CARDIOVASCULAR, PULMONARY, AND RENAL PATHOLOGY

Myofilament Remodeling and Function Is More Impaired in Peripartum Cardiomyopathy Compared with Dilated Cardiomyopathy and Ischemic Heart Disease

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Peripartum cardiomyopathy (PPCM) and dilated cardiomyopathy (DCM) show similarities in clinical presentation. However, although DCM patients do not recover and slowly deteriorate further, PPCM patients show either a fast cardiac deterioration or complete recovery. The aim of this study was to assess if underlying cellular changes can explain the clinical similarities and differences in the two diseases. We, therefore, assessed sarcomeric protein expression, modification, titin isoform shift, and contractile behavior of cardiomyocytes in heart tissue of PPCM and DCM patients and compared these with nonfailing controls. Heart samples from ischemic heart disease (ISHD) patients served as heart failure control samples. Passive force was only increased in PPCM samples compared with controls, whereas PPCM, DCM, and ISHD samples all showed increased myofilament Ca²⁺ sensitivity. Length-dependent activation was significantly impaired in PPCM compared with controls, no impairment was observed in ISHD samples, and DCM samples showed an intermediate response. Contractile impairments were caused by impaired protein kinase A (PKA)—mediated phosphorylation because exogenous PKA restored all parameters to control levels. Although DCM samples showed reexpression of EH-myomesin, an isoform usually only expressed in the heart before birth, PPCM and ISHD did not. The lack of EH-myomesin, combined with low PKA-mediated phosphorylation of myofilament proteins and increased compliant titin isoform, may explain the increase in passive force and blunted length-dependent activation of myofilaments in PPCM samples. (Am J Pathol 2017, 187: 2645–2658; https://doi.org/10.1016/j.ajpath.2017.08.022)

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Dilated cardiomyopathy (DCM) and peripartum cardiomyopathy (PPCM) are distinct forms of cardiac disease that share aspects of clinical presentation, such as dilated ventricle(s), reduced systolic function, and cardiac rhythm disorders.\(^7\) PPCM is a cardiac disease that presents itself during the last month of pregnancy, during delivery, or within the first 5 months postpartum.\(^2\) A genetic cause for both DCM and PPCM has been demonstrated before.\(^3\)–\(^6\) Evidence has accumulated showing that approximately 15% to 20% of PPCM patients carry cardiomyopathy-causing mutations, mainly in \(TTN, MYH7,\) and \(SCN5A,\) genes in which mutations are associated with DCM as well.\(^7\) PPCM patients are often young and show fast cardiac deterioration after disease onset, leading to the need for heart transplantation or death.\(^7\)–\(^10\) In contrast, DCM patients develop cardiac disease at an older age, with a milder disease progression compared with PPCM patients.\(^11\) Although PPCM patients can show fast cardiac deterioration, they often can be stabilized or even show complete recovery on treatment.\(^7\)–\(^14\) In a cohort of 182 PPCM patients in the United States, 27% showed full recovery, whereas 25% experienced at least one major adverse event (defined as death or complications that were life threatening or lead to long-term morbidity).\(^15\) Of the patients who survived a major adverse event, but did not receive a cardiac transplantation, 7 (32%) experienced residual brain damage. Of the cohort, 13 patients died within 8 years, of whom 5 died suddenly, and 11 underwent cardiac transplantation within 2 years after diagnosis. Cardiopulmonary arrest occurred in six patients either during delivery or within the first 6 days after delivery. Severe pulmonary edema requiring intensive care occurred in 16 patients within 1 week after delivery.\(^15\) This cohort clearly illustrates the fast cardiac deterioration in PPCM patients. A German cohort of 115 PPCM patients reported an improvement rate of 85%, a full recovery rate of 47%, a failure to recover rate of 15%, and a death rate of 2%,\(^15\) demonstrating the high recovery rate from PPCM.\(^15\) DCM patients may be stabilized, but recovery after treatment is unlikely.\(^6\)\(^,\)\(^14\)

Interestingly, DCM patients who become pregnant are more likely to experience adverse cardiac events, such as heart failure, sustained arrhythmia, or stroke, earlier in pregnancy than PPCM patients. This is likely caused by an increased hemodynamic load on an already troubled heart. In contrast, increased cardiac strain does not appear to be a disease trigger for PPCM because the highest incidence of disease onset is not at the time of the largest increase in hemodynamic load (approximately 24 weeks), but rather in the first month postpartum.\(^16\) In addition, it has been shown that the hormone prolactin plays a key role in PPCM pathogenesis. High levels of oxidative stress induce cleavage of the breastfeeding hormone prolactin into a 16-kDa fragment that has detrimental effects on cardiac function.\(^7\)\(^,\)\(^17\)\(^,\)\(^18\) The peak of prolactin levels coincides with the peak incidence of disease onset in PPCM. Although prolactin levels are low in nonpregnant and nonnursing individuals, it is unlikely that prolactin plays a role in DCM. A recent study suggests detrimental effects of catecholamine treatment, specifically with the \(\beta1\)-adrenergic receptor agonist dobutamine, in PPCM patients.\(^13\)\(^,\)\(^20\) In fact, experimental data showed that the \(\beta1\)-receptor agonist isoproterenol impaired fatty acid and glucose uptake.\(^20\) The resulting energy depletion increased production of reactive oxygen species and led to cardiomyocyte death and progression of heart failure in STAT3 knockout mice, which could be attenuated with the \(\beta\)-blocker metoprolol.\(^20\) On the basis of differences in disease onset and progression, we hypothesize that different cellular remodeling underlies PPCM and DCM.

