venous blood collected into EDTA tubes was used. The EDTA tubes were immediately placed on melting ice and centrifuged within 30 min at 2000 × g for 20 min to obtain platelet-poor plasma. Immediately following centrifugation, the vials were stored in several aliquots at -80 °C for less than six months and thawed only once prior to analyses.

2.3. Targeted lipid mediator profiling

Plasma samples for liquid chromatography-tandem mass spectrometry (LC-MS/MS)-based LM profiling were extracted using solid-phase (C-18) extraction columns as previously described [14]. Prior to sample extraction, deuterated internal standards, representing each region in the chromatographic analysis (500 pg each) were added to facilitate quantification in 4 vol of cold methanol. Samples were kept at −20 °C for a minimum of 45 min to allow protein precipitation. Supernatants were subjected to solid phase extraction, methyl formate fraction were collected, brought to dryness and suspended in phase (methanol/water, 1:1, vol/vol) for injection on a Shimadzu LC-20 AD HPLC analyzer, Roche Diagnostics. An Agilent Poroshell 120 EC-C18 column (100 mm × 4.6 mm × 2.7 μm) was kept at 50 °C and mediators eluted using a mobile phase consisting of methanol-water-acetic acid of 20:80:0.01 (vol/vol/vol) that was ramped to 50:50:0.01 (vol/vol/vol) over 0-5 min and then to 80:20:0.01 (vol/vol/vol) from 2 to 11 min, maintained till 14-5 min and then rapidly ramped to 98:2:0.01 (vol/vol/vol) for the next 0-1 min. This was subsequently maintained at 98:2:0.01 (vol/vol/vol) for 5-4 min, and the flow rate was maintained at 0.5 ml/min. QTrap 6500 plus was operated using a multiple reaction monitoring method. Each LM was identified using established criteria including matching retention time to synthetic and authentic材料s and at least 6 diagnostic ions. Calibration curves were obtained for each using synthetic compound mixtures at 0.78, 1.56, 3.12, 6.25, 12.5, 25, 50, 100, and 200 pg that gave linear calibration curves with an r² values of 0.98–0.99 and co-efficient of variation of 1–4% [14,15].

2.4. Cytokine analyses

Interleukin (IL)-6, IL-8, and tumour necrosis factor (TNF) were analysed by an enzyme immunoassay (EIA) with the Meso Scale (Rockville, MD) V-Plex Human proinflammatory kit. Intra- and inter-assay coefficients of variation were < 10%.

2.5. Biochemical analyses

Blood samples were analysed consecutively using routine methods in Medical Biochemistry laboratories at Oslo University Hospital Rikshospitalet including analyses of hsCRP on a MODULAR platform (Roche Diagnostics, Basel, Switzerland) and hsTnT by electrochemiluminescence immunoassay (ELICA; Elecsys 2010 analyzer, Roche Diagnostics).

2.6. Statistical analyses

Multivariate analysis (PLS-DA) was performed using SIMCA 14.1 with mean centring and unit variance scaling. All other statistical calculations were performed with Prism 7 for Mac OS X (GraphPad Software, SD, California). All continuous variables were compared with paired or unpaired Student's t-tests and categorical variables with the Chi-square test or Fisher exact test for observations < 5. For comparison of more than two groups, ANOVA and repeated measures ANOVA were used and subsequent analyses were performed only if the one-way analysis of variance was significant. Association of SPMs to cardiac and inflammatory parameters was performed only on significantly elevated SPMs after MI onset with non-parametric statistics (Spearman-rho correlation). A value of p ≤ 0.05 was considered statistically significant, but should be interpreted with caution.

Ethics statement

The study was approved by the Regional Committee for Medical and Health Research Ethics of South-Eastern Norway and conducted according to the Helsinki Declaration. All participants provided written, informed consent.
3. Results

3.1. Study population

Baseline clinical characteristics were comparable in STEMI patients and stable CAD patients. However, the stable CAD patients were slightly older, had higher body mass index (BMI), and hypertension was more prevalent (Table 1). Further, all patients with stable CAD and STEMI had received aspirin (300 mg) prior to baseline sampling. Healthy controls were matched for age, gender and actual blood pressure with the two patient groups and for BMI with the STEMI group (Table 1). As shown in Fig. 1, STEMI patients had hsTnT and hsCRP peak levels on day 1. In contrast to this pattern, neutrophil counts peaked and were tripled already at MI onset compared with stable CAD and healthy volunteers, demonstrating an early inflammatory response during MI, with decreasing levels throughout the observation period. All recorded biochemical parameters are listed in Table 2.

