

Impaired myocardial capillarogenesis and increased adaptive capillary growth in FGF2-deficient mice

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Basic fibroblast growth factor (FGF2) plays a major role in angiogenesis and capillary growth. In contrast to vascular endothelial growth factor, which is required for proliferation and survival of endothelial cells, FGF2 does not seem to be essential since the *Fgf2* knockout is not lethal. Therefore, the precise genetic and physiological roles of FGF2 for capillary development and adaptation remain to be determined. Here we show that myocardial capillary supply is normal at birth, but significantly reduced by ~25% in adult *Fgf2*^{+/-} and *Fgf2*^{-/-} mice as compared with wild-type littermates. In contrast, after induction of myocardial hypertrophy by continuous infusion of angiotensin II (ANG II) for 6 days marked capillary growth was seen in both *Fgf2*^{+/-} and *Fgf2*^{-/-} mice, but not in wild-type littermates. These data demonstrate that two intact *Fgf2* genes are necessary for normal capillary development after birth, whereas FGF2 seems to be dispensable for adaptive myocardial capillary growth in the adult mouse.

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Basic fibroblast growth factor (FGF2) belongs to the family of heparin binding growth factors (FGFs) which are widely distributed in richly vascularized tissues (brain, kidney, retina, adrenal glands, heart, placenta).^{1,2} FGF2 exerts various physiological roles in vascular development^{3,4} and control of blood pressure.^{5–7} It is also involved in cerebral cortex development and function,^{5,8,9} bone formation¹⁰ and cardiac growth.¹¹

Together with acidic fibroblast growth factor (FGF1) and vascular endothelial growth factor (VEGF) FGF2 is involved in angiogenesis^{3,4,12,13} particularly during growth of collaterals in response to hypoxia and to myocardial ischemia.^{14–18} FGF2 is expressed in vascular smooth muscle cells,¹⁹ endothelial cells²⁰ and postmitotic cardiomyocytes;²¹

it stimulates proliferation of endothelial cells, vascular smooth muscle cells and fibroblasts.^{19–21} It is chemotactic for endothelial cells *in vitro*, thus inducing capillary tube formation and regulating blood vessel growth.²⁰

FGF2 has been isolated from and localized in the heart.^{21–23} Increased mechanical load to the heart leads to FGF2 release from cardiomyocytes, thus inducing adaptive myocardial growth.²⁴ *In vitro*, FGF2 increased DNA synthesis and proliferation of neonatal rat cardiac myocytes indicating an important role for myocyte growth and differentiation.²⁵

Although it is clear that FGF2 is involved in the capillary response to ischemia, it has not been determined whether FGF2 is necessary for capillarogenesis during physiological growth and/or for adaptive capillary growth during cardiac hypertrophy. In contrast to VEGF, which is required for proliferation and survival of endothelial cells,^{26–29} FGF2 does not seem to be essential since the *Fgf2* knockout is not lethal.^{5,7–9}

In the present study, two experimental strategies were used in order to investigate the role of

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FGF2 for normal capillary development and adaptation during cardiac hypertrophy. In a combined developmental and physiological approach myocardial capillary supply was analyzed using stereological techniques in adult heterozygous and homozygous FGF2-deficient mice and in wild-type littermates under baseline conditions and after 6 days of continuous angiotensin II (ANG II) infusion to induce myocardial hypertrophy. Specifically, we investigated (i) whether adult FGF2-deficient mice have a distinct cardiac phenotype at the microscopical level, that is, cardiac capillaries, arteries, and interstitial tissue, and (ii) whether capillary adaptation in response to myocardial hypertrophy is altered by a reduced formation of FGF2.

Materials and methods

Generation of *Fgf2*^{+/-} and *Fgf2*^{-/-}

Male and female *Fgf2*^{-/-} mice were generated through gene targeting in R1-ES cells as described in detail previously.⁵ Briefly, mice of mixed C57BL/6J × 129/Sv background were generated as follows: chimeras (ES cells of 129/sv background) were crossed to C57BL/6J females and their heterozygous (+/-) F1 progeny was crossed to generate all three genotypes. These mice and subsequent offspring were used for all studies. Genotyping was carried out in every animal either by PCR using specific oligonucleotides or by Southern blotting. FGF2 serum levels were decreased by 50% in *Fgf2*^{+/-} mice and they were zero in *Fgf2*^{-/-} animals.⁵ The cardiovascular studies were performed in 4–5 months old sibling animals.

In a separate experiment myocardial capillarization was analyzed in newborn wild-type, *Fgf2*^{+/-}, and *Fgf2*^{-/-} mice on postnatal day 1. All mice were re-genotyped after the experiments using either PCR or Southern blotting.