We studied myofilament functional and structural remodeling in PPCM and DCM compared with ischemic heart disease (ISHD) and nonfailing controls to define common and unique pathomechanisms for PPCM and DCM. We show that both PPCM and DCM patients experience increased myofilament \(Ca^{2+}\) sensitivity and reduced myofilament length-dependent activation compared with controls. The changes in PPCM samples were more severe than observed in DCM samples, whereas ISHD samples only showed an increased \(Ca^{2+}\) sensitivity without an impairment in length-dependent activation. In addition, PPCM patients showed a significant increase in passive force \((F_{pass})\) development compared with controls, whereas DCM and ISHD samples showed similar \(F_{pass}\) as controls. Also, myofilament remodeling was different between DCM and PPCM patients. Although both patient groups showed a similar increase in compliant titin and fibrosis, only DCM patients expressed the EH-myomesin isoform, which could have contributed to the observed differences in contractile force of myofilaments. The lack of a stabilizing effect of EH-myomesin in PPCM may play a key role in the short-term fast disease progression observed in PPCM. On the other hand, limited remodeling might also explain total recovery in a large group of PPCM patients.

Materials and Methods

Ethical Approval

Left ventricular tissues from DCM, PPCM, and ISHD samples were acquired from the University of Sydney (Sydney, NSW, Australia), with the ethical approval of the Human Research Ethics Committee (number 2012/2814). The codes of used samples are as follows: DCM, 3.107, 4.036, 3.133, 4.125, 2.082, and 3.042; PPCM, 2.048, 3.058, 3.118, and 4.127; and ISHD, 4.111, 4.108, 4.070, and 4.091. Explanted left ventricular heart tissue of healthy donors were used as control samples; the donors died from a noncardiac cause, typically motor vehicle accidents. These healthy donor samples were also acquired from the University of Sydney, and samples included are as follows: 3.162, 6.042, 3.141, 3.164, 5.086, 6.034, 8.004, 7.040, 7.054, 6.008, 5.128, 7.044, 4.104, 6.020, 3.160, and 6.056. In addition, two DCM samples were acquired from the Biobank of the University Medical Center Utrecht (Utrecht, the Netherlands). This study was approved by the Biobank Research Ethics Committee, University Medical Center Utrecht (protocol number WARB 12/387). Written informed consent was obtained. Of the PPCM samples, two were acquired from the University of Hannover (Hannover, Germany). One sample was obtained during implantation of a left ventricular assist device during the...
acute phase of PPCM and the other after cardiac transplantation after chronic heart failure. PPCM tissue analyses were approved by the local ethics commission of the Hannover Medical School (Hannover, Germany); both patients provided written informed consent. All samples were stored in liquid nitrogen or at −80°C until use.

Fibrosis

Cryosections (5 μm thick) were stained with Picrosirius red. The amount of fibrosis was quantified using ImageJ version 1.49 (NIH, Bethesda, MD; http://imagej.nih.gov/ij). The fibrotic area was selected using the Huang thresholding method, threshold color red, color space red green blue, wide open blue and red filters, and adjusted green filter to distinguish stained from nonstained areas. The percentage of fibrosis was calculated as the percentage of red-stained tissue relative to the total area of the image analyzed. Of each sample, at least three images were taken, and data are shown as the average per sample.

Cardiomyocyte Force Measurements

Maximal force (F_{max}) and F_{pass} of sarcomeres were measured at pCa 4.5 and pCa 9.0, respectively, in single membrane-permeabilized cardiomyocytes mechanically isolated from heart tissue, as previously described.21,22 Briefly, a small piece of tissue (10 to 15 mg) was defrosted in 4°C isolation relax solution containing 1 mmol/L free Mg, 139.6 mmol/L KCl, 2 mmol/L EGTA, 5.95 mmol/L ATP, and 10 mmol/L imidazole, with the pH adjusted to 7.0 with KOH. The tissue was cut in small pieces and mechanically disrupted with a Teflon piston for 5 to 10 seconds at 900 × g to obtain a suspension of single cells, small clumps of cells, and cell fragments. To permeabilize the membranes, the cells were incubated with 0.5% Triton X-100 (Millipore, Burlington, MA) for 5 minutes at 4°C. Cells were washed with isolation relax solution to remove Triton X-100 and kept at 4°C until measurement on the same day. Single cardiomyocytes were selected for measurement on the basis of size (100 to 150 μm long and 10 to 35 μm in diameter) and uniformity of striation pattern. Single cardiomyocytes were attached to stainless steel needles attached to a force transducer and a length motor with silicon-based glue (DB-025, Zwaluw; Den Braven, the Netherlands) while being viewed with an inverted microscope at ×320 magnification. Sarcomere length was determined by spatial Fourier transformation and set at the desired sarcomere length before force measurement, starting with the smallest sarcomere length. The relax (pCa 9.0) and activation (pCa 4.5) solutions contained 6.48 and 6.28 mmol/L MgCl, 5.89 and 5.97 mmol/L Na2ATP, 6.97 and 0 mmol/L EGTA, and 0 and 7 mmol/L Ca^{2+}-EGTA, respectively. Both relax and activation solution also contained 14.5 mmol/L creatine phosphate and 100 mmol/L N,N-bis[2-hydroxyethyl]-2-aminoethane-sulphonic acid, of which the pH was adjusted to 7.1 with KOH. The ionic strength of the solutions was adjusted to 200 mmol/L with potassium propionionic acid. The submaximal pCa solutions were generated by appropriate mixing of the relax and activation solutions. A single batch of pCa solutions was used for all active force measurements in this study to eliminate bias attributable to batch-to-batch variation. All passive and active force measurements were performed at 15°C to ensure stability of the cell during measurement, and temperature was controlled with a circulating water bath. For active force measurements, cells were transferred to the desired pCa (-log[Ca^{2+}]) and force development was recorded until steady state had been reached. Cells were then shortened with 30% of cell length to detach cross bridges and to determine total force development. Cells were then transferred back to a relax solution and shortened again with 30% of cell length to calculate passive force development. Active force development was calculated by subtraction of passive force from total force. Force was measured in activation solution (pCa 4.5) first to obtain F_{max} development and thereafter at different pCa solutions in a random order. Finally, force was measured again in activation solution to confirm the cardiomyocyte did not have inappropriate reduction in F_{max} during measurement protocol. Relative forces during the experiment were corrected for decrease in F_{max} during the protocol, and measurements were excluded if the decrease in force was >30%. Passive force curves were measured by stretching the cell to various sarcomere lengths and measurement of passive force in relax solution. In a set of passive force measurements, 2,3-butanedione monoxime was used as an inhibitor for actin-myosin interaction at a concentration of 50 mmol/L in relax solution. All forces were normalized to cross-sectional area of the cardiomyocytes, calculated as (width × depth × π)/4, at a sarcomere length of 2.3 μm. Protein kinase A (PKA) incubations were performed, as previously described.23 In short, membrane-permeabilized cardiomyocytes were incubated with 80 μL of 1 μL/μL PKA (P5511; Sigma, St. Louis, MO) with 0.006 mmol/L cAMP (Sigma). They were incubated in isolation relax solution at 20°C for 40 minutes before measurement of force at 15°C, according to the same protocol described above, without prior PKA incubation. Relative force-[Ca^{2+}] curves were constructed. Ca^{2+} sensitivity was measured as the [Ca^{2+}] needed to achieve 50% of F_{max} (EC_{50}), and length-dependent activation was measured as the shift in EC_{50} (ΔEC_{50}) at a sarcomere length of 1.9 and 2.3 μm. A representation of how EC_{50} and ΔEC_{50} were calculated can be found in Supplemental Figure S1.

