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Familial Hypertrophic Cardiomyopathy – is the Frank-Starling Law kaput?

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Building on work by Otto Frank in the late 19th century, Ernest Starling described in the early 20th century the ability of the heart to change its force of contraction and therefore stroke volume in response to changes in venous return, which has been called the Frank-Starling law of the heart in honor of these two physiologists. The Frank-Starling effect enables the heart to match cardiac output to venous return on a beat-to-beat basis.¹ A major mechanism responsible for the Frank-Starling effect and hence the beat-to-beat autoregulation of cardiac output is the *sarcomere length-dependent activation* of force development.²

In this issue of *Circulation Research*, Sequeira and coworkers³ investigate both the Ca and length-dependent regulation of force development in cardiomyocytes harvested from patients with Hypertrophic Cardiomyopathy (HCM), a familial disorder commonly associated with mutations in genes encoding sarcomeric proteins.⁴ Consistent with previous animal and human studies⁵, myofilament Ca sensitivity was increased in all 38 HCM patient samples compared to 12 non-failing donor hearts. Notable, the authors observed that length-dependent activation was impaired, which, unlike the Ca sensitivity increase, appeared to be a primary effect of the mutant sarcomeric proteins, because 1), it could not be rescued by increasing protein kinase A (PKA) phosphorylation, and 2), was normalized when mutant troponin proteins were replaced with wild-type protein. These new results suggest that impaired length dependent activation and hence a defective Frank-Starling effect importantly contributes to the pathogenesis of HCM.

Reduced maximal Ca-activated force and increased Ca sensitivity are central findings in human HCM

HCM is a familial disorder commonly associated with mutations in genes encoding sarcomeric proteins. Mutation carriers develop progressive heart hypertrophy and fibrosis,

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DISCLOSURES

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and are at a high risk for sudden death due to ventricular arrhythmia. More than 1000 mutations in >11 genes have been identified that can lead to this disease. The current report³ together with two previous smaller human studies^{6, 7} indicates that *regardless of the disease-causing mutation*, human HCM cardiac muscle exhibits *reduced* maximal Ca-activated force and *increased* Ca sensitivity of muscle activation. This result raises the possibility that HCM mutations trigger a common cascade of molecular events that result in a similar phenotype (Figure 1).

Based largely on in vitro studies and animal models, myofilament Ca sensitization had previously been associated with HCM (reviewed in^{8, 9}). The study by Sequeira et al.³ suggests that myofilament Ca sensitization indeed is a universal finding in human HCM. However, the authors find that Ca sensitization is often not a primary property conferred by the mutant protein itself. Rather, the increased Ca sensitivity was largely due to an apparent hypophosphorylation of PKA targets in the myofilaments: Although all HCM groups had higher myofilament Ca sensitivity at baseline, in tissue from mutation carriers in myosin binding protein C (MyBPC), troponin I (TnI), tropomyosin and from HCM patients without sarcomeric mutations, Ca sensitivity was restored to normal levels by treatment with PKA. The two exceptions were one patient with a mutation in TnT and another patient with a MHC mutation, where the Ca sensitivity increase appeared to be a property directly conferred by the mutant protein. As with any patient study, differences in drug therapy between the HCM and the normal donor group may have contributed to the reduced PKA phosphorylation. Nevertheless, this result suggests that the increased Ca sensitivity of human HCM muscle is mostly the result of post-translational modifications, and potentially helps to maintain systolic function despite the reduction in maximal Ca-activated force caused by the HCM mutation (Figure 1).⁷ Regardless of the underlying cause, the Ca sensitivity increase is likely of major importance for HCM pathophysiology, with experimental evidence indicating that regardless of how myofilaments are Ca sensitized, an increase in myofilament Ca sensitivity can contribute to diastolic dysfunction,¹⁰ focal energy deficits¹¹ and arrhythmogenesis (Figure 1).¹²

Impaired length-dependent activation – a trigger of HCM pathogenesis?

The authors discovered a new common finding in human HCM muscle – impaired length-dependent activation, which, in contrast to increased Ca sensitivity, remained after PKA treatment for all sarcomeric *missense* mutations studied. Furthermore, replacement of mutant with wild-type troponin corrected the impaired length dependence in the two mutations that could be studied. Hence, *impaired length-dependent activation* may be a direct effect of many mutations that cause HCM in humans (Figure 1). What is the significance of this finding for the pathogenesis of HCM? Since a functional length-dependent activation is responsible for the Frank-Starling effect, it is intriguing to speculate that HCM mutations generate an intermittent inability to match cardiac ejection to dynamic changes in ventricular filling volume (i.e., during the respiratory cycle or during exercise). The resulting increase in wall stress due to intermittent impaired ventricular emptying could trigger neuroendocrine activation and a hypertrophic gene program analogous to acquired heart disease with increased wall stress and an impaired Frank-Starling effect such as post-MI cardiomyopathy. While not tested directly in the report by Sequeira, activation of neuroendocrine signaling could be the main culprit for the decrease in PKA phosphorylation of myofilament target proteins found universally in HCM patients (Figure 1). In addition, some of missense HCM mutation (i.e., in TnT or TnI) may directly interfere with PKA's ability to phosphorylate TnI (Figure 1, dashed arrow *b*). PKA phosphorylation of TnI not only regulates myofilament Ca sensitivity, but also is important for physiological length-dependent activation of cardiac muscle.¹³ Hence, PKA hypophosphorylation can independently contribute to impaired length-dependent activation, as was demonstrated by

the authors for two groups of HCM patients (HCM due to truncation mutations of MyBPC and genotype-negative HCM). In those patients, this can generate a vicious cycle that could be central to the pathogenesis in HCM, as shown in Figure 1.

