

ORIGINAL ARTICLE

Low-grade inflammation and the phenotypic expression of myocardial fibrosis in hypertrophic cardiomyopathy

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Accepted 17 January 2012 Published Online First 24 March 2012

ABSTRACT

Objective To investigate the role of inflammation in the phenotypic expression of myocardial fibrosis in hypertrophic cardiomyopathy (HCM).

Design Clinical study.

Setting Kuopio University Hospital and University of Eastern Finland, Kuopio, Finland.

Subjects Twenty-four patients with a single HCM-causing mutation D175N in the α -tropomyosin gene and 17 control subjects.

Main outcome measures Endomyocardial biopsy samples taken from the patients with HCM were compared with matched myocardial autopsy specimens. Levels of high-sensitivity C-reactive protein (hsCRP) and proinflammatory cytokines were measured in patients and controls. Myocardial late gadolinium enhancement (LGE) in cardiac MRI (CMRI) was detected.

Results Endomyocardial samples in patients with HCM showed variable myocyte hypertrophy and size heterogeneity, myofibre disarray, fibrosis, inflammatory cell infiltration and nuclear factor kappa B (NF- κ B) activation. Levels of hsCRP and interleukins (IL-1 β , IL-1RA, IL-6, IL-10) were significantly higher in patients with HCM than in control subjects. In patients with HCM, there was a significant association between the degree of myocardial inflammatory cell infiltration, fibrosis in histopathological samples and myocardial LGE in CMRI. Levels of hsCRP were significantly associated with histopathological myocardial fibrosis. hsCRP, tumour necrosis factor α and IL-1RA levels had significant correlations with LGE in CMRI.

Conclusions A variable myocardial and systemic inflammatory response was demonstrated in patients with HCM attributable to an identified sarcometric mutation. Inflammatory response was associated with myocardial fibrosis, suggesting that myocardial fibrosis in HCM is an active process modified by an inflammatory response.

INTRODUCTION

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online under the BMJ Journals unlocked scheme, see http:// heart.bmj.com/site/about/ unlocked.xhtml Hypertrophic cardiomyopathy (HCM), most commonly caused by mutations in sarcomeric genes, is a myocardial disease characterised by left ventricular hypertrophy, myocyte disarray and myocardial fibrosis.¹ The cardiac phenotype in HCM varies between unrelated individuals and also between family members with an identical diseasecausing sarcomeric mutation, suggesting that HCM is a complex inherited disease modified by other genetic and environmental factors.^{1–5} The molecular events triggered by the genotype or other factors that induce the cardiac phenotype, particularly fibrosis, remain to be determined.^{4–6}

Previous studies have shown that chronic inflammation decreases myocardial contractility, induces hypertrophy and promotes apoptosis and fibrosis, thus contributing to myocardial remod-elling.^{7–9} Inflammatory cytokines, tumour necrosis factor α (TNF α), interleukin (IL)-1 and IL-6, are not constitutively expressed in the normal heart, but are upregulated and produced as a stress response to myocardial injury or mechanical stress.⁸ TNFa and IL-6 can directly attenuate myocardial contractility and induce the development of myocyte hypertrophy, collagen deposition and fibrosis.⁸ IL-1 β has been correlated with myocardial collagen deposition, cardiomyocyte apoptosis and inflammation.¹⁰ The expression of both inflammatory and antiinflammatory cytokines is increased in heart failure.⁹ In addition, nuclear factor κ B (NF- κ B), a transcription factor regulating inflammatory genes, has been shown to be activated in patients with heart failure of various causes.¹¹

Circulating inflammatory cytokines are raised in HCM,¹²⁻¹⁴ but none of the previous studies has investigated the association of cytokine levels with cardiac phenotype determined by cardiac MRI (CMRI). Furthermore, there are no previous studies on NF-KB activation in HCM. Even more importantly, there are few data on myocardial histopathological findings or inflammatory response in patients with HCM caused by a specific sarcomeric gene mutation. Therefore, to investigate the role of inflammation in the phenotypic expression of myocardial fibrosis in HCM, we investigated cardiac histopathology, inflammatory cytokine and high-sensitivity C-reactive protein (hsCRP) levels and cardiac phenotype by CMRI in patients with HCM attributable to an identical HCM-causing mutation D175N in the α -tropomyosin gene (TPM1-D175N).