Protein Expression and Modification

Titin isoforms were separated on a 1% (w/v) agarose gel and stained with SYPRO Ruby protein gel stain (Invitrogen, Carlsbad, CA), as described previously;24,25 samples were measured in triplicate. The size of N2BA titin was calculated, as described previously.24 In short, homogenates of the samples of interest (PPCM, DCM, and ISHD) were loaded adjacent to a nonfailing control heart and soleus muscle of an adult mouse on a 1%
agarose gel. Proteins were separated in the same way as described above for titin isoform composition. A calibration curve was constructed on the basis of mobility of proteins of known molecular weight (N2A soleus, 3690 kDa; N2B non-failing heart, 2970 kDa; nebulin soleus muscle, 748 kDa; and myosin heavy chain, 200 kDa) relative to myosin heavy chain. This calibration curve was used to calculate the molecular weight of the N2BA titin band of the sample of interest on the basis of relative mobility to myosin heavy chain. An example of a calibration curve and the associated calculation to determine titin N2BA size is depicted in Supplemental Figure S2, A and B. To avoid gel-to-gel variation or variation in mobility within a gel, only the adjacent lanes were used for the construction of the calibration curve. For all samples, the same reference control sample was used, and all samples were measured in duplicate.

To assess titin phosphorylation, homogenates were loaded on a 2.1% acrylamide, 0.5% agarose-strengthened gel and separated, as previously described.\textsuperscript{27} Proteins were transferred to a polyvinylidene difluoride membrane, blocked with 3% bovine serum albumin, and incubated with phosphorylation site–specific antibodies directed to Ser4010 (N2B unique sequence domain; PKA and extracellular signal–regulated kinase 2 target) or Ser12022 (PEVK domain; protein kinase C and Ca\textsuperscript{2+}/calmodulin-dependent protein kinase II target), both from Eurogentec (Liège, Belgium). Incubation of primary antibodies was followed by incubation with secondary horseradish peroxidase–conjugated antibody. An enhanced chemiluminescence detection kit (Amersham; GE Healthcare Life Sciences, Chicago, IL) was used to detect phosphorylated proteins with the Fusion FX 7 (Vilber Lourmat, Collégien, France). Membranes were stripped to remove phosphorylation-specific antibody and blocked again with 3% bovine serum albumin. Membranes were then incubated with an antibody directed to total titin (Eurogentec) and visualized with an enhanced chemiluminescence detection kit (Amersham).

Phosphorylations of cardiac troponin I (cTnI) and cardiac myosin binding protein C (cMyBP-C) were assessed, as previously described.\textsuperscript{27} Phosphorylation of cTnI was further studied by Phos-tag analysis, in which non-phosphorylated, mono-phosphorylated, and bis-phosphorylated cTnI samples (MA1-22700; Pierce, Appleton, WI) were separated by polyacrylamide-bound Mn\textsuperscript{2+}–Phos-tag gel electrophoresis Western blotting, as previously described.\textsuperscript{28} Glutathionylation of cMyBP-C was assessed, as previously described,\textsuperscript{29} with minor deviations from protocol. In short: A nonreducing radioligand precipitation assay buffer was used for protein isolation, to which 25 mmol/L N-ethylmaleimide (Sigma) was added before isolation. After protein isolation, the homogenate was diluted in a 1:1 ratio with 2× Laemmli sample buffer and loaded on Criterion TGX Precast gels, 8% to 16% (Bio-Rad Laboratories, Hercules, CA). Proteins were transferred to nitrocellulose membrane (Bio-Rad Laboratories) and blocked with 5% blocking-grade buffer containing 2.5 mmol/L N-ethylmaleimide. Membranes were cut and incubated with antibodies against glutathione (ab19534; Sigma) or glyceraldehyde-3-phosphate dehydrogenase (14C10; Cell Signaling, Danvers, MA). Membranes were stripped (Restore Western Blot Stripping Buffer; Thermo Fisher Scientific, Waltham, MA), blocked with 5% blocking-grade buffer, and incubated with antibody against cMyBP-C (sc-67354; Santa Cruz Biotechnology, Dallas, TX), followed by incubation with secondary horseradish peroxidase–conjugated antibody. An enhanced chemiluminescence kit (Amersham) was used to detect proteins with the Amersham imager 600. As positive and negative controls for the glutathione Western blot, cardiomyocytes were isolated from a male Wistar rat (weight, 200 g). The animal experiments were performed in accordance with the guidelines from Directive 2010/63/EU of the European Parliament on the protection of animals used for scientific purposes. Liberase TM (Roche, Basel, Switzerland)\textsuperscript{30} was used for isolation of adult rat cardiomyocytes, and cells were suspended in plating medium containing medium 199 (Lonza, Basel, Switzerland), penicillin/streptomycin (1%), and fetal bovine serum (5%). Cardiomyocytes were carefully pipetted on laminin-coated glass coverslips. One hour after plating, cells that were not attached were removed by replacing the plating medium with culture medium that consisted of medium 199, penicillin/streptomycin (1%), and ITS supplement (insulin, 10 mg/L; transferrin, 5.5 mg/L; and selenium, 5 μg/L; Sigma Aldrich, St. Louis, MO). Positive controls were rat cardiomyocytes incubated for 1 hour with various concentrations of the membrane-permeable thiol-oxidizing agent diamide (Sigma). Negative controls were homogenates from rat cardiomyocytes treated with the reducing agents 100 mmol/L dithiothreitol and 1.4 mol/L β-mercaptoethanol before loading. The glutathione Western blot was confirmed to work because adding dithiothreitol and β-mercaptoethanol reduced the glutathione signal and adding diamide in nonreducing conditions increased the glutathione signal in a dose-dependent manner (Supplemental Figure S3).

