



# Plasma growth differentiation factors 8 and 11 levels in cats with congestive heart failure secondary to hypertrophic cardiomyopathy<sup>☆</sup>

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## KEYWORDS

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metry

**Abstract Objectives:** Growth differentiation factor (GDF) 11 has been shown to reduce cardiac hypertrophy in mice. Low levels of GDF-11 are associated with cardiac hypertrophy in humans. The authors hypothesized that plasma GDF-11 level is decreased in cats with hypertrophic cardiomyopathy (HCM). Given the close homology between GDF-11 and myostatin/GDF-8, GDF-8 levels were also assessed.

**Animals:** Thirty-seven client-owned cats were enrolled, including cats with normal cardiac structure (n = 16), cats with HCM or hypertrophic obstructive

<sup>☆</sup> A unique aspect of the Journal of Veterinary Cardiology is the emphasis of additional web-based materials permitting the detailing of procedures and diagnostics. These materials can be viewed (by those readers with subscription access) by going to <http://www.sciencedirect.com/science/journal/17602734>. The issue to be viewed is clicked and the available PDF and image downloading is available via the Summary Plus link. The supplementary material for a given article appears at the end of the page. To view the material is to go to <http://www.doi.org> and enter the doi number unique to this paper which is indicated at the end of the manuscript.

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cardiomyopathy (HOCM;  $n = 14$ ), and cats with HCM and congestive heart failure (CHF;  $n = 7$ ).

**Methods:** Plasma samples were analyzed for GDF-8 and GDF-11 using liquid chromatography tandem-mass spectrometry. Levels of GDF-8 and GDF-11 were compared between cats with normal cardiac structure, HCM or HOCM, and CHF.

**Results:** No differences in GDF-11 concentrations were found between cats with normal cardiac structure and cats with HCM/HOCM, with or without history of CHF. Decreased GDF-8 concentrations were detected in cats with CHF compared to cats with HCM/HOCM without history of CHF ( $p=0.031$ ) and cats with normal cardiac structure ( $p=0.027$ ). Growth differentiation factor 8 was higher in cats with HOCM compared to those with CHF ( $p=0.002$ ). No statistical difference was noted in GDF-8 level as a function of age, weight, or body condition score.

**Conclusions:** Plasma GDF-11 was not different between cats with HCM/HOCM and cats with normal cardiac structure regardless of age. Plasma GDF-8 was decreased in cats with CHF compared to cats with normal cardiac structure and cats with asymptomatic HCM/HOCM, suggesting a possible role in CHF development.

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### Abbreviations

CHF	congestive heart failure
FA	formic acid
GDF	growth differentiation factor
HCM	hypertrophic cardiomyopathy
HOCM	hypertrophic obstructive cardiomyopathy
LC-MS/MS	liquid chromatography tandem- mass spectrometry
LV	left ventricle

### Introduction

Hypertrophic cardiomyopathy (HCM) is the most common cardiac disease in cats with a prevalence of 14–16% [1,2]. This disease is found in a variety of cat breeds, with high incidence in Ragdoll cats and Maine Coons. In both breeds, mutation in myosin binding protein C has been identified [3], although involvement of additional genetic mutations is suspected. Given the need to better understand the causes of ventricular hypertrophy and to develop appropriate therapy for both people and veterinary patients, rodent models have been studied and have demonstrated some encouraging results. Loffredo et al. [4] showed that age-related left ventricular (LV) hypertrophy in mice is associated with decreased levels of circulating growth differentiation factor (GDF) 11 as measured with an antibody that also recognizes GDF-8, or myostatin. Growth differentiation factor 11 is a member of the transforming growth

factor- $\beta$  superfamily. It is found postnatally in the spleen, pancreas, kidney, and skeletal muscle [5]; and it is also found in cardiac tissues in mice but at a lower level [6]. Resupplying GDF-11 to aged mice reduces cardiac hypertrophy in a dose-dependent fashion [4,6,7]. This response to GDF-11 treatment raises the possibility that levels of circulating GDF-11 and/or GDF-8 are related to the degree of cardiac hypertrophy [4].

There has been considerable controversy involving methods used to measure GDF-11 in previous studies, which have included polymerase chain reaction, enzyme-linked immunosorbent assay, and proteomic microarrays [4,8,9]. The close homology between GDF-11 and GDF-8 may result in cross-reactivity in certain assays as these two proteins share approximately 90% sequence identity in their mature forms [5]. Both proteins belong to the activin/myostatin subclass of the transforming growth factor- $\beta$  family, activate similar receptors, and therefore share many redundant functions [10]. Growth differentiation factor 8, or myostatin, is expressed postnatally by skeletal and cardiac tissues and suppresses skeletal muscle mass [5]. Expression of GDF-8 is associated with a reduction in cardiomyocyte proliferation without a negative effect on systolic function [8]. Conflicting results regarding levels of either GDF-8 or GDF-11 with age, heart disease, therapy, and prognosis may be the result of technical difficulties associated with separate detection of the two proteins [4,7,10–20]. This may also have added confusion as to how GDF-8 or GDF-11 overexpression may alter cardiac morphology and function [4,6,21–23]. Overcoming the limitations of traditional analytical techniques in