3.2. Patients with STEMI have a distinct and early increase in peripheral SPM concentrations

SPMs and eicosanoids from the eicosapentaenoic acid (EPA), n-3 docosapentaenoic acid (n-3 DPA), docosahexaenoic acid (DHA), and arachidonic acid (AA) bioactive metabolomes were identified and quantified in accordance with published criteria, including matching retention times on liquid chromatography (LC) and tandem mass spectrometry (MS/MS) fragmentation spectra [15]. Supplementary Fig. S1a-b depict representative multiple reaction monitoring chromatograms of selected ion pairs for protectin (PD)1 and PD2n-3 DPA along with representative MS/MS spectra and diagnostic ions employed for their identification. To assess if STEMI was linked to changes in SPM profiles, we first performed a Partial least squares discriminant analysis (PLS-DA) (Fig. 2a top) with results obtained from LC-MS/MS profiling. The PLS-DA plot shows the systematic clusters among observations (closer plots presenting higher similarity in the data matrix) [16] and demonstrates a separation between the healthy controls, stable CAD, and STEMI clusters (actual levels with SEM for all the SPMs in each patient group are shown in Supplementary Table S1). The corresponding loadings plot (Fig. 2a below), that describes the magnitude and manner the SPMs contribute to the cluster separation in the score plot [16], demonstrated that plasma from STEMI patients was characterized by higher levels of several individual LMs such as PD1 and PD2n-3 DPA (VIP score ≥ 1.0).

Combining (cumulative values) all individual SPMs (DHA-, DPA-, EPA-derived and lipoxins) and pro-inflammatory LM (AA-derived leukotrienes, prostaglandins, and thromboxanes), respectively, STEMI patients had an almost doubling of SPM levels (Fig. 2b top) only hours after onset of MI symptoms and before the observed peak in hsTnT, compared with both healthy controls and stable CAD, with declining levels throughout the observation period. Of note, the levels of SPMs in plasma were identified within physiologically relevant concentrations: 1 PM-10 nM [17,18]. In line with the administration of aspirin, patients with stable CAD (p = 0.001) had decreased plasma eicosanoid concentrations (Fig. 2b, below) and in STEMI patients this was seen at onset of MI (p = 0.003) and during follow-up. Our findings so far show that STEMI patients have increased levels of SPMs immediately after MI onset and this response is most likely part of the endogenous damage limitation pathway to minimise secondary damage post MI.

3.3. STEMI patients have elevated levels of DHA- and n-3 DPA-derived protectins

Having found distinct SPM profiles in STEMI patients when compared with both CAD patients and healthy volunteers, we next examined the contribution of each of the LM families to the overall profiles. The definitions of which individual SPMs that are included in each SPM family are defined in Supplementary Table S1. In STEMI patients we observed an increase of DHA-derived SPMs (p = 0.03 and p = 0.02 vs CAD and controls, respectively) when compared with stable CAD and healthy volunteers (Fig. 3a). Of note, we also found increases in both n-3 DPA and DHA-derived protectins during the early hours post MI that decreased during follow-up to levels comparable with those measured in both stable CAD patients and healthy volunteers (3- and 4-fold increase, respectively; Fig. 3b-c).

Assessment of individual mediator concentrations with the protectin metabolomes demonstrated an increase in the DHA-derived SPM PD1 and its double deoxygenation isomer 10S,17S-diHDP in STEMI patients compared with stable CAD and controls, with a decrease during follow-up (Fig. 4a, right). The regulation of the n-3 DPA derived protectins followed similar dynamics as those observed for the DHA-derived congeners, whereby PD2n-3 DPA and the protectin pathway marker 10S,17S-diHDP, increased by three-fold and ten-fold, respectively, in STEMI patients at onset of MI compared with stable CAD and controls with normalization during follow-up (Fig. 4b).