**EH-Myomesin Expression Immunohistochemistry**

Human heart tissue pieces were cut on a Leica CM 1950 cryostat (Leica Biosystems, Wetzlar, Germany), and sections (12 μm thick) were retrieved on polylysine-coated slides and dried overnight. Sections were fixed for 5 minutes in acetone at −20°C, followed by brief rehydration in phosphate-buffered saline, and 30 minutes blocking in 5% preimmune goat serum (Sigma)/1% bovine serum albumin (Sigma) in gold buffer (20 mmol/L Tris-HCl, pH 7.5, 155 mmol/L NaCl, 2 mmol/L EGTA, and 2 mmol/L MgCl\textsubscript{2}). Mouse monoclonal anti-myomesin (clone B4\textsuperscript{31}) and rabbit polyclonal anti-human EH-myomesin\textsuperscript{32} [generously donated by Dr. Irina Agarkova (Institute of Cell Biology, ETH Zurich, Switzerland)] were diluted in 1% bovine serum albumin/gold buffer and incubated overnight in a humid chamber. After 3 × 5 minutes wash in phosphate-buffered saline, secondary antibodies (Cy3-goat anti-mouse immunoglobulins and Cy2-goat anti-rabbit immunoglobulins; both multilabeling quality from Jackson Immunochemicals, West Grove, PA; via Stratech Scientific, Suffolk, UK) and DAPI were...
incubated for 30 minutes at room temperature in a humid chamber. After 3 × 5 minutes wash in phosphate-buffered saline, slides were mounted in Tris-buffered glycerol containing n-propyl gallate with coverslips and were sealed with nail polish. Imaging was performed using a Leica SP5 confocal microscope (Leica Biosystems) equipped with a 405-nm blue diode and argon laser, a helium neon laser, and oil immersion lenses.

**EH-Myomesin Expression Western Blot**

After SDS-PAGE on a 4% to 8% TruPAGE Precast Gel (Sigma) at 120 V, the human heart tissue samples were transferred overnight onto nitrocellulose (Amersham; GE Health Care) using a Bio-Rad Wet Blot system (Bio-Rad Laboratories) at 55 mA. Transfer was visualized with Ponceau Red, followed by blocking in 5% nonfat dry milk (Sainsbury’s) in low-salt buffer for 1 hour. After incubation with primary antibody against human EH-myomesin and secondary antibody (horseradish peroxidase—conjugated goat anti-rabbit immunoglobulins; Calbiochem; Millipore), the chemiluminescence signal was visualized using Clarity Western ECL Substrate (Bio-Rad Laboratories) and detected with a Bio-Rad ChemiDoc Gel Imaging System (Bio-Rad Laboratories). For quantification, blots were reprobed with a polyclonal antibody against all actin isoforms (Sigma), and densitometry was performed using a Mathematica algorithm generously provided by Dr. Mark Holt (Randall Division, King’s College London, London, UK).

**Statistical Analysis**

Graphpad Prism software version 7 (GraphPad Software Inc., San Diego, CA) was used for statistical analysis. \( F_{\text{max}} \) of patient cardiomyocytes was compared with control cardiomyocytes with one-way analysis of variance. Ca\(^{2+}\) sensitivity was calculated as \( EC_{50} \), and length-dependent activation was calculated as \( \Delta EC_{50} \). The amount of fibrosis, \( EC_{50} \), \( \Delta EC_{50} \), \( N2BA/N2B \) ratio, phosphorylation of cTnl, and glutathionylation of cMyBP-C were compared with one-way analysis of variance. Phosphorylation of cMyBP-C was compared between groups with the nonparametric Kruskal-Wallis test. \( F_{\text{pass}} \) values in patient cardiomyocytes were compared with control cardiomyocytes by two-way analysis of variance. Correlations between \( N2BA/N2B \) and \( N2BA/N2B \) size, \( N2BA/N2B \) and fibrosis, and glutathionylation and phosphorylation of cMyBP-C were studied with linear regression. All values are shown as means ± SEM. Statistical analysis was performed with Graphpad Prism version 7. \( P < 0.05 \) was considered to represent a significant difference. \( P \) values are indicated in figures versus controls.