differentiating GDF-8 and GDF-11 requires liquid chromatography tandem-mass spectrometry (LC-MS/MS). This method accurately differentiates GDF-8 and GDF-11 in humans and rats [18,24]. This method resolves multiple distinct residues within GDF-8 and GDF-11, allowing differentiation between these proteins despite their high percentage of homology [18]. In feline samples, LC-MS/MS would be expected to accurately differentiate between GDF-8 and GDF-11 given that there is 100% homology between the mature domains of feline and human GDF-8 and GDF-11 (Fig. 1) [6,25]. In contrast to the mature domains of feline GDF-8 and GDF-11 that exhibit 90% homology, the prodomains exhibit greater differences with only a 54.7% homology between the two proteins [25].

Few studies have investigated the effects and levels of circulating GDF-11 in veterinary patients. Ahn et al. [26] showed that circulating GDF-11 levels did not change significantly in dogs that developed congestive heart failure (CHF) secondary to myxomatous mitral valve disease. Furthermore, GDF-11 did not change as a function of patient age, body weight, and echocardiographic variables. Studies of GDF-11 have not been reported in cats. Cardiac hypertrophy in the form of LV free wall or septal wall thickening is often observed in cats with primary HCM, systemic hypertension, or hyperthyroidism. Given the possible relationship between GDF-11 and cardiac

hypertrophy, the authors investigated circulating GDF-11 in cats with HCM. The authors hypothesized that cats with HCM will have decreased circulating GDF-11 levels compared to cats with normal cardiac structure. We also speculated that there may be a decrease in GDF-11 levels in older cats.

## Animals, materials and methods

Client-owned cats presented to the Foster Hospital for Small Animals at the Cummings School of Veterinary Medicine at Tufts University were enrolled in this study. The study protocol was approved by the Tufts University Clinical Studies Review Committee, and owner consent was obtained before sample collection. Cats received a routine physical examination and echocardiogram examination. Cats with hyperthyroidism, history of systemic hypertension, or aortic stenosis were excluded from the study. Cats with comorbidities including diabetes, neoplasia, and azotemia before starting diuretic therapy were also excluded. Patients were divided into the following five groups: cats with normal cardiac structure <6 years of age (termed 'young normal'); cats with normal cardiac structure ≥6 years of age (termed 'old normal'); cats with HCM without left ventricular tract obstruction; cats with hypertrophic obstructive cardiomyopathy (HOCM); and cats with HCM and clinical signs or history of CHF. Congestive heart

### Mature domains:

GDF8:	DFGLDCDEHS	TESRCCRYPL	TVDFEAFGWD	WIIAPKRYKA	NYCSGCECFV	FLQKYPHTHL
GDF11:	NLGLDCDEHS	SESRCCRYPL	TVDFEAFGWD	WIIAPKRYKA	NYCSGGCEYM	FMQKYPHTHL
GDF8:	VHQANPRGSA	GPCCTPTKMS	PINMLYFNGK	EQIYYGKIPA	MVVDRCGCS	
GDF11:	VQQANPRGSA	GPCCTPTKMS	PINMLYFNDK	QQIYYGKIPG	MVVDRCGCS	

### Prodomains:

GDF8:	QKLQIYVYIY	LFMLIVAGPV	DLNENSEQKE	NVEKEGLCNA	CTWRQNTKSS	RIEAIKIQIL
GDF11:	AEGPAAAAAA	AAAAGA---G	GERSSRPAPS	VAPEPDGCPV	CVWRQHSREL	RLESIKSQIL
GDF8:	SKLRLETAPN	ISKDAIRQLL	PKAPPLRELI	DQYDVQRDDS	-SDGSLEDDD	YHATTETIIT
GDF11:	SKLRLEAPN	ISREVVQQLL	PKAPPLQQL	DLHDFQGDAL	QPEDFLEDE	YHATTETVIS
GDF8:	MPTESDLIMQ	VEGKPKCCFF	KFSSKIQYNK	VVKAQLWIYL	RPVKPTTTFV	VQILRLIKPM
GDF11:	MAQETDPAVQ	TDGSPLCCHF	HFSPKVMFTK	VLKAQLWVYL	RPVPRPATVY	LQILRLKPLT
GDF8:	KD-----	GTRYTGIRSL	KLDMNPGTGI	WQSIDVKTVL	QNWLKQPESN	LGIEIKALDE
GDF11:	GEGTAGGGGG	GRRHIRIRSL	KIELHSRSGH	WQSIDFKQVL	HSWFRQPQSN	WGIEINAFDP
GDF8:	NGHDLAVTFP	GPGEDGLNPF	LEVKVTDTPK	RSRR		
GDF11:	SGTDLAVTSL	GPGAEGLHFP	MELRVLENTK	RSRR		

