

# A molecular mechanism improving the contractile state in human myocardial hypertrophy

Oliver Ritter MD<sup>1</sup>, Nico Bottez MS<sup>1</sup>, Natalie Burkard TA<sup>1</sup>, Hagen D Schulte MD<sup>2</sup>, Ludwig Neyses MD<sup>3</sup>

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**BACKGROUND:** Various molecular mechanisms are operative in altering the sarcomeric function of the heart under increased hemodynamic workload. Expression of the atrial isoform (ALC-1) of the essential myosin light chain, a shift from  $\alpha$ -myosin heavy chain (MHC) to  $\beta$ -MHC, increased phosphorylation of the regulatory myosin light chains and increased troponin I (TnI) phosphorylation have been reported to modulate cardiac contractility in rodents.

**METHODS:** To assess a possible contribution of these sarcomeric proteins to cardiac performance in human myocardial hypertrophy, two different forms of cardiac hypertrophy were investigated: 19 patients with hypertrophic obstructive cardiomyopathy (HOCM)

and 13 patients with aortic stenosis (AS) with marked left ventricular hypertrophy and normal systolic function.

**RESULTS:** There was no change in MHC gene expression, regulatory myosin light chain or TnI phosphorylation status in normal heart (NH), HOCM and AS patients. However, patients with hypertrophied myocardium expressed ALC-1 that was not detectable in NH. ALC-1 protein expression correlated positively with the left ventricular ejection fraction. In patients with hypertrophied myocardium, there was a mean ALC-1 protein expression of  $12.7 \pm 3\%$  (range 3.6% to 32%).

**CONCLUSION:** In humans, ALC-1 expression is in vivo a powerful molecular mechanism of the sarcomere to maintain or improve myocardial contractility under increased hemodynamic demands.

**Key Words:** Myocardial hypertrophy; Myosin light chains

Cardiac hypertrophy is an adaptive response that normalizes wall stress and compensates for increased workload. It is accompanied by qualitative and quantitative changes in the expression of sarcomeric proteins. These

alterations have an impact on contractility, intracellular  $Ca^{2+}$  homeostasis and metabolism. Changes in the myosin subunit composition improve contractility by an increase in force generation at a given  $Ca^{2+}$  concentration (1).

<sup>1</sup>Department of Medicine, University of Wuerzburg, Germany; <sup>2</sup>Department of Thoracic Surgery, Heinrich Heine University Duesseldorf, Germany; and <sup>3</sup>University Department of Medicine, Manchester, United Kingdom

Correspondence and reprints: Dr Ludwig Neyses, University Department of Medicine, Manchester Heart Centre, Manchester Royal Infirmary, Oxford Road, Manchester M13 9WL, United Kingdom. Telephone +44-161-276-6631, fax +44-161-276-5152, email [ludwig.neyeses@mhc.cmht.nwest.nhs.uk](mailto:ludwig.neyeses@mhc.cmht.nwest.nhs.uk), cc: [labneyeses@medizin.uni-wuerzburg.de](mailto:labneyeses@medizin.uni-wuerzburg.de)

Phosphorylation of the regulatory myosin light chain (MLC) can also improve cardiac performance (2). Additionally, phosphorylation of troponin I (TnI) leads to a decreased affinity of contractile proteins to  $Ca^{2+}$  (3). In this study, the focus is on the functional significance of myosin heavy chain (MHC) and MLC expression, and the phosphorylation of TnI and the regulatory MLC in the hypertrophied human ventricle. The myosin muscle motor consists of two heavy chains (MHC) and two pairs of light chains (MLC). The MHCs can be divided into the  $\alpha$ - and the  $\beta$ -MHC isoforms (4). During development and under pathophysiological conditions, the expression of the two cardiac isoforms is regulated in a tissue-specific manner. In all mammalian species,  $\alpha$ -MHC is expressed in the atria throughout life. In rodents,  $\alpha$ -MHC is also the predominant isoform in the ventricle of adult hearts.  $\beta$ -MHC is expressed in the embryonic ventricular tissue. In rodents, there is a switch from  $\alpha$ -MHC to  $\beta$ -MHC under hemodynamic stress (5).  $\beta$ -MHC has reduced ATPase activity and increased economy of isometric force development (6). In humans,  $\beta$ -MHC is the predominant isoform in the adult heart (7). Whether there is a switch under increased workload in the human ventricle is still a matter of debate and is investigated in this study.