PATIENTS AND METHODS

The study protocol was approved by the ethics committee of the Kuopio University Hospital. All subjects gave written informed consent. Consent to the use of cadaver myocardial samples was

obtained from the Finnish Center for Legal Proctection in Health Care. Data of endomyocardial biopsy and cytokine findings have not been published in any form. Previously, we have published studies on CMRI-derived myocardial perfusion, myocardial late gadolinium enhancement (LGE), myocardial contractile impairment and inducibility of life-threatening arrhythmias in this same patient population.² ^{15–17}

Patients with HCM

Twenty-four patients from five Finnish families (11 men, 13 women; mean age, 42 ± 13 years; age range, 17-68 years) with HCM attributable to the mutation D175N in the α -tropomyosin gene $(TPM1-D175N)^{18}$ ¹⁹ were included in the study. In Finland, TPM1-D175N is one of the two founder mutations causing about 11% of all, and 25% of familial, HCM cases in eastern Finland.¹⁹ Of 24 subjects with TPM1-D175N, two did not meet the diagnostic criteria for clinical HCM and were regarded as healthy mutation carriers. All subjects with TPM1-D175N, however, are called 'patients with HCM' in the subsequent analysis. The study protocol described below was performed during one 2-day visit to the Kuopio University Hospital.

Control subjects

Seventeen healthy volunteers not related to patients with HCM and without a previous cardiac disease or drug treatment and with race, gender and age similar to the patients with HCM, were included into our study. Study protocol was identical to that for the patients with HCM, except for cardiac catheter-isation and endomyocardial biopsy.

For histopathological analyses, control myocardial specimens were obtained from 20 cadavers matched for age, gender and race without known cardiac disease (no history of a cardiac disease, normal macroscopic myocardial findings and no clear evidence of a specific myocardial disease on microscopic examination).

Clinical and echocardiographic evaluation

All patients with HCM and control subjects underwent an interview, physical examination, 12-lead ECG recording and echocardiography, as previously described.^{2 15-17}

CMRI protocol

CMRI cine, perfusion and LGE imaging were performed in patients with HCM and controls as previously described.^{2 15 16} CMRI cine imaging was performed in all patients, perfusion imaging in 17 patients with HCM and LGE imaging in 22 patients with HCM.

CMRI image analysis

Left ventricular (LV) characteristics and myocardial perfusion by CMRI were evaluated. LGE image analysis was performed in LV short-axis images at the levels of tips of the mitral valve leaflets and papillary muscles.¹⁶ In statistical analyses, the maximal value of the six segmental LGE heterogeneity values was used. Figure 1 shows LGE images of one control subject and three patients with HCM.

Coronary angiography

Coronary angiography was performed on 21 of 24 patients with HCM using standard angiographic techniques by the same cardiologist (JK).

Endomyocardial biopsy and autopsy myocardial samples

Myocardial specimens for histological analysis and immunohistochemistry were obtained from 20 patients with HCM.



Figure 1 Contrast-enhanced T1-weighted inversion recovery images in (A) a 52-year-old control subject; (B) a 45-year-old male patient with hypertrophic cardiomyopathy (HCM); (C) a 37-year-old female patient with HCM; (D) 19-year-old male patient with HCM. Patients with HCM (B–D) show intramyocardial focal high signal areas (arrows) with increased segmental late gadolinium enhancement.

Endomyocardial biopsy samples were obtained under fluoroscopic guidance from the right ventricle side of the interventricular septum with the standard endomyocardial bioptome. Control cadaver samples were selected from the archives of the department of clinical pathology of Kuopio University Hospital. Representative myocardial routine autopsy specimens from the anterior, posterior or septal wall of the heart were taken from 20 matched cadavers.

Histological and immunohistochemical methods

Histological methods and immunochemistry analyses are described in detail in the online supplementary information.