**Figure 1 Protein structure of GDF 8 and 11 in cats.** The mature domain and prodomain amino acid sequences of GDF-8 and GDF-11 in cats are shown and compared. There is a 90% homology between the mature domains of the two proteins and a 54.7% homology between the prodomains. Amino acid differences are highlighted in red. GDF = growth differentiation factor.

failure was defined as presence of pulmonary edema on thoracic radiographs or pleural or pericardial effusion on echocardiogram, and requiring treatment with furosemide. Echocardiography was performed without sedation by either a board-certified cardiologist or a cardiology resident under the supervision of a board-certified cardiologist. Measurements recorded included M-mode measurements of interventricular septal thickness in diastole and in systole (outer edge to outer edge), LV internal dimension in diastole and in systole (inner edge to inner edge), LV free wall thickness in diastole and in systole (outer edge to outer edge), and fractional shortening, all from the right parasternal short-axis view. The left atrial-to-aortic ratio was measured from both M-mode and 2-D right parasternal short-axis view (leading edge to leading edge). Cats in the normal groups were >1 year of age and had LV wall thicknesses <6 mm in diastole. Cats with a heart murmur as a result of dynamic right ventricular outflow tract obstruction were included in the normal groups if the LV walls in diastole did not measure thicker than 6 mm. Hypertrophic cardiomyopathy was defined as LV free wall or interventricular septum thickness in diastole, either focal or generalized,  $\geq 6$  mm without evidence of LV outflow tract obstruction. Cats in the HOCM group had evidence of LV outflow tract obstruction defined as late systolic LV outflow tract velocity >2.5 m/s [27] in addition to LV wall thicknesses <6 mm in diastole. One mL of whole blood was collected into EDTA collection tubes. Plasma was isolated by centrifugation of whole blood sample at  $1,320\times g$  for 10 min at room temperature immediately after blood collection to remove cells and platelets. Plasma was stored at  $-80$  °C until LC-MS/MS analysis.

Samples were denatured and reduced in 6 M urea and 20 mM dithiothreitol at 60 °C for 40 min and alkylated with 35 mM iodoacetic acid at 37 °C for 30 min. After acidification using 1.5 mL 0.1% formic acid (FA) solution, the samples were loaded into conditioned strong cation exchange solid phase extraction 96 well plates<sup>f</sup> using 1 mL of methanol, followed by 1 mL of 0.1% FA, and washed by 1 mL of 0.1% FA and 1 mL of 10 mM bis-Tris, pH 5.8 buffer. Bound proteins were eluted in 1.5 mL of buffer solution containing 10% methanol in 50 mM Tris at pH 10.5. Eluents were adjusted to pH 8 using 0.1–1 M hydrochloric acid and mixed with 50  $\mu$ L of 100  $\mu$ g/mL trypsin in 50 mM Tris buffer at pH 8.

Then, 50  $\mu$ L of 5 ng/mL isotope-labeled peptides prepared in 0.1% FA was added as internal standard. The digested solution was then loaded into a Strata-X 33  $\mu$ m solid phase extraction system<sup>g</sup> conditioned with 1 mL methanol followed by 1 mL of deionized water. Samples were eluted with 1 mL methanol after washing with 1 mL deionized water and evaporated to dry. Residues were reconstituted with 50  $\mu$ L of 5% methanol and 0.1% FA in deionized water for LC-MS/MS analysis. An ultra-performance liquid chromatography system consisting of LC-20 AD<sub>XR</sub> Binary Pump and SIL-20 AC<sub>XR</sub> Autosampler<sup>h</sup> was used for liquid-chromatographic separation. Chromatographic separation was achieved using Aeris Peptide 3.6  $\mu$ m XB—C18.<sup>i</sup> Column and autosampler temperatures were set at 45 °C and 10 °C, respectively. The mobile phase consisted of 0.1% FA in deionized water and 0.1% FA in methanol. A hybrid triple quadrupole/liner ion trap mass spectrometer<sup>j</sup> equipped with a Turbo V ion source was used for detection. The mass spectrometer was operated in positive electrospray ionization mode using an ion-spray voltage of 5500. The most intense unique surrogated peptides and their multiple reaction monitoring transitions were identified by peptide mapping using an information-dependent acquisition experiment; the ideal multiple reaction monitoring transitions were further optimized by the autotune function of Analyst Software by infusion of desalted tryptic digest of GDF-11 and GDF-8 proteins. Ion source optimization was performed by flow injection using 0.1% FA in deionized water/methanol at 70/30 (v/v) at 600  $\mu$ L/min or 200  $\mu$ L/min depending on the columns used. Calibration curves were prepared by assaying commercial pooled human plasma spiked with eight concentrations of GDF-11 or GDF-8 (0.5, 1, 5, 10, 25, and 50 ng/mL). In addition, isotope-labeled IPGMVVDR and NLGLDEHSSESER peptides were added as internal standards to a final concentration of 5 ng/mL after pH-based fractionation by conditioned strong cation exchange solid phase extraction and tryptic digestion. The lower limit of quantitation is 1 ng/mL. Growth differentiation factor 8 and immunoglobulin G have no detectable cross-reactivity in the GDF-11 assay. The inter-assay coefficient of variation was less than 10%.