MLCs are classified into essential MLC (MLC-1) and regulatory (phosphorylatable) MLC (MLC-2) (8,9). (Table 1) In the human heart, five MLC isoforms exist that are expressed in a tissue-specific manner: there are three regulatory MLCs, namely two ventricular specific (VLC-2 and VLC-2\*) and an atrial specific (ALC-2) (10). Two essential light chains (MLC-1) have been described: ALC-1 in the atrium and VLC-1 in the ventricle (Figure 1). Human embryos express large amounts of ALC-1 in the whole heart and in skeletal muscle. For this reason the ALC-1 was designated as the embryonic MLC (MLC-1emb) (10). A positive correlation between pressure overload in adult patients with valvular heart disease and left ventricular ALC-1 expression exists. Normalization of the

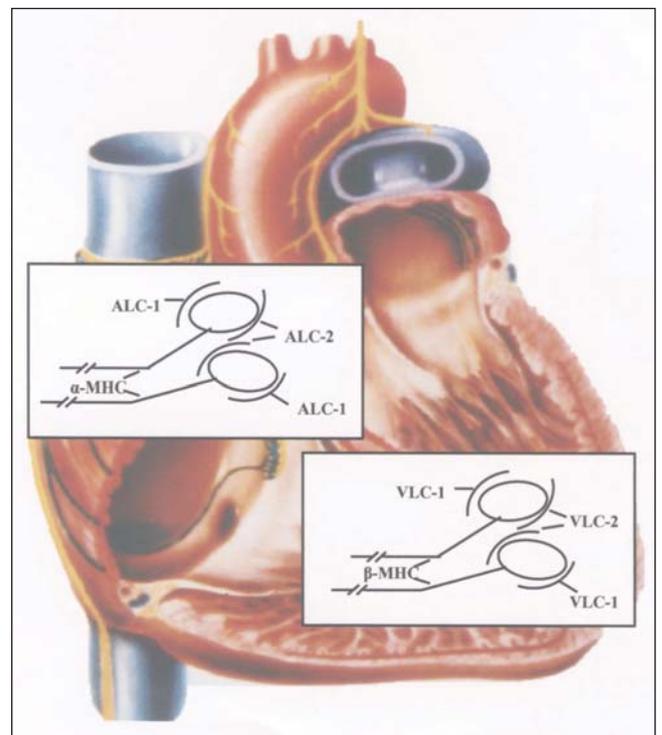
**Table 1**  
**Nomenclature of myosin light chains and heavy chains**

MHC	Myosin heavy chain
MLC	Myosin light chain
MLC-1	Essential myosin light chain
MLC-2	Phosphorylatable (regulatory) myosin light chain
ALC-1	Essential myosin light chain, atrial isoform
ALC-2	Phosphorylatable (regulatory) myosin light chain, atrial isoform
VLC-1	Essential myosin light chain, ventricular isoform
VLC-2	Phosphorylatable (regulatory) myosin light chain, ventricular isoform
VLC-2*	Phosphorylatable (regulatory) myosin light chain, additional ventricular isoform

hemodynamic state by valve replacement significantly reduced hypertrophy and ALC-1 expression (11).

Phosphorylation of the MLC-2 has been shown to increase tension development and  $Ca^{2+}$  sensitivity (12). Recently, it has been demonstrated that there is a spatial gradient across the heart in MLC phosphorylation altering tension production in different wall sections (2).