Table 2

|                      | Healthy Volunteers (n = 10) | Stable CAD (n = 10) | STEMI (n = 15) |
|----------------------|-----------------------------|---------------------|----------------|
| hsTnT baseline, ng/L | 8 (0-9)                     | 15 (3-4)            | 79 (22-1)      |
| NT-proBNP, ng/L      | 64 (6-6)                    | 295 (58-2)          | 251 (106-3)    |
| hsCRP, mg/L          | 14-6 (0-3)                  | 1-2 (0-4)           | 146-0 (0-3)    |
| Leucocytes, 10^9/L   | 6-6 (0-5)                   | 6-9 (5-3)           | 13-4 (7-7)     |
| Neutrophils, 10^9/L  | 2-9 (0-6)                   | 3-9 (0-4)           | 12-9 (1-4)     |
| Monocyte, 10^9/L     | 0-6 (0-1)                   | 0-8 (0-1)           | 10-2 (0-1)     |
| Lymphocyte, 10^9/L   | 2-6 (0-5)                   | 1-8 (0-2)           | 1-9 (0-3)      |
| Platelets, 10^9/L    | 260 (19-6)                  | 260 (15-5)          | 244 (19-3)     |
| Haemoglobin, g/DL    | 81-1 (5-8)                  | 81-3 (4-6)          | 76-6 (2-8)     |
| Total cholesterol, mmol/L | 5-4 (0-3)                  | 4-2 (0-3)           | 4-9 (0-3)      |
| Triglycerides, mmol/L | 1-5 (0-2)                   | 1-3 (0-4)           | 1-6 (0-3)      |
| LDL cholesterol, mmol/L | 3-3 (0-3)                  | 3-5 (0-3)           | 3-2 (0-2)      |
| HDL cholesterol, mmol/L | 1-7 (0-2)                  | 1-6 (0-2)           | 1-3 (0-3)      |
| AST, U/L             | 26 (0-8)                    | 26 (1-1)            | 149-6 (22-1)   |
| ALT, U/L             | 25 (2-1)                    | 34 (4-3)            | 40 (4-6)       |
| HbA1c, %             | 5-4 (0-1)                   | 5-6 (0-1)           | 5-7 (0-1)      |

Data are given as mean (SEM). hsTnT, high-sensitivity Troponin T; BNP, brain natriuretic protein; hsCRP, high-sensitivity C-reactive protein; Hb, haemoglobin; LDL, low density lipoprotein; HDL, high-density lipoprotein; AST aspartate aminotransferase; ALT, alanine aminotransferase.
In contrast with the protectin pathways, the production of RvD5n-3 DPA, which we recently found to exert vasculo-protective actions [21], was reduced in all STEMI patients at all three intervals tested and in stable CAD patients when compared with healthy volunteers.

Taken together, these findings strongly indicate a modulated biosynthesis of several SPMs with a temporal up regulation of DHA- and n-3 DPA-derived protectins immediately after vascular injury suggesting that these mediators may play a role in limiting further tissue damage. In contrast, the potential vasculo-protective RvD5n-3 DPA was reduced.

3.4. Association of SPMs with markers of inflammation and hsTnT in the STEMI population

We next assessed whether there was a correlation between individual significantly regulated SPMs and markers of inflammation. Whereas we found no associations between significantly regulated individual SPMs and hsCRP, we found significant correlations between PD1 \( (r = 0.26, p < 0.05) \), 10S,17S-diHDA (PDX) \( (r = 0.28, p < 0.05) \), and PD2n-3 DPA \( (r = 0.32, p < 0.01) \) with neutrophil counts. Furthermore, data on cytokine levels (i.e. IL-6, IL-8 and TNF) are shown in Supplementary Fig. S2. While IL-8 and TNF were not correlated with the individual protectins, IL-6 levels was correlated with PDX \( (r = 0.35, p < 0.01) \) was also associated with increased levels of hsTnT.