Statistical analysis was performed using commercially available statistics software.<sup>k</sup>

<sup>f</sup> StrataTM—XL-C 100  $\mu$ m, PN: 8E-S044-TGB, Phenomenex Inc., Torrance, CA.

<sup>g</sup> PN: 8E-S100-AGB, Phenomenex Inc., Torrance, CA.

<sup>h</sup> Shimadzu, Framingham, MA.

<sup>i</sup> Phenomenex Inc., Torrance, CA.

<sup>j</sup> AB Sciex QTRAP® 5500, Framingham, MA.

<sup>k</sup> IBM SPSS Statistics for Windows, Released 2017, Version 25.0. Armonk, NY.

Shapiro–Wilk test was used to test for normality; and Kruskal–Wallis test was used for non-parametric statistical testing of differences among more than two groups. Mann–Whitney *U* test or *t*-test (normally distributed data) was used for two-way comparisons. Bonferroni correction was applied to multigroup comparison.

## Results

Thirty-seven cats were enrolled into five groups: old normal group (*n* = 7), young normal group (*n* = 9), HCM group (*n* = 8), HOCM group (*n* = 6), and HCM with CHF group (*n* = 7). The demographics of the cats are reported in Table 1. There was no difference in body weight or body condition score (on a scale of 1–9) among the five groups. Cats in young normal group (mean age 2.4 years ± 1.1) were significantly younger than cats in the old normal group (mean age 10 years ± 3.3, *p*=0.0003), cats in the HCM or HOCM groups (mean age 8.8 years ± 3.2, *p*=0.00001), and cats in the CHF group (mean age 9.1 years ± 2.9, *p*=0.001).

Echocardiographic findings are reported in Table 2. The *p*-values for comparisons are listed in Table 3. All six cats in the HOCM had focal hypertrophy of the interventricular septum in diastole measuring >0.65 cm and systolic anterior motion of the mitral valve. Cats in the young normal group had larger LV internal dimension in diastole measurements compared to older normal cats (*p*=0.018). Fractional shortening was also

lower in cats with CHF compared to cats in the old normal group (*p*=0.004) and cats with HOCM (*p*=0.008).

Circulating GDF-11 levels were not different between the normal, HCM, HOCM, and CHF groups (Figs. 2–4). In contrast, circulating GDF-8 levels were decreased in cats with HCM and CHF compared to cats (both young and old) with normal cardiac structure (*p*=0.027), and cats with HCM or HOCM without CHF (*p*=0.031). In particular, cats with HOCM had significantly higher GDF-8 levels than those in the CHF group (*p*=0.002). Differences were not identified based on age, weight, or sex for either GDF-8 or GDF-11 (Figs. 5–7).

## Discussion

This is the first report in cats of the association between LC-MS/MS measured circulating GDF-8 and GDF-11 and HCM/HOCM or CHF. Age-related changes in GDF-8 or the GDF-11 levels were not observed in cats enrolled in this study. Circulation levels of GDF-11 were not different between normal cats compared to cats with asymptomatic HCM/HOCM or CHF. Circulating levels of GDF-8 and GDF-11 were not different based on sex, in contrast to humans where GDF-8 and GDF-11 are higher in males [10]. The most significant finding of this study was that circulating levels of GDF-8 were decreased in cats with CHF secondary to HCM when compared to cats with normal cardiac structure or cats with HCM/HOCM without CHF.

**Table 1** Demographics of study cats.

	Total	Median age (range), y	Average weight (range), kg	Average BCS (range)	Gender	Total	Breed	Total
Young normal	9	2.5 (1–4)	5.2 (2.8–7.6)	6 (4–9)	CM	5	Abyssinian	2
					SF	1	DSH	7
					F	2		
Old normal	7	12 (6–14)	4.7 (3.4–5.5)	5.8 (4–7)	CM	3	DSH	5
					SF	5	DLH	2
HCM	8	7 (4–12)	6.5 (5–8.8)	6.5 (6–8)	CM	8	DSH	6
							DMH	1
							DLH	1
HOCM	6	11.5 (6–15)	5.4 (4.3–6.6)	6.0 (6–7)	CM	2	DSH	4
					SF	4	DLH	2
HCM with CHF	7	10 (4–13)	5.6 (3.6–9)	4.5 (4–9)	CM	6	Abyssinian	1
					SF	1	Tonkinese	1
							DSH	2
							DMH	2
						DLH	1	

BCS = body condition score, on a scale of 1–9; CHF = congestive heart failure; CM = castrated male; DLH = domestic long hair; DMH = domestic medium hair; DSH = domestic short hair; HCM = hypertrophic cardiomyopathy; HOCM = hypertrophic obstructive cardiomyopathy; SF = spayed female.

