

Hepatocyte RXR alpha deletion in mice leads to inhibition of angiogenesis

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Abstract To investigate the effect of RXR α deficiency in liver on angiogenesis, hepatocyte RXR α -deficient and control wild-type mice were fed either standard or high-fat diet (HF) for 7 weeks. In the 6th week of feeding, Matrigel model of in vivo angiogenesis was applied. Matrigel plug infiltrating cells were then used for visualization of vessels (CD31 staining) as well as for gene expression analysis. HF diet appeared to decrease angiogenesis in hRXR α -deficient mice. Genes related to angiogenesis (*Nos3*, *Kdr*) were down-regulated, whereas genes connected with adipogenesis (*Cebpb*, *Srebf1*), apoptosis (*Gzmb*, *Bcl2*) and proinflammatory pathway (*NfkB*, *Tnfa*) were up-regulated by HF diet in hRXR α -deficient mice in the microarray study. Our results suggest that impaired fatty acid metabolism in liver (hRXR α -deficient mice) leads to impaired angiogenesis due to lipotoxicity and promotion of adipogenesis.

Keywords Adipogenesis · Angiogenesis · FFA · Retinoid X receptor

Introduction

Angiogenesis, the process of the new blood vessels formation, is a combined result of tissue growth and tissue remodeling, which also includes adipose tissue [2, 9]. Several factors such as vascular endothelial growth factor (VEGF), basic fibroblast growth factor (bFGF), nitric oxide (NO) as well as insulin-like growth factor 1 (IGF-1), insulin, adipokines (leptin, adiponectin) and several others play important roles in the angiogenic response [8]. The substances listed above regulate the most important steps in angiogenesis like proliferation, migration, homing and differentiation of the vascular wall progenitor cells. Our previous studies indicated that beta-carotene exerted proangiogenic activity by stimulating endothelial cell migration and homing [4]. The retinoic acid derived from beta-carotene cooperates with free fatty acids (FFA) and its metabolites in lipid metabolism by regulating heterodimer PPAR/RXR activation. Thus, targeted deletion of RXR α gene in hepatocytes leads to dysfunction of enzymes participating in cholesterol, FFA, bile acids, steroids and xenobiotics metabolism [7, 15]. Mice with hepatocytes RXR α deficiency fed a high-fat diet displayed elevated serum triglycerides, increased apo CIII mRNA levels, and higher cholesterol and leptin concentrations in serum [10–12]. The present study was undertaken to define the effect of dyslipidemia induced by RXR alpha deficiency in the liver on angiogenic response in vivo, in mice model.

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Methods

Mice

Mice deficient in hepatocyte RXR α -gene were described elsewhere [11]. The hepatocyte RXR α deletion was

introduced in the mixed genetic background of C57/Bl6, 129/svEvTac and DBA-2 mice [11]. The animals used in the experiments were kindly supplied by Dr. Yu-Jui Yvonne Wan (University of Kansas Medical Center).

Mice were age-matched (males), housed in cages at 22°C with 12/12-h light/dark cycle with free access to food and water. The animals were fed either standard lab chow diet containing 3% fat or high-fat (HF) diet (high saturated fat diet, coconut oil based, 39en% fat) (MP Biomedical Research, USA) for 7 weeks. The study protocol was approved by the local University Ethical Committee (JUMC).

Angiogenesis in vivo

In the 6th week of standard and HF diet, the mice were subcutaneously dorsally injected with Matrigel (Becton Dickinson; 2 × 500 µl) that contained bFGF [25 nM] for 6 days. Matrigel plugs were then removed from the mice. Matrigel infiltrating cells were analyzed for specific gene expression and they were immunohistochemically stained with rat anti-mouse CD31 antibodies specific for platelet/endothelial cell adhesion molecule (anti-PECAM1, Becton Dickinson) [13]. The number of newly formed capillaries (with and without lumen) was counted using a “hot spot” method described previously [14]. The images were analyzed in five fields in three slides taken from different parts of each plug.

Microarray analysis

The differences in angiogenic gene expression in Matrigel infiltrating cells were monitored by means of a microarray analysis (Affymetrix 430A_2 GeneChips), of which results were evaluated with Affymetrix microarray analysis suit. The genes selected for further analysis included the ones with significant differences in signal intensity ($P < 0.05$) and relative change in their expression greater than 1.4-fold.