Laboratory determinations of cytokines and hsCRP

Plasma concentrations of TNF α , IL-6, IL-10, IL-1 β and IL-1RA were measured using assay kits from R&D Systems (Minneapolis, USA). hsCRP was measured using an Immulite analyser and a DPC hsCRP assay (DPC, Los Angeles, California, USA) in all 24 patients with HCM and in 17 controls.

Statistical analysis

Data are given as mean±SD. Statistical analyses were performed with a statistical software package (SPSS Win V.11.5., SPSS Inc). Because of skewed distribution, LV mass and all cytokines were analysed after logarithmic transformation. The differences between the patients and controls were assessed by Student t test. Mann—Whitney test was used to investigate the association between endomyocardial findings and proinflammatory cytokines. Pearson's correlation coefficients were calculated to investigate the association of cytokines with LGE in CMRI. Spearman's correlation coefficient was used to investigate the association of different histopathological features, the association between histopathological inflammation and fibrosis and between histopathological inflammation/fibrosis and LGE in CMRI.

RESULTS

Clinical, echocardiographic and CMRI characteristics

Clinical, echocardiographic and CMRI characteristics of the patients with HCM and controls have been published previously² ^{15–17} and are summarised in table 1. Patients with HCM had stable mild to moderate symptoms (90% of patients had New York Heart Association functional class I–II). None of the subjects with HCM had a history or clinical symptoms or signs of decompensated heart failure, myocarditis, systemic infection, chronic inflammatory disease or life-threatening arrhythmias. None of the patients had an intracardiac defibrillator. About one-third of the patients used cardiac medication, mostly β -blocking agents. None of the patients was taking ACE inhibitors or AT1 receptor antagonists or medication for heart failure.

None of the subjects with HCM had a significant LV outflow tract obstruction (>30 mm Hg) at rest. In CMRI, LV maximal wall thickness was increased in patients with HCM compared with controls. No difference was seen in global LV ejection fraction between patients with HCM and control subjects, but the number of hypokinetic segments was increased in patients with HCM compared with controls.² The LV perfusion reserve was lower and the maximal LV LGE increased in patients with HCM compared with controls.¹⁵

Coronary angiography in patients with HCM

Patients with HCM had normal coronary arteries except for one patient with HCM, who had <50% stenosis in the left anterior descending coronary artery and another patient who had <50% stenosis in the left anterior descending, intermediate and right coronary arteries.

Histological findings in endomyocardial samples

Table 2 shows the histopathological findings in haematoxylin–eosin stained endomyocardial samples in patients with HCM. Sufficient endomyocardial biopsy samples for histology were available for 16 of 20 patients with HCM. Histological samples showed variable amounts of heterogeneity of myocyte size, myocyte hypertrophy, myofibre disarray, myocardial fibrosis, inflammatory cell infiltration and intramyocardial small artery narrowing. Interstitial and perivascular fibrosis was found in about 90% of cases. Inflammatory cell infiltration, including

Table 1	Clinical,	echocardio	graphic a	nd CMRI	characteristics in	
control s	subiects a	nd patients	with hyp	ertrophic	cardiomvopathy	

Characteristics	Control subjects (n = 17)	Patients with HCN (n=24)	
Men/women	8/9	11/13	
Age, years	38±12	42±13	
V _{max} , m/s	1.3±0.2	1.4 ± 0.5	
LV MRI findings			
Maximal wall thickness, mm	9.7±1.7	19.5±4.9**	
Mass, g	123±32	151 ± 57	
End-diastolic volume, ml	146±25	122±45**	
End-systolic volume, ml	56 ± 15	52±29	
Hypokinetic segments, %	12±12	37±20**	
Ejection fraction, %	61±7	58±7	
Perfusion reserve	$1.80 {\pm} 0.58$	1.12±0.35*	
The heterogeneity of late-enhancement, %	12±3	18±11*	

 V_{max} indicates maximum velocity in the Doppler signal from the jet in left ventricular outflow tract.

Data are means \pm SD.

*p<0.05, **p<0.001

 $\mathsf{CMRI},$ cardiac magnetic resonance imaging; HCM, hypertrophic cardiomyopathy; LV, left ventricular.