**Table 2** Summary of echocardiographic measurements.

	Young normal	Old normal	HCM	HOCM	HCM with CHF
IVSd, cm	0.45 (0.08)	0.47 (0.04)	0.58 (0.1)	0.65 (0.08)	0.66 (0.21)
Average (SD)					
LVIDd, cm	1.67 (0.12)	1.43 (0.10)	1.38 (0.35)	1.34 (0.15)	1.52 (0.51)
Average (SD)					
LVPWd, cm	0.47 (0.06)	0.43 (0.05)	0.58 (0.10)	0.56 (0.06)	0.63 (0.25)
Average (SD)					
IVSs, cm	0.69 (0.08)	0.75 (0.09)	0.77 (0.14)	0.88 (0.11)	0.80 (0.23)
Average (SD)					
LVIDs, cm	0.86 (0.14)	0.67 (0.22)	0.77 (0.11)	0.60 (0.11)	0.99 (0.56)
Average (SD)					
LVPWs, cm	0.77 (0.88)	0.73 (0.14)	0.89 (0.13)	0.84 (0.12)	0.78 (0.31)
Average (SD)					
FS, %	49.1 (5.5)	58.5 (9.8)	50.4 (5.0)	55.5 (5.0)	38.1 (16.0)
Average (SD)					
LA:Ao	1.24 (0.23)	1.27 (0.17)	1.24 (0.15)	1.37 (0.25)	2.10 (0.40)
Average (SD)					
LA:Ao (M-mode)	1.37 (0.27)	1.41 (0.25)	1.30 (0.24)	1.43 (0.18)	1.91 (0.46)
Average (SD)					
LVOT velocity, m/s	0.8 (0.2)	1.1 (0.6)	0.9 (0.1)	4.0 (0.9)	1.0 (0.2)
Average (SD)					

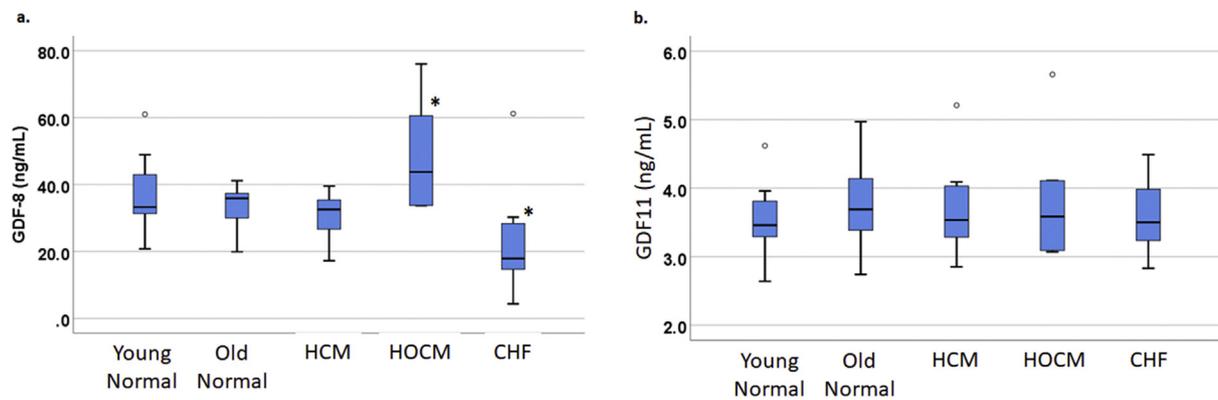
CHF = congestive heart failure; HCM = hypertrophic cardiomyopathy; HOCM = hypertrophic obstructive cardiomyopathy; IVSd = interventricular septum in diastole; LV = left ventricle; LVIDd = LV internal dimension in diastole; LVPWd = LV free wall in diastole; IVSs = interventricular septum in systole; LVIDs = LV internal dimension in systole; LVPWs = LV free wall in systole; FS = fractional shortening; LA:Ao = left atrium-to-aorta ratio in short axis; LVOT = left ventricular outflow tract; SD = standard deviation.