**Results**

**High \( F_{\text{pass}} \) and Blunted Length-Dependent Activation of PPCM Myofilaments**

The 16 control samples used for experiments were derived from 7 females and 9 males (age, 44.1 ± 2.9 years), the PPCM samples were from 6 females (age, 35.7 ± 3.2 years), the DCM samples were from 3 females and 5 males (age, 50.0 ± 2.9 years), and the ISHD samples were from 4 males (age, 56.0 ± 2.7 years). To establish changes in functional remodeling of cardiomyocytes, myofilament force measurements were performed in membrane-permeabilized cardiomyocytes. \( F_{\text{max}} \) was similar in all groups (Figure 1A). \( F_{\text{pass}} \) was significantly higher in PPCM samples compared with controls, and DCM and ISHD samples showed similar \( F_{\text{pass}} \) as controls (Figure 1B). This was not attributable to altered resting sarcomere length because this was similar in cardiomyocytes from all groups (Figure 1C). The same differences in \( F_{\text{pass}} \) between groups were obtained when measured in relax solution containing 50 mmol/L actin-myosin inhibitor 2,3-butanedione monoxime (Figure 1D), which implies the differences observed are caused by a passive myofilament component. Ca\(^{2+}\) sensitivity of myofilaments was significantly higher in all patient groups compared with controls, evident from the leftward shift of the force-Ca\(^{2+}\) relationship (Figure 1E). There was a significantly lower \( EC_{50} \) in PPCM, DCM, and ISHD samples compared with controls (Figure 1F). Length-dependent activation of myofilaments, indicated by the slope of the curve in Figure 1G, was lower in PPCM and DCM samples, but not in ISHD samples, compared with controls. The decrease in length-dependent activation, expressed as \( \Delta EC_{50} \), was largest in PPCM samples and significantly different from controls (Figure 1H).

**More Compliant Titin and Fibrosis in PPCM and DCM but Not in ISHD**

Titin is a giant sarcomeric protein that spans the entire sarcomere from the Z-disk to the M-band and is important in the regulation of \( F_{\text{pass}} \) through isoform switching and phosphorylation.\(^{32,33} \) In addition, it has been suggested that a switch to the more compliant titin isoform leads to reduced length-dependent activation.\(^{34,35} \) Titin can exist in the small and stiff N2B isoform and in longer and more compliant N2BA isoforms. It has been shown that changes in titin isoform composition and phosphorylation contribute to DCM pathogenesis.\(^{36,37} \) However, it is unknown if titin isoform composition and phosphorylation is altered in PPCM patients. Titin isoforms were separated with gel electrophoresis (Figure 2A). Although PPCM samples showed a nonsignificant increase and DCM samples showed a significant increase in the N2BA/N2B ratio compared with controls, ISHD samples did not differ from controls (Figure 2B). The size of the N2BA titin was also increased in PPCM and DCM samples, whereas a smaller increase in N2BA size was observed in ISHD samples, compared with controls (Figure 2C). The size of the N2BA titin isoforms significantly correlated with the N2BA/N2B ratio in DCM and PPCM samples, but not in ISHD samples and controls (Figure 2D). This implies that there is more N2BA titin and also larger N2BA isoforms in PPCM and DCM samples.
The increases in titin size and N2BA/N2B ratio imply that the heart induces a response to increase cellular compliance, possibly in response to stiffening of the heart by fibrosis. We, therefore, measured the amount of fibrosis by Picrosirius red staining in the same tissue samples in which N2BA/N2B ratio and titin size were measured. Figure 2E shows representative images of tissue samples in which fibrosis is stained in red. A significant increase in fibrosis was found in PPCM and DCM samples compared with controls (Figure 2F), whereas the increase in fibrosis observed in ISHD samples was not significant compared with controls (Figure 2F). Figure 2G illustrates that a higher level of fibrosis coincides with higher N2BA/N2B ratios in PPCM and DCM samples.

Reduced Phosphorylation and Preserved Glutathionylation of cMyBP-C in End-Stage Dilated Cardiomyopathies

Oxidative stress has been shown to play an important role in PPCM pathogenesis and is also reported in other forms of heart failure; it may underlie the changes in cellular function. Oxidative stress can cause glutathionylation of various proteins and thereby alter their function. Glutathionylation of cMyBP-C has been shown to impair phosphorylation of cMyBP-C and subsequently decelerate Ca²⁺-mediated force kinetics in DCM and ISHD. ProQ-stained (phosphorylation; Thermo Fisher Scientific) (Figure 3A) and SYPRO-stained (total protein) (Figure 3A) gel analysis showed decreased phosphorylation of cMyBP-C (Figure 3B) in PPCM, DCM, and ISHD samples compared with controls. Western blots for glutathionylated proteins (glutathionylated cMyBP-C; total cMyBP-C) (Figure 3C) showed there was no significant increase in glutathionylated cMyBP-C (Figure 3D) in PPCM, DCM, and ISHD samples compared with controls. Also, no correlation was found between the amount of glutathionylated cMyBP-C and phosphorylation of cMyBP-C (Figure 3E). This implies that the decreased phosphorylation of cMyBP-C in these samples was not caused by glutathionylation of cMyBP-C. The decreased phosphorylation of cMyBP-C could have been caused by desensitization of the β-adrenergic receptors and subsequent impaired PKA-mediated phosphorylation, a mechanism confirmed in various forms of heart failure.