3.5. Low pro-inflammatory LM levels in STEMI and stable CAD primarily reflect effects of aspirin

Aspirin influences biosynthesis of certain LM and the aspirin triggered pathway for AA is shown in Fig. 5a. As expected and shown by others [22], in accordance with the observed decrease in pro-

**Fig. 1.** STEMI patients present with elevated troponin T and inflammatory markers. Plasma from healthy volunteers \( (n = 10) \) and patients with stable CAD \( (n = 10) \) and STEMI \( (n = 15) \) were collected after MI onset. Repeated samples were drawn from STEMI patients. The figure show plasma levels of hSTnT, hsCRP, and neutrophils. All results are expressed as mean ± SEM. *\( p < 0.05 \), **\( p < 0.01 \), ***\( p < 0.001 \) vs. STEMI baseline. *\( p < 0.05 \), **\( p < 0.01 \), ***\( p < 0.001 \) for repeated measures ANOVA for STEMI-baseline, day 1, and 8.

**Fig. 2.** Patients with STEMI have a distinct and early increase in SPM biosynthesis. Plasma from healthy volunteers \( (n = 10) \) and patients with stable CAD \( (n = 10) \) and STEMI \( (n = 15) \) were collected after MI onset. Repeated samples were drawn from STEMI patients. LM profiles were obtained using LC-MS/MS. (a) Partial least squares discriminant analysis of the LM-profiles. Top panel, two-dimensional score plot; lower panel, two-dimensional loading plot. Gray ellipse in the score plot denotes 95% CI regions. (b) Cumulative values of SPMs and pro-inflammatory LMs (prostaglandins, leukotrienes, and thromboxane B\(_2\) ) after MI onset and during study follow up. All results are expressed as mean ± SEM. *\( p < 0.05 \), **\( p < 0.01 \), ***\( p < 0.001 \) vs. STEMI baseline. *\( p < 0.05 \), **\( p < 0.01 \), ***\( p < 0.001 \) for repeated measures ANOVA for STEMI-baseline, day 1, and day 8.
inflammatory LM (Fig. 2d), STEMI and CAD patients had lower plasma levels of the n-6 PUFA AA-derived PG family (i.e., PGD₂, PGE₂ and PGF₂α) and thromboxane (Tx)B₂ compared with healthy volunteers (Fig. 5b). In fact, as all patients in the stable CAD and STEMI group were given 300 mg aspirin prior to the first blood sampling the observed pattern is in line with the mechanism of action of aspirin [23]. In addition to a decrease in AA-derived inflammatory LM, aspirin may trigger the biosynthesis of the pro-resolving aspirin triggered (AT)-SPM and notably, patients with STEMI had raised levels of AT-PD₁ and AT-LXA₄ (Fig. 5c-d; Supplementary Table S1). There was a decline in AT-LXA₄ levels at day 1 following STEMI, but the differences were not significant, and forthcoming studies should clarify if these changes are real and not by chance.

3.6. Modulation of 5-LOX dependent mediators

As both pro-inflammatory leukotrienes (LT) and several SPM families, including Rv of the E-series, D-series, and T-series, are 5-LOX dependent pathways [7], we lastly looked into the balance and timing of their biosynthesis in STEMI. In terms of AA-derived 5-LOX products, LT levels as a whole were comparable in healthy controls and STEMI patients, but for 12-epi-6-trans-LTB₄, a biosynthetic pathway marker of the potent leukocyte chemoattractant LTB₄, the increase was significant (Fig. 6). Interestingly, SPMs in all the major 5-LOX dependent families were detected in circulation during STEMI and especially, RvT₄ was the most abundant with peaking levels on day 8 (p = 0.03 vs healthy volunteers). Of note we observed that the ratio of RvT₄ to LTB₄ was higher after one week compared with MI onset, potentially reflecting a shift in the product profile of the 5-LOX enzyme from LTB₄ to the pro-resolving RvT₄.

Having observed changes in both classic eicosanoid pathways as well as omega-3 derived SPM we next assessed whether there were changes in circulating of unesterified essential fatty acids that are substrates in the formation of the lipid mediators. In STEMI patients at baseline we found a ~5-fold increase in plasma n-3 DPA concentrations and ~2-fold increase in the concentrations of AA, EPA and DHA when compared with concentrations measured in healthy volunteers. Of note, levels of these fatty acids returned to those measured in healthy volunteers after 1 day. Thus, these findings highlight an acute increase in the plasma concentrations of lipid mediator substrates following STEMI.