ANG II Infusion

The effect of a continuous ANG II infusion (2 ng/g body weight/min for 6 days, via osmotic minipump,

Alzet, model 1007 D) on myocardial growth and on capillary adaptation to myocardial hypertrophy was investigated in wild-type, *Fgf2*^{+/-} and *Fgf2*^{-/-} mice ($n = 6-3$ animals per group). The dose of ANG II was adjusted to induce an increase in blood pressure of 20 mmHg in normal mice as determined in pilot experiments and previous studies.⁵ On the day of recording, mice were instrumented with a carotid artery catheter during general anesthesia (fentanyl 0.03 µg/g body weight, fluanisone 10 µg/g body weight) and midazolam 10 µg/g body weight) and mean blood pressure and heart rate were measured in awake, resting mice (see *Dono et al*⁵). Infusion or overproduction of ANG II is known to induce myocardial hypertrophy by an increase in blood pressure and thus an increased myocardial work load as well as by direct growth stimulatory effects on cardiac myocytes, interstitial cells and vascular smooth muscle cells.³⁰

Cardiovascular Studies

Implantation of carotid catheters (polyethylene catheter with outer diameter of 400 µm), measurement of mean intra-arterial blood pressure and heart rate were performed as described previously.⁵ All animal experiments were approved by the local authorities and performed according to the EMBL guidelines on animal experimentation (for animal numbers per group see Tables 1 and 2).

Table 1 Blood pressure, body weight, heart weight and left ventricular weight in untreated wild-type, *Fgf2*^{+/-} and *Fgf2*^{-/-} mice

Groups	Mean arterial blood pressure (mmHg)	Body weight (g)	Heart weight (mg)	Left ventricular weight (mg)
Wild-type (n = 10)	124 ± 4	25.4 ± 0.9	133 ± 6	105 ± 6
<i>Fgf2</i> ^{+/-} (n = 11)	133 ± 4	26.2 ± 0.5	134 ± 2	105 ± 2
<i>Fgf2</i> ^{-/-} (n = 13)	118 ± 3	26.8 ± 0.8	141 ± 5	110 ± 6
ANOVA	NS	NS	NS	NS

NS = not significant.
Data are mean ± s.e.m.

Table 2 Blood pressure and cardiac phenotype in response to ANG II infusion in wild-type, *Fgf2*^{+/-} and *Fgf2*^{-/-} mice

Groups	Mean arterial blood pressure (mmHg)	Left ventricular weight (mg/g)	Myocyte diameter (µm)	Volume density of interstitial tissue (%)	Length density of capillaries (mm/mm ³)	Wall thickness of arteries (µm)	Wall-lumen ratio of arteries (µm/µm)
Wild-type ANG II (n = 6)	144 ± 5	4.85 ± 0.04	18.7 ± 0.49	3.48 ± 0.86	3521 ± 290	2.80 ± 0.18	0.09 ± 0.004
<i>Fgf2</i> ^{+/-} ANG II (n = 12)	151 ± 4	5.49 ± 0.26	20.3 ± 0.59	4.47 ± 0.64	3373 ± 289	2.80 ± 0.13	0.11 ± 0.009
<i>Fgf2</i> ^{-/-} ANG II (n = 13)	146 ± 4	5.26 ± 0.24	19.8 ± 0.23	4.70 ± 0.53	3933 ± 232	3.62 ± 0.33	0.09 ± 0.002
ANOVA	NS	NS	NS	NS	NS	NS	NS

NS = not significant.
Data are mean ± s.e.m.

Analysis of Myocardial Capillarization in Newborn Wild-Type, Fgf2^{+/-} and FGF2^{-/-} Mice

To investigate myocardial capillarization in newborn mice an immunohistochemical approach was used since perfusion fixation and orientator technique cannot be used in newborn mice. For capillary staining a CD34 rat polyclonal antibody (BD Pharmingen, concentration 1:50) was used. Pretreatment was performed using a pressure cooker for 5 min in preheated target retrieval solution pH 6.0 (Dako Cytomation, Denmark). The expression of CD34 was analyzed on paraffin sections by computer-aided planimetry in a blinded fashion, that is, the observer was unaware of the group the specimen belonged to. Three to five digital images of each sample ($n = 10\text{--}15$ per group) were obtained and quantitatively analyzed by analySIS (Soft Imaging System GmbH) using standardized settings. The mean number of CD34-positive capillary profiles were counted in the entire tissue sections and expressed as capillaries per mm² tissue (magnification $\times 200$).

Analysis of Myocardial Capillarization after ANG II Infusion

At the end of the experiment, the abdominal aorta was catheterized under general anesthesia (see above). Then, retrograde vascular perfusion was performed at a controlled pressure as described in detail.³¹ First, 10% dextran with procain hydrochloride was used for 2 min to relax the vasculature, then 3% glutaraldehyde was used for 5 min as fixative. After the perfusion, the heart of each animal was taken out for determination of weight and volume. In all, 12 small pieces ($2 \times 2 \times 2$ mm³) of the left ventricle including the septum were sampled according to a modification of the orientator method,^{31,32} and embedded in Epon-Araldite. This technique was developed to determine the capillary number per area and to allow a simple estimation of capillary length density (by multiplying capillary density by 2) as a three-dimensional parameter of myocardial capillarization.^{31,32} It has been used for determination of myocardial capillary supply in several rat and mice models. Semithin sections (0.8 μ m) were cut and stained with methylene-blue/basic fuchsin and examined using light microscopy with oil immersion and phase contrast at a magnification of 1000:1. In several animals per group ultrathin sections were cut, stained with lead citrate and uranyl acetate and qualitatively investigated using a Zeiss EM 10 electron microscope (Zeiss Co., Oberkochen, Germany) at various magnifications.