Table 2	Endomyocardial biopsy findings in 16/20 patients with
hypertropl	hic cardiomyopathy (haematoxylin-eosin staining)

Biopsy findings	Number (%	
Heterogeneity in myocyte size		
No	3 (19)	
Moderate	6 (38)	
Marked	7 (44)	
Myocyte hypertrophy		
No	1 (6)	
Mild	11 (69)	
Moderate	4 (25)	
Myofibre disarray		
No	2 (13)	
Mild	6 (38)	
Moderate	6 (38)	
Marked	2 (13)	
Myocardial fibrosis		
No	2 (13)	
Mild	6 (38)	
Moderate	6 (38)	
Marked	2 (13)	
Inflammatory cell infiltration*		
No	10 (63)	
Mild	4 (25)	
Moderate	1 (6)	
Marked	1 (6)	
Intramyocardial small artery narrowing		
No	12 (75)	
Mild	2 (13)	
Moderate	1 (6)	
Marked	1 (6)	

*Eosinophilic granulocytes and mononuclear inflammatory cells.

mainly mononuclear inflammatory cells and eosinophilic granulocytes, was found in 37% of the patients. Narrowed intramyocardial small arteries were found in one-quarter of the patients. Figure 2 A–C shows typical, mild and marked histopathological findings in patients with HCM, respectively.

In control cadaver myocardial samples, mild heterogeneity of myocyte size was found in one of 20 specimens, mild myocyte hypertrophy in seven samples, mild myofibre disarray in one sample, mild interstitial fibrosis in five samples, mild inflammatory cell infiltration of mononuclear cells in one sample and eosinophilic granulocytes and intramyocardial small artery narrowing in none of the samples.

Immunohistochemical findings in endomyocardial samples

Immunostaining with rabbit anti-human antibody showed that seven of 11 (64%) patient endomyocardial biopsy samples had CD3 positivity implicating T-lymphocytes (two with moderate and five with weak positive staining). We could, however, replicate the CD3 positivity with NCL-CD3-PS1 in only one patient endomyocardial biopsy sample showing marked inflammatory response (figure 2C). In 20 control cadaver myocardial samples, weak CD3 positivity with rabbit antihuman antibody was found in two samples and CD3 positivity with NCL-CD3-PS1 in none of the samples. B-lymphocytes with M755 staining were not found in patients or controls. Occasional macrophages with MO814 were found in one patient sample only.

Picrosirius collagen staining was positive in 12 of 15 HCM samples. Extensive fibrosis was found in two cases (figure 3), moderate fibrosis in four cases and mild fibrosis in six cases. In





Figure 2 (A) Typical histopathology in an endomyocardial biopsy of a patient with hypertrophic cardiomyopathy (HCM). (a) A general view, haematoxylin and eosin (H&E) $\times 200$; (b, c) moderate fibre disarray, interstitial fibrosis (asterisk), moderate myocyte size heterogeneity and hypertrophy and scattered mononuclear inflammatory cells (arrows), H&E $\times 400$, $\times 630$. (B) Mild histopathological findings in a patient with HCM. (a) A general view, H&E $\times 200$; (b, c) mild fibre disarray, fibrosis (asterisk), myocyte hypertrophy and occasional mononuclear inflam20 control cadavers, mild fibrosis in Picrosirius staining was found in two cases.

NF-κB nuclear positivity was detected in eight of 15 patient endomyocardial biopsy samples. NF-κB nuclear activity was found in cardiomyocytes in four cases (three cases showed NF-κB positivity in 20–50% of nuclei, one case in 5% of nuclei) (figure 4). In three other cases, nuclear positivity of NF-κB was found in inflammatory cells. Endothelial NF-κB nuclear positivity was found in one case. In control cadaver myocardial specimens, no nuclear NF-κB positivity was detected.

Cytokines

Levels of hsCRP, IL-1 β , IL-1RA, IL-6 and IL-10 were significantly higher in patients with HCM than in control subjects (table 3). There was a trend towards higher TNF α levels in patients with HCM compared with control subjects (p=NS).

Associations between histopathological myocardial fibrosis, LGE and inflammatory response

In patients with HCM, the degree of histopathological myocardial fibrosis significantly correlated with LGE in CMRI (r=0.568, p=0.034). The grade of myocardial inflammatory cell infiltration correlated with fibrosis in histopathological samples (r=0.614, p=0.011) and with LGE in CMRI (r=0.541, p=0.046).