In addition to the impaired length-dependent activation, which is measured experimentally by the increase in Ca sensitivity (pCa_{50}) induced by increased sarcomere length (Sequeira et al, Figure 2), all sarcomeric *missense* mutation (but not MyBPC truncation mutants and genotype-negative HCM) also impaired the length-dependent increased in maximum force (Sequeira et al, Table 2). The latter is measured at fully-activating $[Ca]$ and reflects the intrinsic force generating ability of the muscle. While there can be many reasons for the reduced length dependence of maximum force (e.g., loss of myofibrils), the net result is an even further compromised length-tension relationship and hence impaired Frank Starling effect.

Open questions

A number of questions remain. Given that genetic background might affect the functional effect of the mutations and many of the individuals studied by Sequeira et al. were family members, the results of the study may not be quite generalizable to the general HCM population. By necessity, all human myocardial samples came from patients with significant cardiac hypertrophy and/or heart failure. Hence, it is still possible that the altered length-dependent activation is the consequence of pathological remodeling of the sarcomeric apparatus in the diseased myocardium. Nevertheless, the rescue with wild-type troponin protein replacement provides compelling evidence that this is not the case, at least for two specific mutations. Furthermore, passive viscoelastic properties of human HCM cardiomyocytes are similar to those of cardiomyocytes from healthy donor hearts,⁷ further supporting the hypothesis that altered Ca and length dependent regulation of muscle activation are central to the pathophysiology of human HCM (Figure 1). Not directly investigated in the current study, but likely also important for HCM pathophysiology is the inefficient ATP utilization (increased “tension cost”) caused by many HCM mutations.¹⁴ It remains to be seen which of these myofilament properties is the most significant driver of the disease. However, altered myofilament properties affect *every* myocyte in the heart starting at *birth* and while some heterogeneity is expected, there are significant questions remaining. For example, what is the reason for the incomplete penetrance of the disease (age dependent, but possibly as low as 41%¹⁵)? What is responsible for the focal nature of hypertrophy and fibrosis in HCM hearts? This clearly shows that the presence of a HCM mutation does not inevitably lead to disease, and even in an affected patient some regions of the heart remain normal. Such a phenotype may imply a “second hit” that is necessary to initiate the disease and remains to be identified. A potential culprit is the focal energy deprivation that occurs in HCM mouse models during stress,¹¹ which can directly contribute to reentry arrhythmias,¹¹ but also has the potential to drive heterogeneous hypertrophy and remodeling.¹⁶ The next frontier will be to test whether therapeutic interventions aimed at normalizing Ca sensitivity, length-dependent activation or inefficient ATP utilization are beneficial in HCM.

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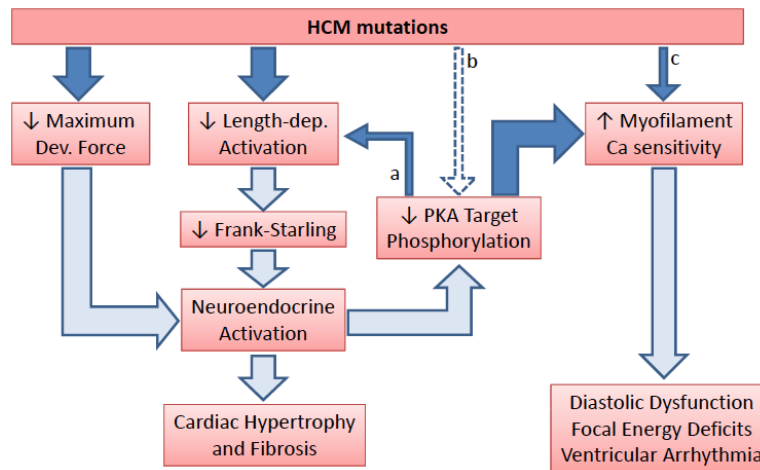


Figure 1. Pathophysiological framework of HCM

Solid arrows indicate experimental findings reported by Sequeira and coworkers. Open arrows indicate the consequences based on experimental evidence from other studies. (a) Impaired length-dependent activation can also be caused by PKA hypophosphorylation (i.e., of TnI). (b) HCM mutations may directly interfere with phosphorylation of myofilament PKA targets. (c) Only a subset of HCM mutations directly alter myofilament Ca sensitivity.