Quantitative Stereology

All investigations were performed in a blinded manner, that is, the observer was unaware of the experimental protocol. Uniformly random sampling was achieved by preparing a set of equidistant slices of the left ventricle and the interventricular septum

with a random start. Three slices of the left ventricle were selected by area-weighted sampling and processed according to the orientator method.^{31,32} Stereological analysis was performed on 12 random samples of differently orientated sections of the left ventricular myocardium including the inter ventricular septum.

In brief, the length density (L_V) of capillaries, that is, the length of capillaries per unit tissue volume, a three-dimensional parameter of myocardial capillary supply, and the volume density (V_V) of cardiac nonvascular interstitium, that is, the volume of nonvascular interstitial cells and fibers per unit myocardial tissue volume (excluding endothelial cells), were measured in eight systematically subsampled areas per section using a Zeiss eyepiece with 100 points for point counting. In all, 12 orientator sections per animal (a total of 96 test areas per animal) were investigated resulting in a single mean value per heart. Volume density (V_V) was obtained according to the equation $P_P = V_V$ (with P_P is point density) and length density (L_V) was determined using the equation $L_V = 2Q_A$ (where Q_A is area density, ie the number of capillary transects per area of myocardial reference tissue). Reference volume was the total myocardial tissue exclusive of noncapillary vessels. Total length of capillaries per heart (L_{captot}) was calculated using the formula: $L_{\text{captot}} = L_V \times V$ with $V = m/\delta$ and $\delta = 1.04$ g/cm³ (Törnig *et al*³¹). Intercapillary distance (ICD) was calculated according to a modification of the formula of Henquel and Honig³³ as $\text{ICD} = (\sqrt[3]{4 \times Q_A/\sqrt{3}}):2$.

In our hands the method of capillary counting on semithin sections was superior to all other methods (ie immunohistochemical staining of capillaries using factor VIII or CD 31) if adequate perfusion fixation was used to open the capillaries.^{31,32} We had shown previously that counting of capillaries is very reliable and reproducible on semithin sections since plastic embedding does not lead to tissue shrinkage and capillaries can easily be identified and counted.³¹ In the present study, using electron microscopy on ultrathin sections we could exclude collapse of myocardial capillaries after perfusion fixation.

Statistics

Data are given as mean \pm standard error (s.e.m.). After testing for normality, one-way ANOVA or Kruskal–Wallis test, respectively, were used for analysis of variance. Significance between groups was assessed using Duncan's multiple range test. Results were considered significant when probability of error (P) was less than 0.05.

Results

Effect of Disruption of the Fgf2 Gene on Myocardial Architecture and Capillarization in Adult Mice

At the end of the experiment body weights were similar in all experimental groups as were absolute

and relative heart weights and left ventricular weights in the three untreated groups (Table 1).

Mean arterial blood pressure tended to be lower in *Fgf2*^{-/-} than in *Fgf2*^{+/-} mice and wild-type littermates. The difference, however, did not reach statistical significance. Also, there was no correlation of blood pressures with heart weights or left ventricular weights, which tended to be higher in *Fgf2*^{-/-} mice (Table 1).

Figure 1 shows that the number of capillary profiles per area myocardium is markedly lower in *Fgf2*^{+/-} (Figure 1b) and *Fgf2*^{-/-} (Figure 1c) mice compared to wild-type animals (Figure 1a). Figure 2 indicates that three-dimensional myocardial capillary supply (assessed as capillary length density (L_V), that is, capillary length per volume myocardium) was significantly lower in *Fgf2*^{+/-} (3228 ± 245 mm/mm³) and *Fgf2*^{-/-} mice (3385 ± 241 mm/mm³), respectively, than in *Fgf2*^{+/+} littermates (4403 ± 290 mm/mm³, $P < 0.05$, Figures 1 and 2). There were no differences in capillary length density between male and female animals of the same genotype.

Total capillary length per heart (L_{captot}) was also significantly lower in *Fgf2*^{+/-} (3.23 ± 0.22 m) and *Fgf2*^{-/-} (3.50 ± 0.20 m) mice than in wild-type controls (4.57 ± 0.38 m, $P < 0.05$). In parallel with the decrease in capillary length density, mean ICD (ie the distance between the capillary and the center of the adjacent myocyte) increased. It was significantly higher in *Fgf2*^{+/-} (19.4 ± 2.67 μ m) and *Fgf2*^{-/-} (18.9 ± 2.66 μ m) than in wild-type animals (16.0 ± 2.65 μ m, $P < 0.05$).

In contrast, mean myocyte diameter was not significantly different between wild-type mice (17.1 ± 0.33 μ m) and FGF2-deficient mice (17.8 ± 0.35 μ m in *Fgf2*^{+/-} and 17.7 ± 0.27 μ m in *Fgf2*^{-/-}). The same was true for wall thickness and wall to lumen ratio of intramyocardial arteries and volume density of myocardial interstitial tissue ($V_{V_{\text{int}}}$) (*Fgf2*^{+/+}: $1.69 \pm 0.18\%$, *Fgf2*^{+/-}: $1.91 \pm 0.18\%$, *Fgf2*^{-/-}: $2.50 \pm 0.23\%$, $P > 0.05$). The number of intramyocardial arteries per area myocardium was not different between the genotypes indicating a normal representation of arterioles in mice lacking FGF2 compared to wild-type controls.