**Table 3** Statistical findings (p-values).

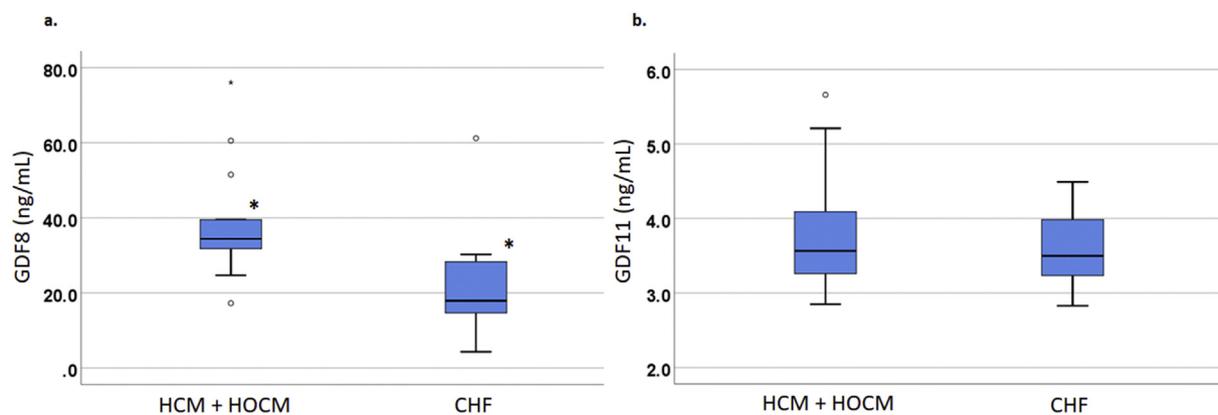
	Young vs. old	Young vs. HCM	Young vs. HOCM	Young vs. CHF	Old vs. HCM	Old vs. HOCM	Old vs. CHF	HCM vs. HOCM	HCM vs. CHF	HOCM vs. CHF
Age	0.0002	0.015	0.0001	0.001	—	—	—	—	—	—
Weight	—	—	—	—	0.01	—	—	—	—	—
BCS	—	—	—	—	—	—	—	—	—	—
IVSd	—	0.023	0.002	0.018	0.038	0.005	0.03	—	—	—
LVIDd	0.018	0.032	0.002	—	—	—	—	—	—	—
LVPWd	—	0.037	—	0.042	0.004	0.01	0.005	—	—	—
IVSs	—	—	0.013	—	—	0.035	—	—	—	—
LVIDs	—	—	0.003	—	—	—	—	0.013	—	—
LVPWs	—	—	—	—	—	—	—	—	—	—
FS	0.047	—	—	—	—	—	0.004	—	—	0.008
LA:Ao	—	—	—	0.002	—	—	0.003	—	0.002	0.023
LA:Ao (M-mode)	—	—	—	0.029	—	—	0.001	—	0.006	—

BCS = body condition score; CHF = congestive heart failure; HCM = hypertrophic cardiomyopathy; HOCM = hypertrophic obstructive cardiomyopathy; IVSd = interventricular septum in diastole; LV = left ventricle; LVIDd = LV internal dimension in diastole; LVPWd = LV free wall in diastole; IVSs = interventricular septum in systole; LVIDs = LV internal dimension in systole; LVPWs = LV free wall in systole; FS = fractional shortening; LA:Ao = left atrium-to-aorta ratio.

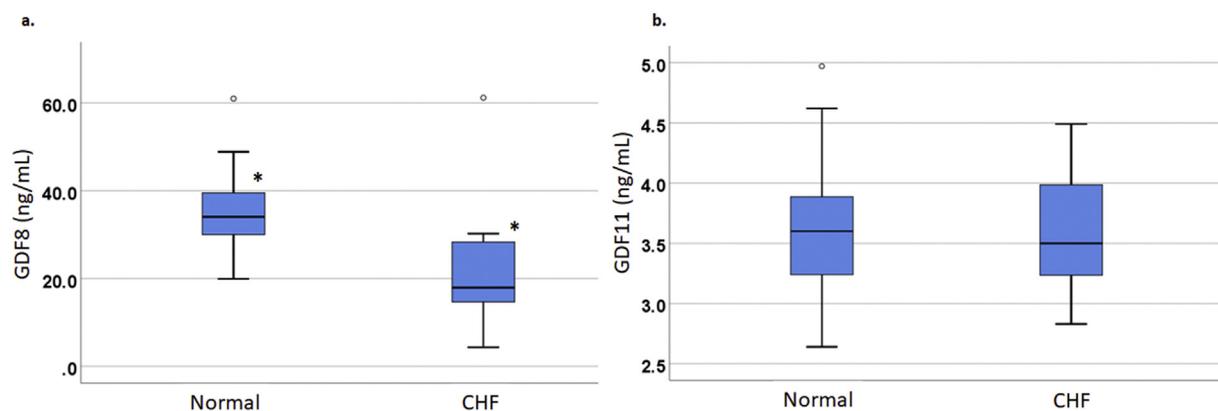
Only p-values that are <0.05 are shown. Bonferroni corrected p-value = 0.005.



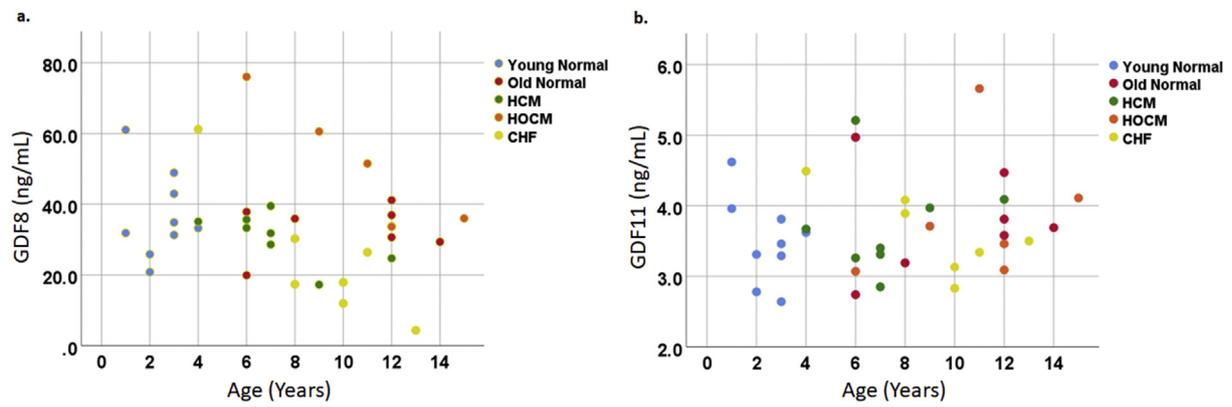
**Figure 2** Growth differentiation factor 8 and 11 levels. The GDF-8 levels for the five different disease groups are shown in (a) ( $n = 7$ /group for old normal,  $n = 8$  for HCM,  $n = 6$  for HOCM,  $n = 7$  for CHF, and  $n = 9$  for young normal). GDF-11 levels for each of the groups are shown in (b). Open circles represent outliers. \* Denotes statistically significant differences. CHF = congestive heart failure; GDF = growth differentiation factor; HCM = hypertrophic cardiomyopathy; HOCM = hypertrophic obstructive cardiomyopathy.