Levels of hsCRP were significantly associated with histopathological myocardial fibrosis (p<0.05). All other cytokine levels tended to be higher in patients with moderate or marked histopathological findings compared with those with no or mild findings (p=NS, data not shown). Cytokine levels did not correlate significantly with NF- κ B activation (data not shown).

Levels of hsCRP, TNF α and IL-1RA significantly correlated with maximal LGE in patients with HCM (table 4). There were no significant associations with IL-1 β , IL-6, IL-10 and CMRI-derived maximal LGE. There were no significant associations of cytokine levels with CMRI-derived maximal LV thickness, LV mass, LV diastolic or systolic volumes, global ejection fraction, or myocardial perfusion (data not shown).

DISCUSSION

Our study demonstrates that a low-grade myocardial and general inflammatory response is present in HCM attributable to a single well-documented causative sarcomeric mutation (*TPM1-D175N*). Variable low-grade myocardial inflammation in the patients with HCM was indicated by the presence of myocardial inflammatory cell infiltration and enhanced nuclear NF- κ B activity in the myocardium and by increased levels of hsCRP and circulating inflammatory cytokines. Myocardial inflammatory cell infiltration and levels of hsCRP significantly correlated with histopathological myocardial fibrosis and LGE in CMRI and TNF α and IL-1RA correlated with LGE, suggesting that myocardial fibrosis in HCM may be an active process modified by an inflammatory response.

[Continued]

matory cells (arrow), H&E \times 400, \times 630. (C) Severe HCM. (a) A general view: Weigert van Gieson staining highlights marked fibrosis (red, shown by asterisk), \times 200; (b) multiple mononuclear inflammatory cells (arrow); (C) showing CD3 positivity in immunohistochemistry (staining in brown, arrow). Original magnification \times 400. The patient had severe symptoms, marked left ventricular hypertrophy and inducible ventricular arrhythmia in ventricular stimulation. An intracardiac defibrillator was subsequently implanted.



Figure 3 Endomyocardial fibrosis in hypertrophic cardiomyopathy. Increased fibrosis is shown by Picrosirius Red staining (in red, asterisk). Cardiomyocytes are pale yellow stained. Original magnification \times 400.

Potential pathogenic mechanism for myocardial fibrosis formation in HCM

Based on our findings we suggest a pathogenic mechanism for myocardial fibrosis formation in HCM. We propose that myocardial fibrosis in HCM is likely to be an active process, in which primary injury-for example, mechanical stress,²⁰ due to disorganised sarcomeric and cellular architecture,²¹ myocardial ischaemia²² or neuroendocrinological activation,⁷ induces NF- κ B upregulation in the myocardium. NF- κ B, in turn, activates production of proinflammatory cytokines, inflammatory cell invasion into the myocardium and activation of fibroblasts, finally leading to myocardial fibrosis.^{7 8 11}

Histopathological myocardial phenotype and the causative mutation in HCM

Endomyocardial samples in our patients with HCM showed variable amounts of myocyte hypertrophy, myocyte size heterogeneity, myofibre disarray, myocardial fibrosis, low-grade



Figure 4 Nuclear NF- KB activation is detected in cardiomyocytes (arrows) in hypertrophic cardiomyopathy. $\times 400$.

Table 3	Circulating	levels of	⁻ cytokines	in	control	subjects	and	in
patients v	with hypertro	phic car	diomyopat	hy				

	Control subje	cts (n=17)	Patients with HCM (n=24)		
Cytokines	Mean±SD	Range	Mean±SD	Range	
hsCRP, mg/l	1.29±1.50	0.09-5.69	4.93±8.38**	0.33—38.90	
TNFα, pg/ml	2.33 ± 1.22	1.38-6.43	3.24 ± -2.98	1.56-16.50	
IL-1β, pg/ml	$0.18 {\pm} 0.08$	0.09-0.34	$0.32 \pm 0.25^*$	0.09-1.00	
IL- 1RA, pg/ml	216±76	101-391	490±577**	194-3023	
IL-6, pg/ml	1.11 ± 0.72	0.38-2.80	2.15±1.67**	0.75-7.55	
IL-10, pg/ml	0.86 ± 0.55	0.49-2.04	2.98±4.04***	0.49-20.86	