The ultrastructure of myocytes and capillaries was studied by qualitative electron microscopy in several randomly chosen animals per group ($n = 2-4$ per group). The findings are illustrated in Figure 3a-c. Electron microscopical investigations showed no structural abnormalities of cardiomyocytes and capillaries in the heart of *Fgf2*^{+/-} (Figure 3b) or *Fgf2*^{-/-} (Figure 3c) mice compared to wild-type animals (Figure 3a).

In order to determine, whether the reduced myocardial capillarization observed in adult FGF2-deficient mice results from an altered embryonic development, myocardial capillary supply was assessed in newborn mice by immunohistochemical staining of CD34 as the number of capillary profiles per area myocardium (Q_A). No significant differ-

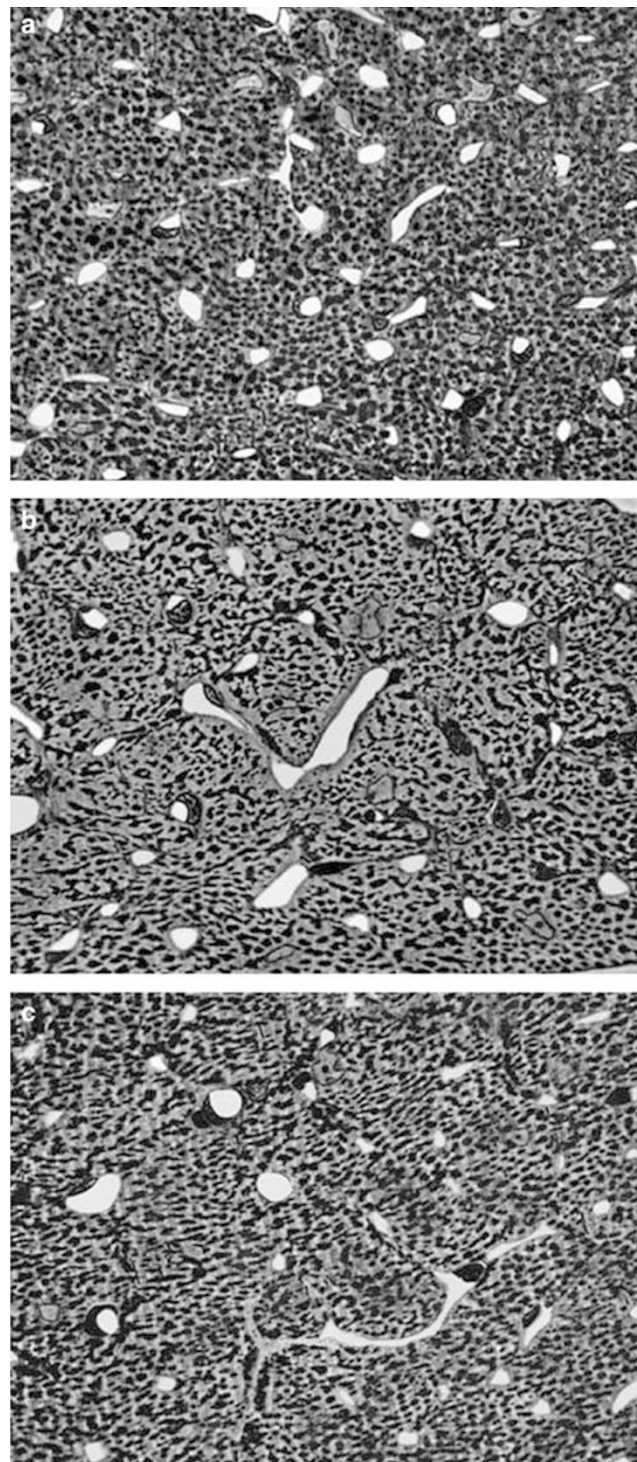


Figure 1 Reduced myocardial capillarization in *Fgf2*^{+/-} and *Fgf2*^{-/-} mice. (a-c) Light microscopical aspect of myocardial capillary supply in wild-type (a), *Fgf2*^{+/-} (b), and *Fgf2*^{-/-} (c) mice. The number of capillary profiles (white structures) per area myocardium is markedly lower in *Fgf2*^{+/-} (b) and *Fgf2*^{-/-} (c) mice compared to wild-type mice (a). Semithin section, magnification: 1:500.

ences were observed between wild-type, *Fgf2*^{+/-} and *Fgf2*^{-/-} mice. The number of capillary profiles per area myocardium ($1/\text{mm}^2$) was 1969 ± 377 in

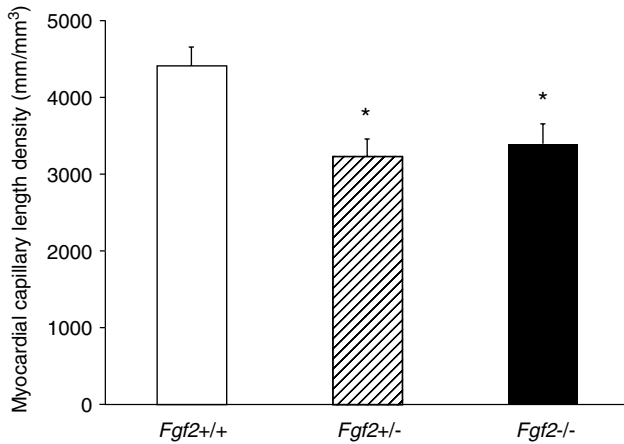


Figure 2 Statistical comparison of capillary length densities as an index of myocardial capillary supply in untreated animals (mean \pm s.e.m.). Capillary length density is significantly lower in *Fgf2*^{+/-} and *Fgf2*^{-/-} than in wild-type mice. *: $P < 0.05$.

wild-type, 2051 ± 109 in *Fgf2*^{+/-}, and 2159 ± 545 in *Fgf2*^{-/-} mice, respectively.