The cardiac troponin complex comprises the inhibitory (TnI), the  $Ca^{2+}$ -binding (TnC) and the tropomyosin-binding (TnT) subunits (13). Phosphorylation of TnI by Protein Kinase A (PKA), which results from  $\beta$ -adrenergic stimulation, leads to a decrease in the sensitivity of the contractile proteins to  $Ca^{2+}$ , which is accompanied by an increased crossbridge cycling rate. These results indicate that PKA-induced phosphorylation of TnI increases the power output-generating capacity of cardiac myocytes (14). In terminally failing hearts, TnI phosphorylation was increased compared with normal heart (NH). This phosphorylation difference could underlie the reported decrease of myofibrillar  $Ca^{2+}$  sensitivity of failing myocardium (15). However, there are no data on TnI phosphorylation in hypertrophied, non-failing human myocardium. Therefore, in the present study TnI phosphorylation status of hypertrophied human myocardium was investigated.



**Figure 1** Scheme of myosin heavy chain (MHC) and myosin light chain (MLC) distribution in normal human heart. On the atrial level there is  $\alpha$ -MHC and the atrial MLC isoforms ALC-1 and ALC-2. On the ventricular level only  $\beta$ -MHC and the ventricular MLC isoforms VLC-1 and VLC-2 can be detected. Each myosin molecule consists of two MHCs, and on the head of each MHC there is one molecule of MLC-1 and MLC-2

## MATERIAL AND METHODS

## Patient population

Tissue from the left ventricular outflow tract was obtained from 13 patients undergoing aortic valve replacement for aortic stenosis (AS) and from 19 unrelated patients with hypertrophic obstructive cardiomyopathy (HOCM) undergoing transaortic subvalvular myotomy-myectomy (16) for severe obstruction of the left ventricular outflow tract. The clinical evaluation consisted of preoperative physical examination, electrocardiography (ECG), echocardiography and cardiac catheterization. All the patients in this study had a normal ejection fractions (EF). Atrial tissue was taken from the auricle of a patient undergoing mitral valve replacement. Tissue from the interventricular septum was taken from six healthy donor hearts that could not be transplanted for technical reasons. Tissue samples were frozen in liquid nitrogen immediately after resection. Informed consent was obtained from all patients.

## Tissue preparation

All tissue samples were immediately chemically skinned or frozen in liquid nitrogen after resection. All biochemical experiments were performed with demembrated multicellular heart fibres (skinned fibres) prepared as described previously (17) or with frozen tissue samples. For the preparation of the skinned fibres, ventricular fibre bundles (1 mm thick, between 2 and 5 mm long) were incubated in a solution containing 20 mM imidazole, 10 mM  $\text{NaN}_3$ , 5 mM ATP, 5 mM  $\text{MgCl}_2$ , 4 mM EGTA, 2 mM dithioerythritol, 50% glycerol and 1% Triton X-100, pH 7, at 4°C for 20 h. Subsequently the fibres were transferred into the same solution except Triton X-100 and stored at -20°C. Tissue specimen for analysis of phosphorylation levels were homogenized in ice-cold trichloroacetate before further experimentation. All tissue samples immediately frozen after resection were stored at -80°C.

## Biochemical analysis

Samples for high-resolution protein gel electrophoresis for MHC analysis were prepared as follows. Briefly, samples were homogenized in low-salt buffer (20 mmol/L KCl, 2 mmol/L  $\text{KH}_2\text{PO}_4$ , 1 mmol/L EGTA, pH 6.8, 1 mmol/L phenylmethylsulphonylchloride and 100  $\mu\text{L}$  *N,N*-dimethyl formamide). The samples were then centrifuged at 900 *g* for 10 min at 4°C. Laemmli's buffer was added to each sample. The preparation and composition of the gel were as described earlier (10). Gel samples (0.25 to 1  $\mu\text{g}$ ) were loaded in a 3  $\mu\text{L}$  volume onto 15-well gels. The stacking and separating gels (0.75 mm thick) consisted of 4% and 8% acrylamide, respectively; the stacking gels included 5% glycerol. The gels were run at a constant voltage of 200 V for 30 h. The gels were fixed and silver-stained. A gel documentation system (Bio-Rad, Germany) was used to scan the stained gels.