3.7. SPM levels in relation to clinical characteristics

One control and one patient in each group (CAD and STEMI) used n-3 supplements prior to inclusion, but importantly, these individuals did not differ from the other individuals in their respective groups in relation to the actual SPM parameters (data not shown). Moreover, we cannot exclude that dietary differences (e.g., intake of fish) between the different patient groups could have influenced our findings. However, Supplementary Fig. S3 on circulating free fatty acid levels between the different patient groups show that the levels of all four fatty acids are elevated at baseline in STEMI patient, levels that rapidly return to concentration levels measured in both healthy volunteers and CAD patients. These data suggest that the increases in SPM concentrations are via an upregulation of SPM biosynthetic pathways and not merely reflecting difference in FA availability. Finally, the fact that only the CAD group included diabetic patients (20%) and only the STEMI group included...
smokers (53%) is clearly a limitation of study, but importantly, patients in these subgroups (diabetic and smokers) did not differ from the other individuals in their respective groups in relation to the individual significantly regulated SPMs (Supplementary table S2).

4. Discussion

While data on the regulation of inflammatory pathways during MI is abundant, this is, to the best of our knowledge, the first report on the regulation of SPMs during acute MI. We show that circulating SPMs are markedly modulated at the onset of STEMI, indicating an early activation of resolution processes during MI. We found that circulating SPM levels peak within hours after onset of MI symptoms, even before the observed maximum release of hsTnT, corresponding with the early neutrophil response in circulation. Moreover, in contrast to this rise in pro-resolving mediators, there was a marked decline in pro-inflammatory PG and TxB2 throughout the observation period, reflecting the use of aspirin as previously shown by others [22]. Finally, in contrast to the rapid increase in overall SPM biosynthesis, there seemed to be a delayed shift from pro-inflammatory to pro-resolving LM in the LOX-dependent pathway during STEMI. Our findings show that pro-resolving mechanisms are activated early during STEMI, underscoring that resolution of inflammation is an active and regulated process.

Our observations suggest that post-MI inflammation and wound healing programs are activated in parallel and indicate that the "inflammation breaks" are activated immediately after MI onset. Whereas this is the first report on resolution mediators during the acute phase of STEMI in humans, SPMs have been shown to modulate post-MI wound healing in experimental animal models [24–26]. Diminished resolution of inflammation worsened prognosis after experimental MI.
as well as promoted plaque instability [5,24]. We have reported data in patients with stable CAD showing diminished levels of several SPMs [27] and demonstrated the diurnal regulation and lower levels of RvDn-3 DPA in CAD patients admitted for PCI [13]. Herein we extend these findings by showing a much more marked regulation of SPMs in STEMI patients. Also, and most importantly, in contrast to the previously reported decreased RvD1 concentrations in stable atherosclerotic disorders, we found marked up-regulation of several SPMs immediately after onset of symptoms in STEMI patients, with a gradual decline during the first week. In contrast, however, the potential vasculo-protective RvD5n-3 DPA was reduced during MI, and therapy that counteract this decrease could be of potential interest in STEMI patients.

The early SPM response in STEMI was mainly driven by the DHA- and n-3 DPA-derived protectin families, and especially PD1, 10S,17S-diHDA (PDX), PD2n-3 DPA, and 10S,17S-diHDPA were higher in STEMI patients shortly after MI onset. DHA derived protectins are formed via the stereoselective conversion of DHA by 15-LOX, with both PD1 and 10S,17S-diHDA (PDX) carrying anti-inflammatory properties [19]. Moreover, 10S,17S-diHDA (PDX) have been shown to inhibit TxA2-induced platelet aggregation [28] and during mouse ischemia-reperfusion injury, PD1 administration before ischemia resulted in a reduction in functional and morphological kidney injury [29]. The n-3 DPA-derived protectins have been found to enhance the resolving capacity of macrophages, partly through induction of efferocytosis and a reprogramming of the macrophage phenotype [30]. Of note, PD2n-3 DPA reduces neutrophil recruitment during sterile inflammation [14] and interestingly, in contrast to maresins that are mostly produced by pro-resolving (M2) macrophages, protectins may also be produced by neutrophils [6]. Indeed, herein PD2n-3 DPA, as well as PD1, were positively correlated with neutrophil counts shortly after MI onset, and it is tempting to hypothesize that the increase in the protectin levels may represent a counteracting mechanism to dampen the harmful effect of the initial rise in neutrophils following STEMI.