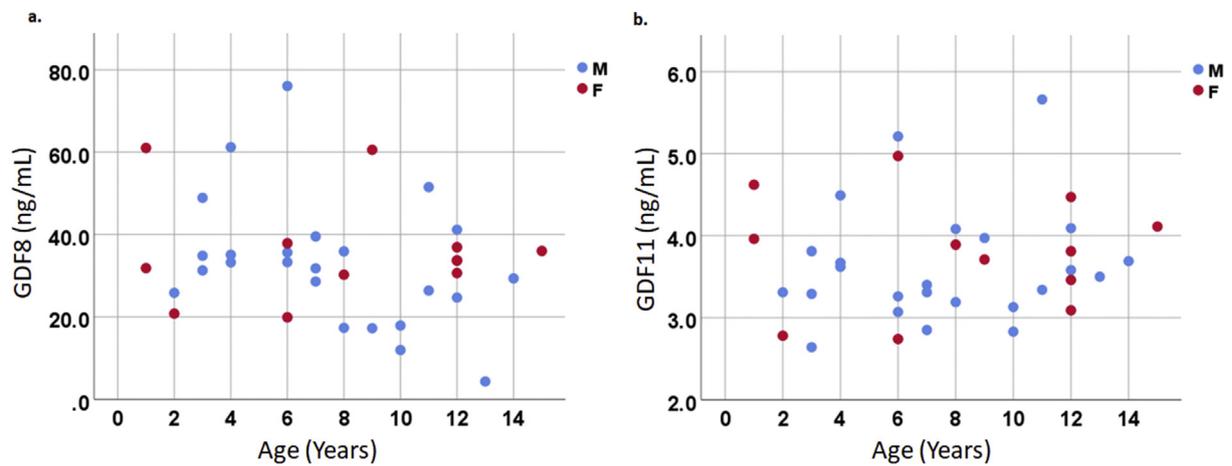
**Figure 3** Comparison of cats with CHF, HCM, and HOCM. Cats with CHF ( $n = 7$ ) had lower GDF-8 levels compared to cats with HCM and HOCM ( $p=0.031$ ,  $n = 14$ ) (a). No difference was detected between groups for GDF-11 (b). Open circles and small asterisk represent outliers. \* Denotes statistically significant differences. CHF = congestive heart failure; GDF = growth differentiation factor; HCM = hypertrophic cardiomyopathy; HOCM = hypertrophic obstructive cardiomyopathy.



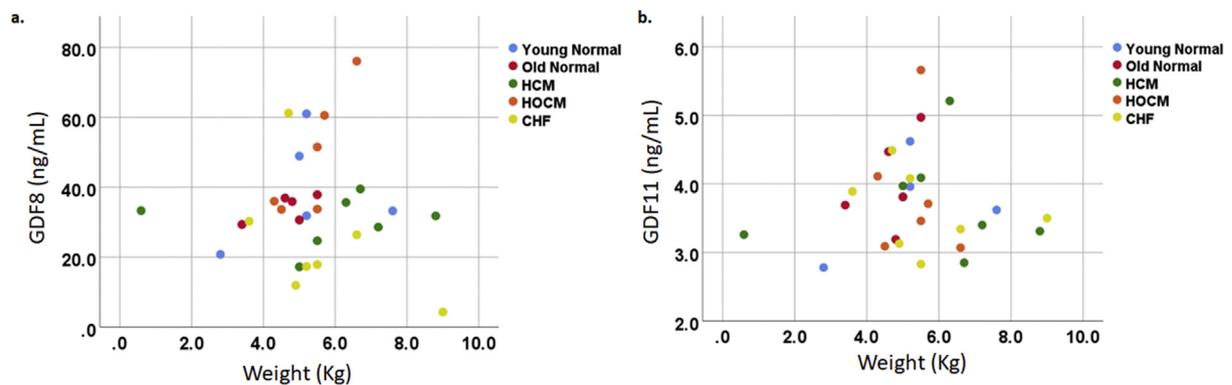
**Figure 4** Comparison of cats with CHF to cats with normal hearts. Cats with CHF ( $n = 7$ ) had lower GDF-8 levels compared to cats with no heart disease ( $p=0.027$ ,  $n = 16$ ) (a); whereas no difference was detected between groups for GDF-11 (b). Open circles represent outliers. \* Denotes statistically significant differences. CHF = congestive heart failure; GDF = growth differentiation factor.