MLCs were analyzed by a high-resolution two-dimensional gel electrophoresis two-dimensional polyacrylamide gel

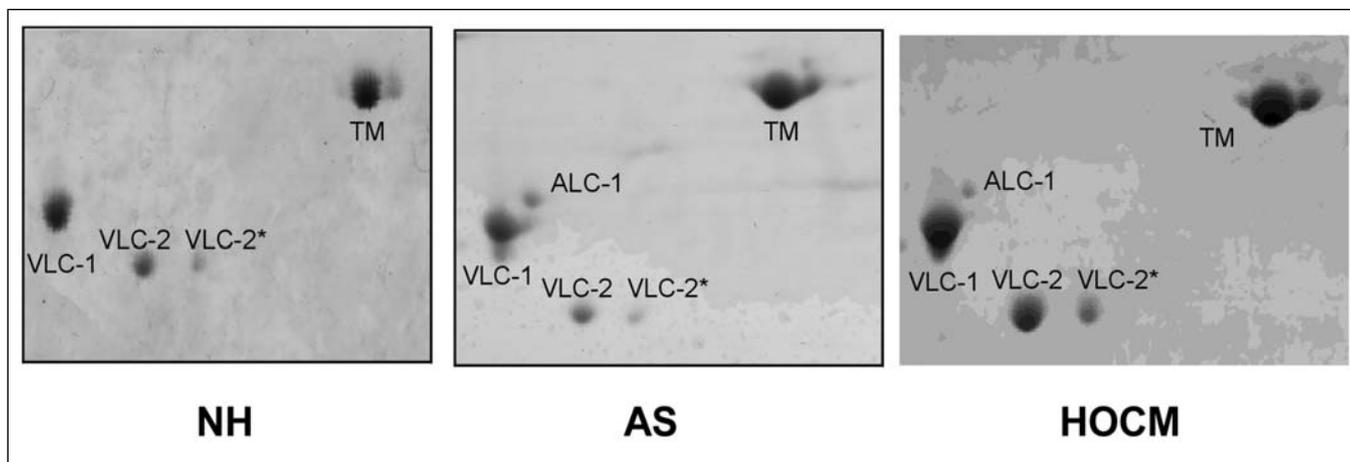
Table 2

Echocardiographic, hemodynamic and biochemical data of patients with hypertrophic obstructive cardiomyopathy (HOCM) or aortic stenosis (AS)

Diagnosis	Age (years)	Sex	IVS (mm)	EF (%)	LVEDP (mmHg)	ALC-1%
HOCM	56	f	17	62	12	1
HOCM	53	m	14	63	21	3
HOCM	38	m	28	66	16	4
HOCM	34	f	24	68	10	2
HOCM	56	f	17	69	6	8
HOCM	46	f	20	73	13	5
HOCM	60	f	24	72	17	14
HOCM	80	f	20	88	15	2
HOCM	63	f	20	76	15	10
HOCM	53	m	31	77	19	6
HOCM	48	m	24	76	1	11
HOCM	43	m	24	81	16	11
HOCM	64	f	22	81	20	13
HOCM	47	m	13	79	11	16
HOCM	46	m	15	72	1	20
HOCM	42	f	17	85	10	20
HOCM	62	f	16	82	20	22
HOCM	70	m	22	86	6	25
HOCM	56	f	17	80	17	32
AS	70	m	22	65	6	1
AS	56	f	17	64	9	2
AS	46	f	20	70	13	2
AS	60	f	24	74	17	2
AS	80	f	20	78	15	5
AS	43	m	15	74	24	6
AS	42	f	17	74	10	7
AS	68	f	16	78	20	9
AS	51	f	17	82	16	6
AS	72	m	22	80	6	16
AS	56	f	17	78	16	18
AS	46	f	20	76	13	24
AS	55	m	33	80	9	23