Aspirin irreversibly inhibits COX-1 with a subsequent reduced biosynthesis of pro-inflammatory LM. All our patients (STEMI and CAD) received aspirin prior to baseline sampling, and as expected we observed suppressed biosynthesis of COX-1 dependent pro-inflammatory LM in these patients (PGs and TxB2). Furthermore, aspirin modifies the enzymatic activity of COX-2, where it promotes a shift in COX-2 activity to 15-LOX-like activity, with the consequential biosynthesis of AT-SPMs contributing to additional anti-inflammatory effects of aspirin [31]. Hence, the effects of low-dose aspirin beyond inhibition of PGs and thromboxane are becoming increasingly apparent. AT-PD1 exhibits significant higher metabolic stability than PD1 [19] and in our study AT-PD1 was clearly elevated throughout the observation period. Chiang et al. [22] showed that aspirin administered in doses ranging from 80 to 650 mg for 8 weeks to healthy volunteers both inhibited PG biosynthesis and increased AT-LXA4 in circulation, the latter a pattern that was also observed when assessing plasma concentrations of STEMI patients when compared to healthy volunteers (Supplementary Table S1). Of note we also found that 1 day after aspirin administration the concentrations of this mediator were markedly reduced. Given that STEMI patients received a single dose of aspirin, these observations are in line with the transient effects that aspirin exerts on changing the activity of leukocyte COX-2. Indeed, at variance to the biological actions of aspirin on platelet COX-1 which last up to 2 weeks since platelet COX-1
is not turned over in these cells, leukocyte COX-2, like many other leukocyte proteins, is turned over and therefore the acetylated protein is readily degraded.

In contrast to the early increase in 15-LOX products, we found a delayed increase in the biosynthesis of the 5-LOX SPMs. While 5-LOX activity may also be increased during the initial stages, this activity produces LTs. Later in the timeline, possibly regulated by protectins, 5-LOX remains in the cytosol where it produces RvTs. Moreover, Fredman et al. [7] also showed decreased levels of 5-LOX SPMs in unstable atherosclerotic lesion in both human samples and murine models of atherosclerosis and one might speculate if the late (1 week after MI) rise in 5-LOX SPMs could inhibit further plaque destabilization. Earlier reports have shown that 5-LOX deletion impairs wound healing and promotes cardiac rupture after MI, indicating a crucial role in post-MI inflammation [32]. If peak level was reached within the study observation period, or if the 5-LOX SPMs will continue to rise even further, is at present not clear and should be explored in forthcoming studies.

4.1. Study limitations

This study has some important limitations. First of all there is a low number of individuals included that limit the impact of our findings. Moreover, the use of aspirin in all patients but not in any controls may clearly have influenced comparative analyses between patients and controls. Furthermore, diet may influence levels of various SPM, and the lack of detailed dietary information of the study population limit the interpretation of the data. Finally, correlation analyses do not necessarily document any causal relationship, and forthcoming studies should include more mechanistic studies.

5. Conclusion

This is the first report on the regulation of SPMs during acute MI in humans. Our findings show that pro-resolving mechanisms are increased early during STEMI, indicating that the “inflammation breaks” are activated immediately after MI onset. These findings could
contribute to the start of a new era in relation to targeting inflammation during MI, focusing not only on anti-inflammatory intervention, but also on enhancing the pro-resolving capacity.

Supplementary data to this article can be found online at https://doi.org/10.1016/j.ebiom.2019.07.024.

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Declaration of Competing interests
None declared.

Author contributions
Fosshaug, Linn E performed and planned study, did lab work, and wrote article.
Colas, Romain A performed lab work and statistical analyses.
Ansensrud, Anne K recruited patients and sampled data.
Gregersen, Ida recruited patients, sampled data, and did lab work.
Nymo, Ståle, performed and advised on statistical analyses.
Sagen, Ellen L performed lab work.
Michelsen, Annika performed lab work.
Vinge, Leif Erik planned study, clinical advice, and wrote article.
Øie, Erik planned clinical advice, and wrote article.
Gullestad, Lars sampling of data, biobanking, and conceptual work.
Halvorsen, Bente sampling of data, biobanking, and conceptual work.
Vinge, Leif Erik planned study, clinical advice, and wrote article.

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