Figure 1 Baseline characteristics. A: Maximal force (Fmax) was not significantly different in PPCM (25.1 ± 2.3 kN/m²), DCM (27.9 ± 1.7 kN/m²), and ISHD (30.9 ± 1.9 kN/m²), compared with controls (29.7 ± 1.8 kN/m²). B: Passive force (Fpass) was significantly increased in PPCM (P < 0.0001), compared with controls. Fpass was not significantly altered in DCM and ISHD, compared with controls. C: Resting sarcomere length was not different between PPCM (1.82 ± 0.012), DCM (1.79 ± 0.009), ISHD (1.79 ± 0.011), and controls (1.80 ± 0.005). D: Fpass measured in relax solution containing 50 mmol/L 2,3-butanedione monoxime showed significantly increased Fpass in PPCM (P = 0.0001), compared with controls, whereas Fpass was not significantly altered in DCM and ISHD, compared with controls. E: The force-calcium curve was shifted to the left in PPCM, DCM, and ISHD samples, compared with controls. F: Ca²⁺ sensitivity, measured as 50% of Fmax (EC₅₀), was significantly increased in PPCM (1.19 ± 0.08 μmol/L, P < 0.08), DCM (1.27 ± 0.08 μmol/L, P < 0.0001), and ISHD (1.25 ± 0.10 μmol/L, P < 0.0001), compared with controls (1.84 ± 0.05 μmol/L). G: Length-dependent activation, indicated by the increase in Ca²⁺ sensitivity on stretch, was decreased in PPCM and DCM, but not in ISHD, compared with controls. H: The shift in Ca²⁺ sensitivity, indicated by ΔEC₅₀, was significantly reduced in PPCM (0.26 ± 0.05 μmol/L, P = 0.0234), compared with controls (0.54 ± 0.08 μmol/L), whereas it was nonsignificantly reduced in DCM (0.41 ± 0.03 μmol/L) and similar to controls in ISHD (0.52 ± 0.06 μmol/L). Data are expressed as means ± SEM. n = 4 PPCM samples (A–C, E, F, and H); n = 16 DCM cardiomyocytes (A); n = 6 DCM samples (A): n = 27 DCM cardiomyocytes (A); n = 4 ISHD samples (A–C, E, F, and H); n = 15 ISHD cardiomyocytes (A); n = 7 control samples (A, B, E, F, and H); n = 30 control cardiomyocytes (A); n = 14 PPCM cardiomyocytes (B); n = 6 DCM samples (B, C, E, F, and H); n = 15 DCM cardiomyocytes (B); n = 9 ISHD cardiomyocytes (B); n = 18 control cardiomyocytes (B); n = 40 PPCM cardiomyocytes (C); n = 36 DCM cardiomyocytes (C); n = 22 ISHD cardiomyocytes (C); n = 6 control samples (C); n = 7 control cardiomyocytes (C); n = 3 PPCM samples (D); n = 13 PPCM cardiomyocytes (D); n = 3 DCM samples (D); n = 8 DCM cardiomyocytes (D); n = 3 ISHD samples (D); n = 8 ISHD cardiomyocytes (D); n = 6 control samples (D); n = 11 control cardiomyocytes (D); n = 8 PPCM cardiomyocytes (E, F, and H); n = 13 DCM cardiomyocytes (E, F, and H); n = 6 ISHD cardiomyocytes (E, F, and H); n = 17 control cardiomyocytes (E, F, and H); *P < 0.05, ****P < 0.0001 versus controls.
of Ser4010 was decreased in PPCM samples compared with controls (Figure 4, A and C). However, Ser4010 phosphorylation was also decreased in DCM samples (Figure 4, A and C), whereas Fpass was not altered in DCM samples. An alternative hypothesis for the discrepant change in Fpass in PPCM and DCM would be that protein kinase Cε mediated phosphorylation of titin is increased in PPCM, leading to an increased Fpass.42 However, protein kinase Cε mediated phosphorylation at Ser12022 was decreased, rather than increased, in PPCM, and unaltered in DCM and ISHD compared with controls (Figure 4, B and D). To determine whether the decrease in Ser4010 phosphorylation was causal to the increase in Fpass in PPCM samples, the experiments were repeated after incubation with exogenous PKA. Indeed, PKA normalized Fpass in PPCM samples (Figure 4E).

cTnI showed prominent monophosphorylation and bisphosphorylation of cTnI in controls, whereas PPCM and DCM samples showed prominent non-phosphorylation and monophosphorylation of cTnI. The ISHD samples showed predominantly monophosphorylated cTnI (Figure 4, G and H). Exogenous PKA normalized myofilament Ca2+ sensitivity in PPCM, DCM, and ISHD samples to control levels (Figure 4, I and J). In addition, exogenous PKA restored ΔEC50 in PPCM and DCM to control levels (Figure 4K).

Expression of EH-Myomesin in DCM, but Not in PPCM and ISHD, Samples

Although Fpass normalized to control levels after PKA in the PPCM samples, we set out to find an explanation for the discrepant finding of Fpass and titin phosphorylation in PPCM, DCM, and ISHD samples. Myomesin is a component of the M-band and is important for sarcomere stability. The fetal form of myomesin, EH-myomesin, has been shown to be a hallmark of DCM and is virtually absent in adult healthy heart tissue.32 It is believed that EH-myomesin is reexpressed to provide stability in

Figure 2 Titin isoform shift in DCM and PPCM and fibrosis in all dilated cardiomyopathies. A: Titin isoforms N2BA and N2B were separated with gel electrophoresis. B: N2BA/N2B was increased in DCM (0.84 ± 0.16, P = 0.0327) and nonsignificantly in PPCM (0.82 ± 0.15), but not in ISHD (0.43 ± 0.07), compared with controls (0.52 ± 0.02). C: The size of N2BA titin was nonsignificantly increased in DCM (3471 ± 39 kDa) and PPCM (3446 ± 39 kDa), compared with controls (3388 ± 16 kDa), whereas a smaller increase in N2BA size was observed in ISHD (3436 ± 7 kDa). D: The size of N2BA was significantly correlated to the N2BA/N2B ratio in PPCM (P = 0.00479, R² = 0.7775) and DCM (P = 0.00365, R² = 0.5450), but not in ISHD, controls, or all groups combined. E: Representative images of cryosections stained with Picrosirius red of a control, PPCM, DCM, and ISHD sample. F: Fibrosis was significantly increased in PPCM (4.51% ± 0.66%, P = 0.0003) and DCM (5.46% ± 0.55%, P < 0.0001), compared with controls (1.21% ± 0.23%), whereas fibrosis was only nonsignificantly increased in ISHD (2.92% ± 0.46%). G: No significant correlation was found between the amount of fibrosis and N2BA/N2B ratio versus controls; P < 0.05 N2BA/N2B ratio versus N2BA size (slope of linear regression). Scale bars = 50 μm (E).
overstretched conditions in DCM. Our study confirmed EH-myomesin was reexpressed in DCM samples (Figure 5, C and G). Expression of EH-myomesin was much lower in ISHD samples (Figure 5, D and H) and almost absent in PPCM samples (Figure 5, B and F). Our control patients did not show expression of EH-myomesin or only showed expression at a low level (Figure 5, A and E). Western blot analysis confirmed high EH-myomesin expression in DCM, but not in PPCM and controls (Figure 5, I and J). Thus, the increase in titin N2BA/N2B ratio is accompanied by an increase in EH-myomesin in DCM, but not in PPCM. The absence of the stabilizing effect of EH-myomesin expression in PPCM may provide an explanation about why PPCM showed more impairment of myofilament function compared with DCM. Titin isoform expression and PKA-mediated myofilament phosphorylation were similar in the two groups.