IVS Diameter of the interventricular septum in mm; LVEDP Left ventricular end-diastolic pressure; EF Ejection fraction; ALC-1% Expression of the atrial isoform *m* (ALC-1) of essential myosin light chain (MLC-1) as a percentage of total MLC-1 (ie, ALC-1 + ventricular MLC isoform = 100%); f female; m male

electrophoresis [2D PAGE] technique. Isoelectric focusing (first dimension) was performed in glass capillaries (12.5 cm long, 1 mm inner diameter) using the pH gradient 4.5 to 5.5 (Pharmalytes, Pharmacia, Sweden). The gels were run overnight at 600 V constant load for the first dimensional separation. The second dimension was a sodium dodecylsulphate electrophoresis, using slab gels 10.5×9.5 cm, 1 mm thick. The gels were stained in Coomassie blue and the MLCs were evaluated by computer-assisted scanner densitometry (ScanPack, Biometra, Germany).



**Figure 2)** Original photographs of myosin light chains (MHC) in the myocardium of normal heart (NH) and in patients with aortic stenosis (AS) or hypertrophic obstructive cardiomyopathy (HOCM). All fibres contained VLC-1, VLC-2 and VLC-2\*, but patients with hypertrophied myocardium contained ALC-1 additionally. ALC-1 Atrial essential light chain; TM Tropomyosin; VLC-1 Ventricular essential light chain; VLC-2 and VLC-2\* Isoforms of the ventricular regulatory light chain. No phosphorylation of the regulatory light chains was detectable

TnI phosphorylation status was analyzed by 2D PAGE technique according to the above protocol. However, the first dimension (isoelectric focusing) was a nonequilibrium gel with inversed current. TnI phosphorylation level was expressed as a percentage of total TnI, ie, TnI unphosphorylated + TnI phosphorylated = 100%.

#### Statistical analysis

All statistical analyses were performed using commercially available statistic programs (Enzfitter, Epistat, Germany) on Apple Macintosh or on IBM compatible PC. Values are expressed as mean  $\pm$  SEM. Significance analysis was performed using the Student's *t* test for unpaired values. Regression analysis was performed by the least-squares method.

## RESULTS

#### Clinical data

In clinical terms there were homogeneous populations of patients with HOCM and with AS. All patients had left ventricular myocardial hypertrophy. The extent of hypertrophy of the interventricular septum was similar in both patient groups. There were no significant differences in hemodynamic data in the patient groups (Table 2).

#### MLC analysis

The MLC isoforms in frozen tissue and skinned fibres were analyzed according to their isoelectric point and molecular weight by 2D PAGE. The atrial isoform of the essential light chain (ALC-1) had a higher molecular weight and was more acidic than the ventricular isoform (VLC-1). Figure 1 shows three original gels of tissue preparations of the left ventricle from patients with HOCM, AS and NH. All the fibres contained VLC-1 and the two ventricular isoforms of the regulatory light chains, VLC-2 and VLC-2\*, but the preparation of

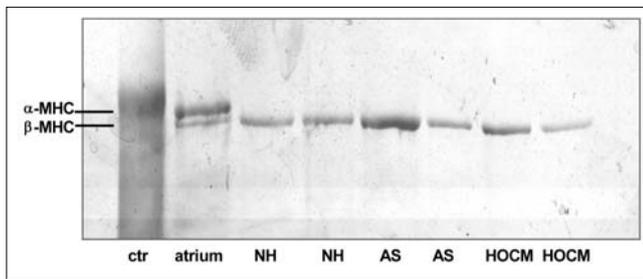
normal myocardium showed no ALC-1 whereas diseased myocardium had ALC-1 (11.4% of total essential light chains [MLC-1]). The mean ALC-1 expression was  $12.7 \pm 1\%$ . Those patients who expressed ALC-1 had very different amounts ranging from 3.6% to 32.2% ALC-1 (expressed as a percentage of total MLC-1, ie, ALC-1 + VLC-1 = 100% MLC-1) (Table 2).