In summary, these data document that two *Fgf2* genes are essential for normal postnatal capillary supply and physiological capillary growth and that loss of one or both alleles results in a substantial reduction (-25%) of myocardial capillary length density as a parameter of three-dimensional capillary distribution in adult mice.

Response to Continuous ANG II Infusion

In order to investigate, whether capillary adaptation to myocardial hypertrophy is altered in *Fgf2*^{+/-} or *Fgf2*^{-/-} mice compared to wild-type animals, ANG II was infused for 6 days.

As expected ANG II infusion led to a significant ($P < 0.05$) increase in mean arterial blood pressure (Table 2), heart weight (data not shown) and relative left ventricular weight, that is, left ventricular weight/body weight ratio (Table 2) in all ANG II-treated groups compared to untreated animals of the same genotype (for comparison see Table 1).

Figure 4 illustrates the increase in relative left ventricular weight after ANG II infusion for each experimental group compared to the untreated animals of the same genotype. Cardiac hypertrophy was also documented at the cellular level by a significant ($P < 0.05$) increase in mean myocyte diameter in all three groups after ANG II treatment (Table 2) compared to untreated animals of the same genotype.

In parallel, marked expansion of nonvascular interstitial tissue (approximately $+100\%$ compared to baseline) developed in all groups indicating that FGF2 is not necessary for interstitial myocardial activation and expansion, respectively (Table 2). Figure 5a–c illustrate focal activation of cardiac fibroblasts and a mild increase in interstitial fibrous tissue in wild-type, *Fgf2*^{+/-} and *Fgf2*^{-/-} mice after ANG II infusion (compared to untreated animals of the same genotype in Figure 1a–c). The results of a

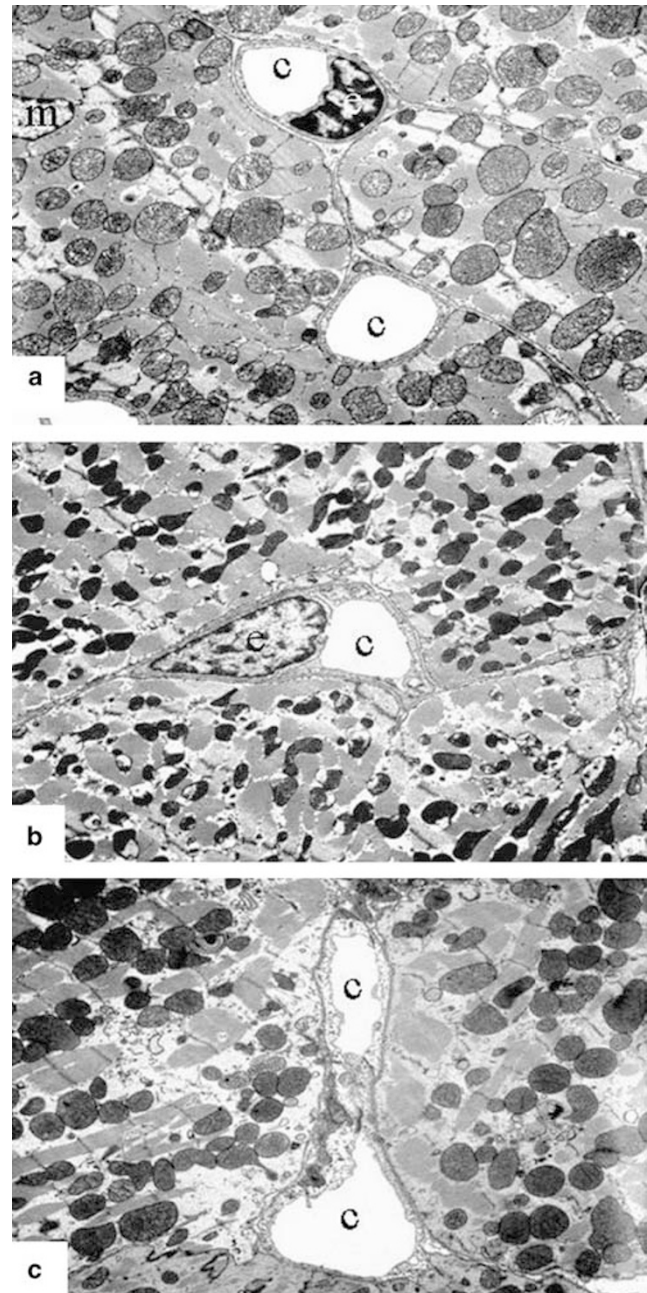


Figure 3 Ultrastructure of the myocardium in wild-type (a), *Fgf2*^{+/-} (b), and *Fgf2*^{-/-} (c) mice. There are no structural alterations of cardiomyocytes (m) or capillaries (c) in *Fgf2*^{+/-} (b) and *Fgf2*^{-/-} (c) compared to wild type (a). Magnification: 1:6.000.

quantitative analysis of volume density of cardiac interstitial tissue are depicted in Figure 6. After ANG II infusion the percentage of interstitial tissue was significantly ($P < 0.05$) higher in all groups compared to untreated animals.