*p<0.05; **p<0.01; ***p<0.001

hsCRP, high-sensitivity C-reactive protein; IL-1B, interleukin 1B; IL- 1RA, interleukin 1 receptor antagonist; IL-6, interleukin 6; IL-10, interleukin 10; TNF α , tumour necrosis factor α .

inflammatory cell infiltration and intramyocardial small artery narrowing. Transgenic mice carrying a single missense mutation in codon 403 of the myosin heavy-chain gene exhibit variable amounts of histopathological hypertrophy, myocyte disarray, fibrosis and susceptibility to induced arrhythmias.²³ In humans, only limited information on histopathological myocardial findings in patients with HCM attributable to an identified sarcomeric mutation has been published.²⁴ As histopathology appears to vary to a great extent in humans and in transgenic animals with identical disease-causing mutations, other factors than the causative mutation necessarily contribute to histopathology in HCM.

Chronic myocardial inflammatory cell infiltration in HCM

Mild to marked interstitial and perivascular inflammatory cell infiltration of mononuclear inflammatory cells, showing CD3 positivity in immunostaining with rabbit anti-human antibody, and of eosinophilic granulocytes, was found in over one-third of histological endomyocardial specimens of the patients with HCM. In one patient with severe HCM, inflammatory cell infiltration was extensive (figure 2C). In contrast, practically no inflammatory cells were recognised in control myocardial samples. Three previous studies have reported that there is mild chronic inflammatory cell infiltration in the myocardium of patients with HCM, $^{3\ 25\ 26}$ but none of them has included genotyped subjects.

Myocardial NF-KB activity in HCM

A new finding in our study was that immunohistochemical staining for nuclear NF-KB activity was positive in half of the endomyocardial samples of patients with HCM. NF-KB activation was detected particularly in cardiomyocytes, but also in inflammatory and endothelial cells. None of the control

Table 4	The association of circulating cytokine levels with myocardial
maximal L	GE heterogeneity at mid-ventricular level in patients with HCM
(Pearson's	s correlation coefficients)

	LGE in CMRI
hsCRP, mg/l	0.517*
TNFα, pg/ml	0.486*
IL-1β, pg/ml	0.032
IL- 1RA, pg/ml	0.593**
IL-6, pg/ml	0.387
IL-10, pg/ml	0.038

Analysis performed using log 10 transformed values. *p<0.05; **p<0.01.

CMRI, cardiac magnetic resonance imaging; HCM, hypertrophic cardiomyopathy; hsCRP, high-sensitivity C-reactive protein; IL-1β, interleukin 1β; IL-1RA, interleukin 1 receptor antagonist; IL-6, interleukin 6; IL-10, interleukin 10; LGE, late gadolinium enhancement; TNF α , tumour necrosis factor α

myocardial samples showed NF- κ B activation. NF- κ B is a pivotal intracellular mediator of inflammatory response, inducing the proinflammatory cytokine expression.¹¹ NF- κ B activation has been previously implicated in cardiac dysfunction and heart failure.¹¹ NF- κ B activation leads to proinflammatory phenotype including upregulation of TNF α , which activates inflammatory cell invasion and fibroblasts, resulting in perivascular fibrosis formation in the myocardium.²⁷ Our finding of NF- κ B activation in the concept that an inflammatory response may play an integral part in the phenotypic expression of myocardial fibrosis in HCM.

Proinflammatory and anti-inflammatory cytokines in HCM

In this study, both proinflammatory and anti-inflammatory cytokines were raised in patients with HCM. TNF α levels have been reported to be increased in HCM in some,¹² ¹⁴ but not in all, previous studies.¹³ IL-6 levels have been shown to be elevated in HCM in two previous studies.¹³ ¹⁴ Decreased myocardial TNF α expression has been reported after non-surgical septal reduction in patients with obstructive HCM.²⁸ There are, however, no previous studies evaluating systematically the levels of both proinflammatory and anti-inflammatory cytokines in patients with HCM and particularly, there are no studies correlating levels of cytokines with LGE and other LV characteristics in CMRI.