The ratio between essential light chains (MLC-1) and regulatory light chains (MLC-2) showed no significant difference between patients with and without ALC-1. In the NH, the MLC-1/MLC-2 ratio was  $1.1 \pm 0.02\%$ . In patients with hypertrophied myocardium, an MLC-1/MLC-2 ratio of  $1.26 \pm 0.78$  and a ratio of  $1.21 \pm 0.48$  in patients without ALC-1 ( $n=6$ ), respectively was observed. The atrial isoform of the regulatory light chain (ALC-2) was never observed in our tissue. No tissue preparation-dependent differences in results were found. In all individuals investigated here, the content of ALC-1 in skinned fibres and in frozen tissue was identical, indicating total incorporation of ALC-1 protein into the sarcomere.

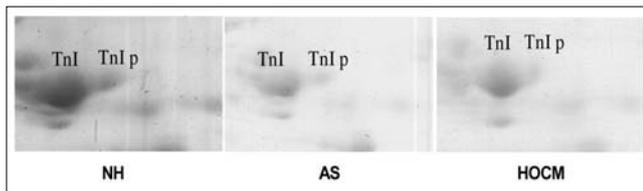
Phosphorylated MLC-2 promotes contractility and thus counteracts the attenuating effects of phosphorylated TnI. Therefore, MLC-2 phosphorylation in 2D PAGEs was investigated. Phosphorylated proteins are more acidic, resulting in a shift of protein spots to the cathode. However, virtually no MLC-2 phosphorylation in hypertrophied or normal human myocardium was found.

#### MHC analysis

The relative proportions of the two forms of the MHC have been shown to be affected by a wide variety of pathological and physiological stimuli. Hearts that express the faster  $\alpha$ -MHC produce more power than those expressing the slower  $\beta$ -MHC, leading to the hypothesis that MHC isoforms play a major role in the determination of cardiac contractility. In the present study, therefore, the MHC



**Figure 3)** Original photographs of myosin heavy chains (MHC) in the myocardium of normal heart (NH) and in patients with aortic stenosis (AS) or with hypertrophic obstructive cardiomyopathy (HOCM). Ctr Rabbit skeletal muscle myosin. Human atrial myocardium contained  $\alpha$ -MHC and  $\beta$ -MHC. Human ventricular myocardium from patients with NH, AS or HOCM contained only  $\beta$ -MHC



**Figure 4)** Original photographs of troponin I (TnI) in the myocardium of normal heart (NH) and in patients with aortic stenosis (AS) or hypertrophic obstructive cardiomyopathy (HOCM). TnI p indicates phosphorylated TnI. There are no differences in the amount of phosphorylated TnI in the three patient groups

protein isoform content of our samples was determined. No  $\alpha$ -MHC was detectable in six NH and in 32 nonfailing but hypertrophied human hearts (Figure 3).

#### TnI phosphorylation

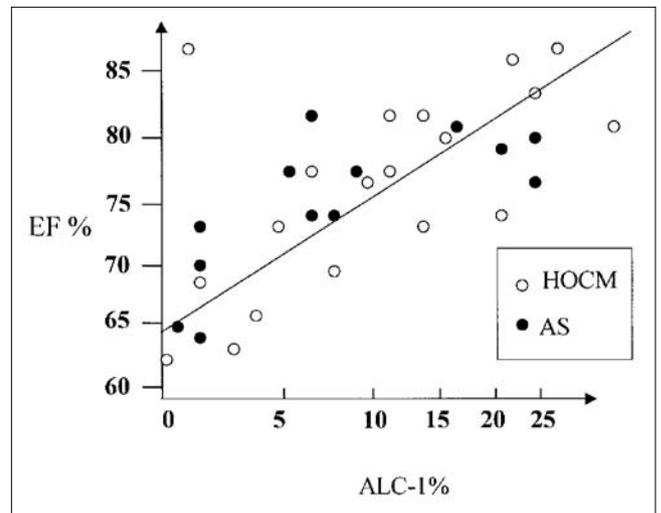
TnI phosphorylation contributes to the impaired contractile response of the myocardium to hemodynamic stress. Using nonequilibrium 2D PAGE, TnI phosphorylation was compared in patients with NH, HOCM and AS. However, not significant change was found in the phosphorylation level of TnI in the three patient groups. In NH, phosphorylated TnI was  $12 \pm 3\%$  ( $n=6$ ), in AS it was  $10 \pm 4\%$  ( $n=13$ ; not significant) and in HOCM it was  $13 \pm 4\%$  ( $n=19$ ; not significant) (Figure 4).