Wall thickness and wall to lumen ratio of intramyocardial arteries were similar in all experimental groups irrespective of ANG II treatment (Table 2). In contrast, the length density of cardiac capillaries was significantly lower in ANG II-treated wild type (3521 ± 290 mm/mm³) than in untreated

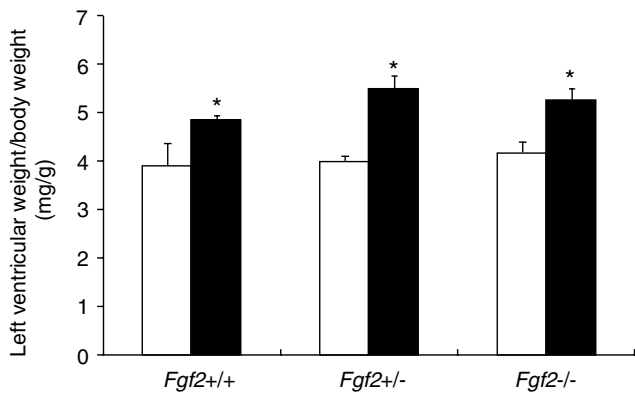


Figure 4 Myocardial hypertrophy in response to continuous ANG II infusion. Left ventricular weight/body weight ratio (mg/g, mean \pm s.e.m.). ANG II induced a significant increase in relative left ventricular weight in all three experimental groups compared to untreated animals of the same genotype. Open bars: no ANG II infusion; black bars: with ANG II infusion, *: $P < 0.05$.

wild-type mice (4403 ± 290 mm/mm³, $P < 0.05$). In contrast, in *Fgf2*^{+/-} and *Fgf2*^{-/-} mice ANG II treatment caused no further decrease in capillary length density (3373 ± 289 and 3933 ± 232 mm/mm³ in ANG II-treated mice compared to 3230 ± 245 and 3500 ± 241 mm/mm³ in untreated animals, respectively). Mean ICD was comparable in all three ANG II-treated groups (18.9 ± 2.21 in *Fgf2*^{+/+} mice, 18.9 ± 3.25 in *Fgf2*^{+/-} mice and 17.4 ± 1.77 in *Fgf2*^{-/-} mice). Figure 7 shows that the total length of capillaries per heart ($L_{\text{cap,tot}}$) was significantly increased in ANG II-infused *Fgf2*^{+/-} and *Fgf2*^{-/-} mice compared to untreated animals.

In summary, these findings demonstrate marked capillary growth in FGF2-deficient mice in response to ANG II-induced myocardial hypertrophy.

Discussion

The present results document that in adult mice even a loss-of-function mutation of a single *Fgf2* gene leads to a substantial myocardial capillary deficit (-25%). Since myocardial capillarization was normal at birth in FGF2-deficient mice, this finding indicates that high levels of FGF2 are required for a normal postnatal development of cardiac capillaries. The capillary deficit appears to be specific for the heart since capillary supply of other organs, for example, the kidney (data not shown), was not impaired. Moreover, the deficit appears to be strictly limited to capillaries since we did not detect any reduction of myocardial arterioles.

FGF2 serum levels are reduced to 50% in *Fgf2*^{+/-} and to zero in *Fgf2*^{-/-} mice.⁵ Although a quantitative relation between FGF2 serum levels and myocardial capillary supply has not yet been demonstrated directly, the present findings strongly suggest that during postnatal cardiac development high levels of FGF2 are required for a normal myocardial capillarization. Interestingly, a haploinsufficiency has also been found for loss-of-function mutations of the