**Discussion**

The clinical similarities and differences in disease onset, progression, and outcome between PPCM and DCM patients suggest that the cellular pathomechanisms overlap, but may also show distinct changes. Herein, we show that myofilament function is more impaired in PPCM compared with DCM with respect to length-dependent activation and $F_{pass}$. In addition, myofilament remodeling was different, because DCM samples expressed EH-myomesin, whereas PPCM samples did not. On the other hand, PPCM and DCM samples showed similar changes in myofilament Ca$^{2+}$ sensitivity and sarcomeric protein phosphorylation.

Fibrosis Content Is Higher in PPCM and DCM Compared with ISHD and Controls

Although fibrosis is consistently reported in DCM, reports on the occurrence of fibrosis in PPCM patients are conflicting; animal models of PPCM show fibrosis. Magnetic resonance imaging data at diagnosis of PPCM in patients frequently display no hint for fibrosis, a feature that would be consistent with the ability of many patients to fully recover from the disease. In the present study, we found fibrosis in both DCM and PPCM samples. However, the assessment of fibrosis was performed in small tissue areas of samples obtained during heart transplantation. It cannot be excluded that the tissue samples were selectively taken from more fibrotic regions and may, thus, not be representative of whole heart fibrosis content. Techniques, such as magnetic resonance imaging, might give a better impression of whole heart fibrosis content. No late gadolinium enhancement by magnetic resonance imaging was shown in PPCM (two acute phase patients and four patients at a later stage) previously. In addition, our fibrosis analysis was only performed in end-stage explanted heart tissue of PPCM patients and might, therefore, have been induced over time and may not reflect the acute or short-term PPCM situation. The ISHD samples were taken from the remote area and not within the infarct zone. Although PPCM and DCM are more likely to show a diffuse fibrosis pattern and, therefore, any part of the heart taken might show fibrosis, the fibrosis in ISHD is more likely to be localized in or close to the infarct area. This could explain why we found little fibrosis in the ISHD samples.
PKA-Mediated Hypophosphorylation in All Dilated Cardiomyopathies

Independent of the initial cause, all forms of dilated cardiomyopathy (PPCM, DCM, and ISHD) exhibited higher myofilament Ca\(^{2+}\) sensitivity compared with controls, whereas only PPCM samples had an increased F\(_{\text{pass}}\). The increase in myofilament Ca\(^{2+}\) sensitivity was attributed to decreased PKA-mediated phosphorylation of cTnI because incubation with exogenous PKA restored Ca\(^{2+}\) sensitivity to controls in all groups. The increased F\(_{\text{pass}}\) in PPCM samples was also restored with exogenous PKA, and a decreased PKA-mediated phosphorylation of titin was confirmed at Ser4010. In addition, we also found a decreased phosphorylation of cMyBP-C, which was not related to glutathionylation of cMyBP-C. These combined results strengthen the suggestion that PKA-mediated phosphorylation was impaired in all heart failure samples. The β-adrenergic receptors were shown to be desensitized, leading to impaired PKA-mediated phosphorylation of sarcomeric proteins in various forms of heart failure.\(^{41}\) We have now confirmed PKA-mediated phosphorylation is impaired in...
end-stage PPCM as well. Despite a similar down-regulation of cTnI phosphorylation, the ΔEC50 was significantly decreased only in PPCM samples compared with controls. Although DCM samples showed a nonsignificant impairment of length-dependent activation, ISHD samples had normal length-dependent activation values compared with controls, despite their increased Ca2+ sensitivity. Decreased length-dependent activation has been reported for samples with compliant titin,35,53 and an increase in compliant titin has been reported in DCM.36,37 We observed an increase in the N2BA/N2B ratio in PPCM and DCM samples compared with controls. In addition, an increase in size of the compliant titin isoform was found, corresponding to 58 kDa in PPCM, 93 kDa in DCM, and 48 kDa in ISHD samples, compared with controls. Although this increase was not significant, it is of a magnitude that should not be ignored because such an increase in molecular weight might alter protein function. The fact that titin N2BA size and the N2BA/N2B ratio were significantly correlated in DCM and PPCM samples indicates that the same regulatory mechanism is responsible for both the shift in isoform and the inclusion of additional exons to make N2BA bigger. This observation is in line with the increase in titin N2BA size and N2BA/N2B ratio found in a patient with a mutation in RBM20.24 The observation that in ISHD, N2BA size was increased, whereas N2BA/N2B ratio was unaltered, implies the isoform shift is preceded by inclusion of additional exons in N2BA, after which both titin size and N2BA expression increase further. Both cTnI phosphorylation and titin isoform composition are regulators of myofilament length-dependent activation.34 The decreased cTnI phosphorylation and increased N2BA/N2B and N2BA size explain the blunted myofilament length-dependent activation in PPCM and DCM samples. The preserved myofilament length-dependent activation in ISHD samples, which showed decreased cTnI phosphorylation and unaltered N2BA/N2B ratio, indicates a synergistic role of cTnI phosphorylation and titin in regulating myofilament length-dependent activation.