#### Correlation between ALC-1 protein expression and EF

EF showed a mean value of  $73 \pm 7\%$ . Notably, there was a statistically significant positive correlation between EF and ALC-1 content of the myocardium. This is compatible with a role of the ALC-1 in contractility in these disease states (Figure 5). The high correlation coefficient of 0.76 ( $P < 0.01$ , F-test) further supports this hypothesis.

### DISCUSSION

Gene expression in the overloaded rodent heart has been studied extensively to highlight the molecular response of the sarcomere. It is generally accepted that, in terms of contractile proteins, a shift of MHC isoforms is part of the response (18). Work from several groups has shown that in



**Figure 5)** Relation between the expression of atrial isoform of the essential myosin light chain (ALC-1) and ejection fraction (EF). Each circle indicates one patient. The ALC-1 content is shown as a percentage of total myosin light chain-1 (ALC-1 + VLC-1). AS Aortic stenosis; HOCM Hypertrophic obstructive cardiomyopathy

the rodent heart, the MHC phenotype switches from the  $\beta$ -MHC isoform to the  $\alpha$ -MHC isoform during maturation and vice versa under increased hemodynamic demands (19-24). This conversion of the phenotype is associated with alterations in the contractile and energetic behaviour of the contractile apparatus. Expression of  $\alpha$ -MHC leads to a higher maximal shortening velocity, paralleled by a decreased economy in tension development (25). In turn, high amounts of  $\beta$ -MHC decrease maximal shortening velocity and increase economy of contraction (26), which is believed to improve the function of the overloaded cardiomyocyte. These findings raised the question of whether this mechanism might also be operative in the human ventricle. Immunohistochemical studies (27-29) and peptide maps (28) revealed a tissue-specific expression of both MHC isoforms, with a predominant expression of the  $\alpha$ -MHC in the atrium and the  $\beta$ -MHC in the ventricle. However, quantification of the relative distribution of both isoforms in the ventricle has proved difficult.

In two previous studies,  $\alpha$ -MHC mRNA was detected in the normal human heart. A downregulation of  $\alpha$ -MHC mRNA and an upregulation of  $\beta$ -MHC mRNA was reported in the failing and hypertrophied human heart in analogy to the rodent heart (29,30). However, these studies did not address the problem of whether this switch at the RNA level is reflected in MHC protein content.

Therefore, we here quantified the expression of  $\alpha$ -MHC protein in the normal and hypertrophied human ventricle using high-resolution gel electrophoresis. We found no traces of  $\alpha$ -MHC protein expression in the normal or in the hypertrophied ventricular human tissue. The MHC switch for regulation of power output is not available in humans. What might be the human correlate? We recently reported that the human ventricular cardiomyocyte regulates its

power output by changing essential MLC gene expression rather than MHC gene expression (31,32). These observations imposed the question of whether an increase of ALC-1 expression is also beneficial for the performance of the human ventricle *in vivo*.

Here, we evaluated the effects of ALC-1 on cardiac performance of the human heart *in vivo* in a clinical homogeneous population of patients with HOCM and AS without systolic dysfunction. We observed a close and statistically significant correlation between essential MLCs and contractility of the human ventricle *in vivo*. Increased ALC-1 protein expression augmented cardiac contractility in the human ventricle.