**Figure 5** Scatter plots of GDF-8 and GDF-11 levels as a function of age and disease group. Individual data points for GDF-8 (a) and GDF-11 (b) are shown. Each patient point is labeled by disease status: blue (young normal), red (old normal), green (HCM), orange (HO CM), and yellow (CHF). CHF = congestive heart failure; GDF = growth differentiation factor; HCM = hypertrophic cardiomyopathy; HO CM = hypertrophic obstructive cardiomyopathy.



**Figure 6** Scatter plots of GDF-8 and GDF-11 levels as a function of age and sex. Individual data points for GDF-8 (a) and GDF-11 (b) are shown. Each patient point is labeled by sex: blue (male) and red (female). GDF = growth differentiation factor.



**Figure 7** Scatter plots of GDF-8 and GDF-11 levels as a function of weight and disease group. Individual data points for GDF-8 (a) and GDF-11 (b) are shown. Each patient point is labeled by disease status: blue (young normal), red (old normal), green (HCM), orange (HO CM), and yellow (CHF). CHF = congestive heart failure; GDF = growth differentiation factor; HCM = hypertrophic cardiomyopathy; HO CM = hypertrophic obstructive cardiomyopathy.

The absence of differences in circulating GDF-11 levels associated with aging or the development of cardiac hypertrophy is in contrast with previous reports in aging mice, aging humans, and humans with advanced heart disease as classified by the New York Heart Association class ranking [4,10,18]. Contrary to the studies in mice and humans, the observed ventricular hypertrophy in cats enrolled in this study was secondary to primary HCM and not as a result of aging alone. Unpublished observations from the authors' laboratory indicate that the age-dependent variations in circulating GDF-8 and GDF-11 in rodents represent changes in specific isoforms of these two proteins rather than the total GDF-8 and GDF-11 levels. Changes in specific protein isoforms were not investigated in this study.

There are conflicting reports in humans regarding the association between CHF and circulating GDF-8 levels. Although some studies report decreased GDF-8 levels associated with ischemic or non-ischemic heart failure [11,13], others report increased GDF-8 levels with CHF secondary to dilated cardiomyopathy [14] or with decompensated CHF [15,20]. In this study, cats with HCM had decreased circulating GDF-8 levels once heart failure had developed. The cause of decreased circulating GDF-8 in cats with CHF might be the result of decreased production from either cardiac or skeletal muscle. Decreased production from the cardiac tissue might be secondary to loss of cardiomyocytes and increased cardiac fibrosis, which is often found in cats with advanced heart disease. Confirmation of these causes for the decreased GDF-8 production would require analysis of heart and muscle tissues. Changes in patient body condition score were not observed in cats with CHF in this study, although muscle condition scores measuring cachexia were not obtained and thus skeletal muscle loss cannot be ruled out.

Despite the decrease in GDF-8 levels seen in cats with CHF, it is unclear if GDF-8 supplementation in cats with CHF would be beneficial. The decrease may be a compensatory outcome needed to maintain cardiac function. In this case, GDF-8 supplementation would be detrimental, although increases in GDF-8 have not been shown to negatively affect systolic function. Cats with CHF had concurrent decreases in GDF-8 level and fractional shortening compared to the other groups; however, a causative relationship between these two findings cannot be established from this study. Studies in human patients showed that those in the highest quartile for combined GDF-11 and GDF-8 levels had markedly lower risk of reaching study end-points related to

cardiovascular mortality [6]. These studies concluded that GDF-11 and/or GDF-8 protect against adverse cardiovascular events. Studies of recombinant GDF-8 supplementation or GDF-8 antagonists might confirm the effect of GDF-8 levels on cardiac function.

One of the main limitations of this study is the small sample size. Larger sample sizes will be needed to confirm the findings in this study. Based on the authors' data from this study, detecting a significant difference in GDF-11 levels between young normal cats and cats with HCM or HOCM would require 60 patients. As discussed above, tissue analysis might also aid in determining the cause of decreased circulating GDF-8. This study also did not investigate changes in specific GDF-8 or GDF-11 isoforms but rather measured the total GDF-8 and GDF-11. Lastly, muscle condition data might have provided additional insight into the changes in the GDF-8 level as this protein is found in skeletal muscles in addition to cardiac muscle.

## Conclusions

Circulating GDF-11 in cats was not associated with age or the development of HCM or CHF. A decrease in circulating GDF-8 was found in cats with CHF secondary to HCM/HOCM. These results suggest that the process by which cardiac hypertrophy develops in laboratory mice may be different from what occurs in cats with HCM or HOCM.

## Conflicts of Interest Statement

The authors do not have any conflicts of interest to disclose.

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## Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.jvc.2019.08.002>.

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