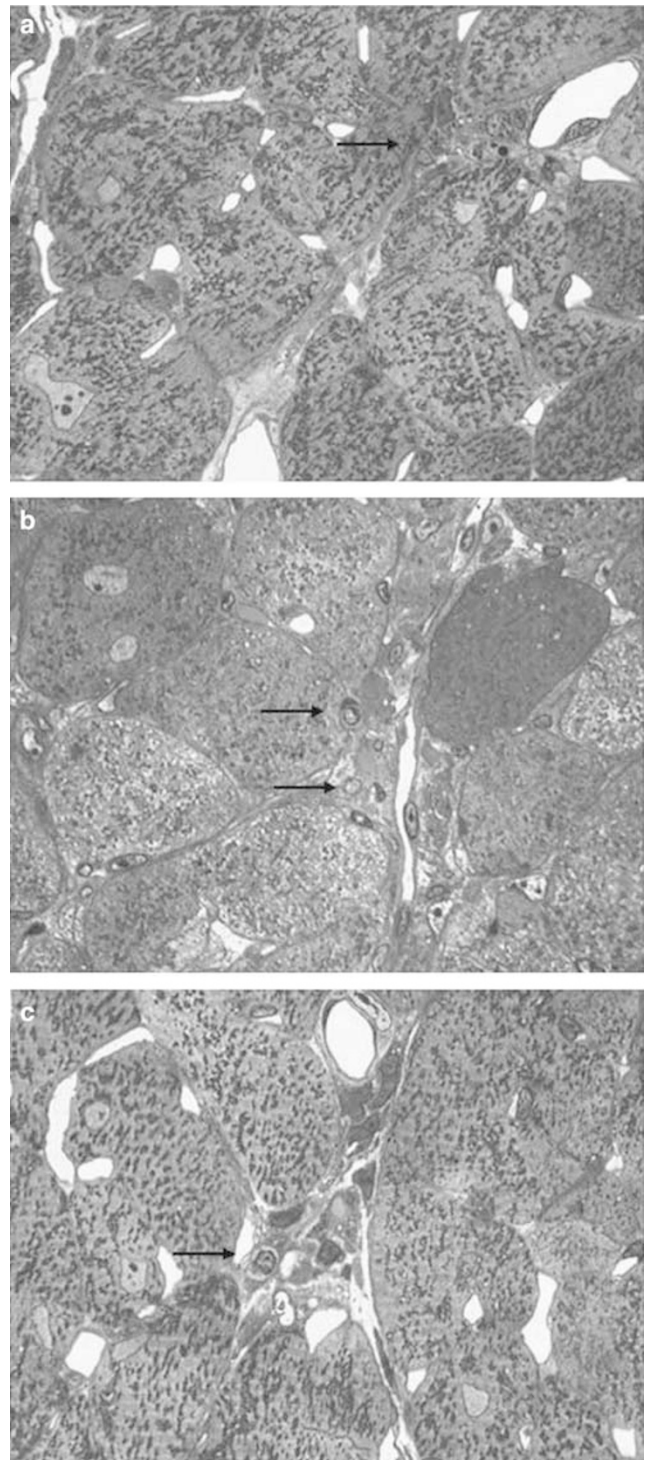


Figure 5 Effect of continuous ANG II infusion on cardiac nonvascular interstitial tissue. (a–c) Light microscopical aspect of the myocardium after ANG II infusion. An increase in cardiac interstitial tissue (ie fibroblasts indicated by arrows and interstitial fibers) was seen in wild-type (a), *Fgf2*^{+/-} (b), and *Fgf2*^{-/-} (c) mice after continuous ANG II infusion compared to the normal morphology in untreated animals (see Figure 1). Magnification: 1:800.

gene encoding interleukin-2 (IL-2)³⁴ which maps physically to a closely related site of *Fgf2* on chromosome 4 (q25–27, see *Dono et al*⁵).

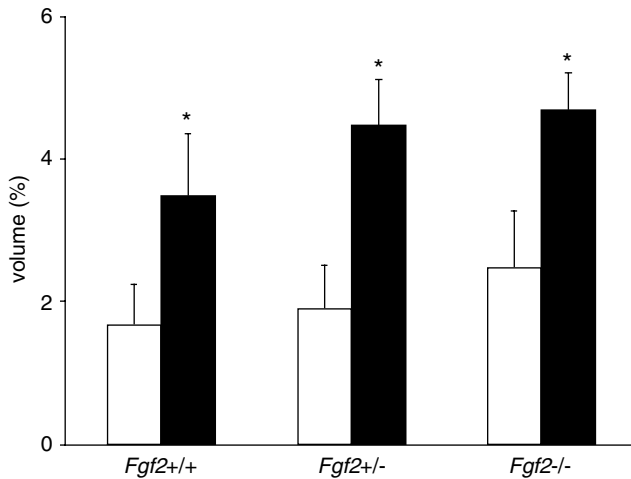


Figure 6 Statistical comparison of volume density of myocardial nonvascular interstitial tissue as an index of myocardial fibrosis after continuous ANG II infusion (mean \pm s.e.m.). The volume percentage of interstitial tissue was significantly higher after continuous ANG II infusion in all groups compared to untreated animals of the same genotype ($P < 0.05$). Open bars: untreated animals, filled: continuous ANG II infusion, *: $P < 0.05$.

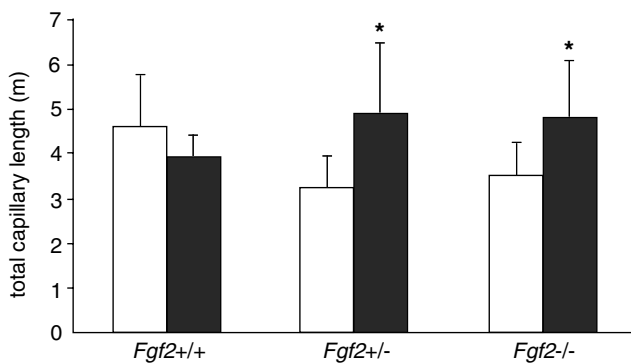


Figure 7 Capillary adaptation in response to continuous ANG II infusion. Total capillary length (m, mean \pm s.e.m.). Total capillary length after continuous ANG II infusion was significantly higher in *Fgf2*^{+/-} and *Fgf2*^{-/-} compared to untreated animals of the same genotype. Open bars: untreated animals, filled: continuous ANG II infusion, *: $P < 0.05$.