Clinical implications of the association of inflammatory response with myocardial fibrosis in HCM

CMR-derived LGE reflects collagenous scar formation in HCM.²⁹ The notion that myocardial fibrosis is a potentially modifiable inflammatory process opens interesting clinical implications in preventing cardiovascular events in HCM, as myocardial fibrosis is a major determinant of malignant arrhythmias and end-stage systolic heart failure in HCM and consequently, increases the risk of cardiac death.⁵ ³⁰

Strengths and limitations of the study

We demonstrated for the first time, in a well-defined genotyped patient population with HCM, consistent signs of low-grade myocardial inflammation by serological, histopathological and immunohistochemical methods. We also showed that inflammatory response is associated with myocardial fibrosis, documented by histopathological methods and LGE in CMRI. Yet, there are some limitations to this study. First, the patient population with HCM is of limited size. Human studies including patients with genotype-verified diagnosis of HCM, especially those with a single causative mutation, are, however, few and generally small, as shown by recent studies.^{31 32} Second, it is possible that the findings of this study may not be applicable to all patients with HCM with different causative mutations in sarcomeric genes. However, myocardial fibrosis is a common manifestation of the HCM, LGE presenting in 80% of cases.¹ Furthermore, according to current knowledge, no particular clinical HCM phenotype is mutation specific.⁵ Probably, the findings of this study apply to HCM caused by other sarcomeric mutations as well, and confirmation of our findings in large genotyped patient populations is warranted. Third, in our CMRI method, severity and not the extent of LGE was measured. However, the method used has been regarded as scientifically valid and shown to be associated with serum amino-terminal propeptide of type III collagen.¹⁶ Fourth, cadaver myocardial specimens were used as controls and compared with endomyocardial biopsy findings of patients with

- In patients with HCM attributable to an identical diseasecausing mutation in the alpha-tropomyosin gene, myocardial inflammatory cells and enhanced NF-kB activity are commonly present, and circulating cytokine and hs-CRP levels are increased.
- This inflammatory response is associated with histopathological and magnetic resonance derived myocardial fibrosis.
- Our findings suggest that a low grade inflammatory response plays a significant role in the phenotypic expression of myocardial fibrosis in hypertrophic cardiomyopathy.

HCM, since it is unethical to obtain endomyocardial samples from healthy controls. Fifth, there was a variable number of patient samples in different immunohistochemical analyses, because the second best endomyocardial biopsy sample designated for immunohistochemical stainings was not sufficient for all microscopic slides in every patient (see online supplementary information). Nevertheless, this study is, to our knowledge, the first human study to perform both endomyocardial biopsy and CMRI in genotyped subjects with HCM.

CONCLUSIONS

In conclusion, we demonstrate that in patients with HCM attributable to an identical disease-causing mutation in the α -tropomyosin gene, myocardial inflammatory cells and enhanced NF- κ B activity are present and circulating cytokine and hsCRP levels are increased. This inflammatory response is associated with myocardial fibrosis. Our findings suggest that a low-grade inflammatory response plays a significant role in the phenotypic expression of myocardial fibrosis in HCM.

Contributors All authors have contributed to the article by participating in the design (JK, VK, PS, KPe, KPu, AN, ML, IK, SY), performance of genetic studies (JK, PJ, ML), clinical and echocardiographic studies (JK, KPe), magnetic resonance studies (PS, JK), histological and immunohistochemical studies (JK, VK, AN, IK, SY), laboratory measurements (KPu), drafting the manuscript and approving the final manuscript (all authors). All authors have contibuted to the study design, collection of and/or analysis of the data, writing and/or editing the manuscript and approving the manuscript.

 ${\bf Funding}$ This study was supported by the Academy of Finland and the Finnish Heart Research Foundation (grants to JK.).

Competing interests None declared.

Patient consent Obtained.

Ethics approval Provided by the ethics committee of the Kuopio University Hospital.

Provenance and peer review Not commissioned; externally peer reviewed.

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