Myocardial performance could be influenced not only by isoform shifting of MHCs and MLCs but also by phosphorylation of sarcomeric proteins. To determine the influence of phosphorylation of sarcomeric proteins, we investigated the phosphorylation levels of two key proteins of myocardial contractility: VLC-2 and TnI in normal and hypertrophied human heart. In our patients, we observed no phosphorylation of VLC-2 in the human ventricle and no significant changes in TnI phosphorylation levels.

Expression of ALC-1 in diseased hearts raises the question of whether there is a compensatory downregulation of the ventricular isoform. According to the generally accepted myosin model, the ratio between MLC-1 and MLC-2 is 1 (ie, each MHC monomer is associated with one essential and one regulatory light chain) (33). In our patients we did not see a difference in the MLC-1/MLC-2 ratio in any of the groups, the ratio being about 1 in each individual. Downregulation of the ventricular form of the MLC-2 is to be expected in order to maintain the normal MLC-1/MLC-2 ratio. This observation is in line with previous studies on ALC-1 expression in the diseased human myocardium. The hypertrophied right ventricle of children with congenital heart diseases has been demonstrated to express large amounts of ALC-1 up to adulthood (34). Similarly, the dilated left ventricle of patients with ischemic or dilative cardiomyopathy re-expressed ALC-1 (35). In adult patients with valvular heart disease, a positive correlation between systolic wall stress and left ventricular ALC-1 expression has been reported. Normalization of the hemodynamic state by valve replacement significantly reduced hypertrophy and ALC-1 expression (11). Recently, some studies investigated the functional consequences of re-expression of essential MLCs on the molecular level. Myosin crossbridges with high amounts of ALC-1 had a higher detachment rate and rate of force generation than myosin with VLC-1 (31). These results showed that ALC-1 increased cycling kinetics of cardiac crossbridges in demembrated multicellular heart fibres, thus regulating the contractile state of the heart. It was also shown that there is a positive correlation between the Ca<sup>2+</sup> sensitivity of isometric force development of skinned human heart fibres and ALC-1 content (32). The inotropic effects of ALC-1 expression could be confirmed *in vivo* by a recent mouse model overexpressing ALC-1 (36).

The molecular mechanism explaining the effects of essential MLC on crossbridges in the heart may reside in the mode of interaction of the essential MLC and actin. It has been demonstrated that the amino terminal domain of MLC-1 isoforms interacts with the carboxyl terminus of actin. This interaction could be of functional importance because the inhibition of this interaction increased force production and shortening velocity of human heart fibres (37). Interestingly, the primary structure of the most amino terminal end of ALC-1 and VLC-1 is different: ALC-1 contains two charges less compared with VLC-1, suggesting a different binding affinity between ALC-1 or VLC-1 and actin. In fact, binding of ALC-1 to actin was found to be weaker than binding of VLC-1 to actin (38). Weakening of the VLC-1-actin interaction by peptide competition accelerated crossbridge cycling kinetics, increased tension production and increased Ca<sup>2+</sup> sensitivity (39). These results are analogous to studies on muscle fibres with an increased content of ALC-1 (31,32,39).

To our knowledge, this is the first study investigating MHC and MLC composition in the following subsets of patients with cardiac hypertrophy and normal systolic function: HOCM and severe AS. In view of the relative rarity of these patients (particularly HOCM with indication for operation) the numbers are large. It is noteworthy that large pieces of tissue were used for measurements as opposed to percutaneous biopsies with the inherent problem of tissue heterogeneity (fibrosis). These two factors may also explain the relatively low variability of our results compared with other studies in human tissue. We demonstrate improved intrinsic contractile function in the human ventricle *in vivo*, which significantly correlated with enhanced ALC-1 protein expression, excluding an MHC type of regulation. Expression of ALC-1 in the hypertrophied human ventricle therefore may be an adaptational molecular mechanism to improve the inotropic state of the human heart. ALC-1 expression turned out to be a powerful compensatory mechanism to enhance power output of the human heart.

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