FGF2 expression is markedly enhanced during the early period of fetal coronary vascularization and during the early neonatal period when capillary proliferation is particularly high.^{4,35} In the mouse, heart transition to the adult stage is associated with a decrease of FGF2 receptor 1 mRNA expression.³⁶ In parallel, cardiac growth is regulated by receptor-coupled FGF signaling only during tubular stages of cardiogenesis and becomes FGF independent after the second week of embryogenesis.³⁷ Thus, FGF2 signaling appears to be particularly activated during early stages of myocardial capillarization. In contrast, FGF2 has only a minor influence on existing vessels as documented in dogs with chronic single-vessel coronary occlusion where administration of FGF2 did not have any effect on mature collateral vessels.³⁸ Similarly, in *Fgf2*^{-/-} mice with targeted disruption of the *Fgf2* gene vascular growth in the

mouse ischemic hindlimb was not different from wild-type mice indicating that endogenous FGF2 is not required for revascularization in the setting of peripheral ischemia.³⁹ Using neutralizing antibodies to VEGF and FGF2 and comparable stereological techniques in perfusion-fixed rat hearts different roles of VEGF and FGF2 in early postnatal coronary angiogenesis were confirmed.⁴⁰

A reduction of capillary length density has been found in several experimental models of cardiac hypertrophy.⁴¹ It is particularly pronounced in left ventricular hypertrophy due to renal insufficiency and can be documented in experimental renal failure in rats^{31,42} as well as in uremic patients.⁴³ A reduction of myocardial capillary length density and intercapillary diffusion distance by $\sim 25\%$, as found for *Fgf2*^{+/-} and *Fgf2*^{-/-} mice in the present study, does not appear to alter heart function in resting rats with subtotal nephrectomy *per se* because of the physiological coronary reserve (unpublished observation). A normal cardiac oxygen supply at rest would also be consistent with the absence of an expansion of myocardial interstitial tissue or myocardial fibrosis in *Fgf2*^{+/-} and *Fgf2*^{-/-} mice (Figure 1), which are histopathological signs of chronic myocardial ischemia. In view of this finding, it is also very unlikely that the capillary changes observed in *Fgf2*^{+/-} and *Fgf2*^{-/-} mice are due to a possible effect of FGF2 on interstitial tissue. In addition, normal cardiac contractility was found in *Fgf2*^{-/-} mice.¹¹ Cardiac oxygen supply, however, may be severely impaired if additional metabolic stress factors like an increased heart rate or cardiac hypertrophy are present.⁴⁴ Such a functional ischemic cardiomyopathy has been recently described to accompany an abnormal capillarization with up to 75% reduction of capillary density in VEGF164 and VEGF188 knockout mice.⁴⁵

In order to investigate the potential role of FGF2 in capillary adaptation in response to myocardial hypertrophy mice received hypertensive doses of ANG II via a continuous infusion. In all genotypes ANG II increased blood pressure and induced myocardial hypertrophy of similar degrees. In contrast to wild-type animals, total capillary length per heart increased significantly in mice deficient for FGF2 (*Fgf2*^{+/-} and *Fgf2*^{-/-}) in response to ANG II treatment. These findings demonstrate that FGF2 is not essential for adaptive capillary growth. The mechanism(s) leading to the marked capillary growth in response to ANG II-induced myocardial hypertrophy in FGF2-deficient mice remains to be elucidated. It is of note, however, that in a recent study on the cardiac transcriptional response to acute and chronic ANG II treatment in mice Larkin *et al*⁴⁶ found that the response to ANG II infusion in mice is somewhat inconsistent and in general not as marked as in rats. After acute ANG II infusions an increase in blood pressure and related changes in myocardial gene expression, but not in heart weight, was found.⁴⁶ Among the genes differently expressed

after acute ANG II infusion were those involved in growth regulation, oxidative stress response and amino-acid metabolism.

It seems very unlikely that the physiological role of FGF2 in the adult organism would be inhibition of adaptive capillary growth. Thus, in the absence of FGF2 other growth factors may be upregulated which take over the role normally played by FGF2, thus facilitating capillary growth. In particular, major roles were shown for VEGF, angiotensin and Tie2 in cardiac vasculogenesis and angiogenesis after ANG II infusion.⁴⁷ In dogs and pigs VEGF infusion stimulated collateral vessel development in the ischemic myocardium resulting in improved cardiac global and regional function.^{48,49} It is also well known that the various FGF family members can substitute for each other in their respective functions.¹³ Thus, it seems possible that in the *Fgf2*^{-/-} mouse some functions of FGF2 are taken over by other family members. This notion is supported by the observation that the FGF-receptor-deficient mouse has a distinct cardiac phenotype.³⁷

The present finding of a preserved adaptive myocardial growth in *Fgf2*^{-/-} mice seems to be in contrast to the observation that cardiac hypertrophy induced by transverse aortic coarctation was severely attenuated in *Fgf2*^{-/-} mice.¹¹ This discrepancy may be explained by the fact that ANG II infusion and transverse aortic coarctation induce cardiac hypertrophy by activation of different intracellular-signaling cascades,^{46,50} partly because pressure overload by thoracic aortic coarctation does not activate the circulating renin-angiotensin system.⁵¹

The present study indicates that two intact alleles of *Fgf2* are required for a normal postnatal cardiac vascularization on the capillary level, but does not appear to be essential for adaptive myocardial capillary growth in the adult mouse. FGF2-deficient adult mice represent a genetic animal model with a specific reduction of myocardial capillaries that may be very useful in cardiovascular research, for example, for the investigation of adaptive and reparative processes in chronic myocardial ischemia or myocardial infarction.

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