Different Myofilament Remodeling of the M-Band in PPCM and DCM

PPCM and DCM samples both had decreased phosphorylation at Ser4010 of titin and a similar increase in N2BA/N2B ratio, whereas Fpass was higher compared with controls only in PPCM. The expression of myomesin isoforms follows the expression of titin isoforms in various muscle types. The
longer compliant isoforms of titin and EH-myomesin are predominantly expressed in the fetal heart and are replaced by the shorter and stiffer myomesin during the progression to adulthood.\textsuperscript{54,55} This was also reported for soleus, where the larger titin N2A isoform\textsuperscript{56} is expressed, and the EH-myomesin dominates.\textsuperscript{57} The reexpression of EH-myomesin in DCM patients could be a result of the induction of the fetal gene program, which has been shown in various forms of heart failure and is believed to be activated to cope with the changes in cardiac demand.\textsuperscript{58} Whether the expression of these fetal genes is beneficial is still under debate. EH-myomesin acts as a minispring and has an additional domain of approximately 200 amino acids, similar to the PEVK domain of titin.\textsuperscript{59} It is believed that EH-myomesin provides stability to the sarcomere in overstretched conditions.\textsuperscript{59} Herein, we presented evidence that expression of EH-myomesin might prevent an increase in $F_{pass}$ and a decrease of length-dependent activation in DCM patients, but not in PPCM patients, who do not express EH-myomesin. Exogenous PKA restored $F_{pass}$ and length-dependent activation in PPCM samples, indicating that the lack of EH-myomesin is overcome by PKA-mediated protein phosphorylation. An \textit{in vitro} study showed phosphorylation of myomesin at Ser482 by PKA and subsequent disturbance of the interaction between titin (through the M4 domain) and myomesin (the My4 to My6 domains).\textsuperscript{60} However, Fukuzawa et al.\textsuperscript{61} could not confirm a titin (M4)—myomesin (My4 to My6) interaction in a pulldown assay or a forced Yeast Two-Hybrid assay. In addition, phosphorylation of myomesin did not affect the myomesin-obscurin interaction. Therefore, the effect of myomesin phosphorylation \textit{in vivo} warrants further research. Nonetheless, in heart failure, where the $\beta$-adrenergic receptors are desensitized and PKA-mediated phosphorylation of sarcomeric proteins, such as cTnI and titin, is decreased, the expression of EH-myomesin may play an important role in stabilization of the sarcomeres to maintain or limit the increase in $F_{pass}$ and impairment of length-dependent activation.

\textbf{Clinical Perspective}

Although PPCM and DCM have similar clinical phenotypes, they differ in their clinical progression and, therefore, have a different prognosis. We show herein that myofilament remodeling is different in PPCM and DCM. We also show that PPCM samples have impaired PKA-mediated phosphorylation, similar to what has been observed in DCM samples, but that the effect on myofilament function is larger in PPCM. These observations could provide a possible explanation about why PPCM patients have such a strong response to stimulation of their $\beta$-adrenergic system.\textsuperscript{20} The notion that we could restore all parameters with exogenous PKA indicates that both PPCM and DCM patients might benefit from an increase in $\beta$-adrenergic stimulation to normalize myofilament function. However, we have discussed before that dobutamine, a $\beta$-adrenergic receptor agonist, worsens heart failure in PPCM patients and STAT3 knockout mice, and is associated with adverse outcome independent of treatment with bromocriptine.\textsuperscript{13,20} These results question whether the $\beta$-adrenergic system is, therefore, a realistic clinical target in PPCM treatment. However, increasing $Ca^{2+}$ sensitivity has been a target for the treatment of systolic dysfunction because a greater force is generated at lower calcium concentrations to achieve sufficient contractile power in systolic dysfunction. We show that the myofilaments are already very $Ca^{2+}$ sensitive and this should be taken into consideration during treatment. In addition, too high $Ca^{2+}$ sensitivity might cause
contraction at diastolic calcium concentrations and thereby contributes to cardiac dysfunction.

Conclusion

We show that different forms of heart failure (PPCM, DCM, and ISHD) share aspects of underlying pathomechanisms, but they also differ in important aspects. An overview of the changes in each form of heart failure is shown in Figure 6. PPCM, DCM, and ISHD have decreased cMyBP-C phosphorylation and increased Ca$^{2+}$ sensitivity induced by decreased phosphorylation of cTnI. However, PPCM and DCM share additional pathomechanisms, such as increased compliant titin, reduced phosphorylation of titin at Ser4010, and impaired myofilament length-dependent activation, observations that were not seen in ISHD. The increased myofilament Ca$^{2+}$ sensitivity and decreased length-dependent activation were more pronounced in PPCM samples. Both parameters could be restored after incubation with exogenous PKA. In addition, DCM, but not PPCM, patients show induction of the expression of EH-myomesin. The lack of EH-myomesin reexpression might have contributed to an increase in $F_{\text{pass}}$ and severely blunted length-dependent activation in PPCM patients in situations of low PKA-mediated phosphorylation because of instability of the sarcomeres in overstretched conditions. Therefore, this study shows myofilament remodeling and function is more impaired in PPCM, compared with DCM and ISHD, and the lack of EH-myomesin reexpression may explain the fast deterioration of some PPCM patients to end-stage heart failure.

Limitation to the Study

No tissue from healthy postpartum women could be explored and, therefore, the condition of a healthy maternal heart in the postpartum phase is not known. In addition, the different heart failure groups could not be completely matched regarding age and sex of patient samples studied. However, no difference was observed between samples derived from males and females in the DCM group.

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Supplemental Data

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References