Statistics

All results were presented as mean values ± standard deviation (SD). The *t* Student's test was applied to determine the significance between analyzed factors; P values lower than 0.05 ($P < 0.05$) were considered to be significant.

Results

Hepatocyte RXR α -deficient mice and their wild-type counterparts were fed regular and HF diet for 6 weeks in

order to elucidate the role of hepatocyte RXR α in angiogenesis under different lipemic conditions. The animals were injected with Matrigel (BD), which was later removed and analyzed for angiogenic response and gene expression in the infiltrated cells. The hRXR α -deficient mice demonstrated a tendency to weaker angiogenic response compared to the wild-type mice when supplied with HF diet (Fig. 1). In addition, high-fat (HF) diet induced down-regulation of a number of angiogenic genes in hepatocyte RXR α -deficient mice as demonstrated by Affymetrix microarray analysis of gene expression in the cells infiltrating Matrigel plugs (Table 1). Gene expression of such growth factor receptors as *Kdr* (vascular endothelial growth factor receptor-2) or *Pdgfrb* (platelet-derived growth factor receptor beta) that induce proliferation and migration of endothelial cells was significantly reduced. Activity of *Nos3* gene (endothelial nitric oxide synthase) that regulates vasodilatation, endothelial cell protection and angiogenesis [1] was also inhibited.

The high-fat diet resulted in up-regulation of pro-apoptotic genes in the Matrigel-infiltrating cells derived from hRXR α -deficient mice. Expression of the genes encoding activators of caspase 3 [*Prf1* (perforin 1), *Gzmb* (granzyme B)], pro-apoptotic *Bcl2L11* (Bcl2 interacting mediator of cell death), NF κ B pathway [*Nfkbia* (nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor), *Nfkbl* (nuclear factor of kappa light polypeptide gene enhancer in B-cells 1)], TNF α receptors [*Tnfrsf21* (tumor necrosis factor receptor superfamily, member 21),

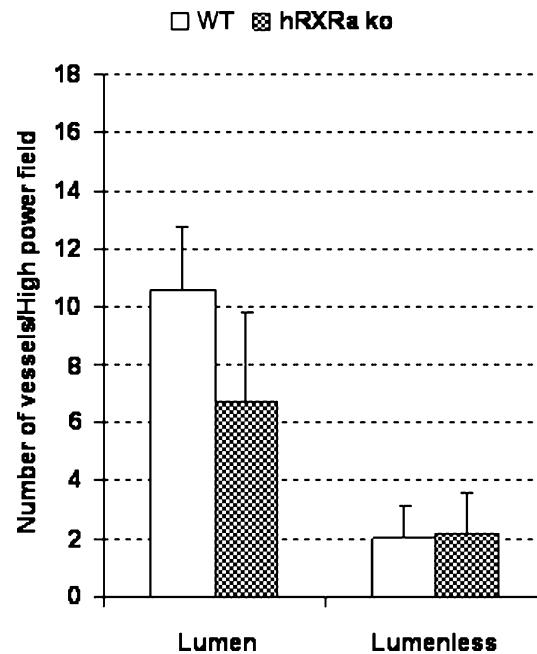


Fig. 1 Number of vessels with lumen and without lumen in Matrigel plug of wild-type and hepatocyte RXR α -deficient (hRXR α ko) mice fed with a high-fat diet. Values are expressed as mean ± SD ($n = 5$)

Table 1 Comparison of microarray changes in relative gene expression (between Matrigel plug cells from mice fed with a standard diet (ST) and matrigel plug cells from mice fed with a high-fat diet (HF)) in RXR α -deficient mice (hRXR α ko) and WT mice

Gene symbol	WT HF vs ST	hRXR α ko HF vs ST
Angiogenesis		
Kdr	nc	−1.87 ^a
Pdgfrb	nc	−1.52 ^a
Nos3	nc	−2.64 ^a
Apoptosis		
Prf1	−4.29 ^a	2.14 ^b
Gzmb	−5.66 ^a	9.85 ^b
Bcl2l11	−1.87 ^a	1.41 ^b
Nfkbie	−2.14 ^a	4 ^b
Nfkbl	−1.32 ^a	1.52 ^b
Nfkbia	−1.52 ^a	1.32 ^b
Tnfrsf21	−1.52 ^a	2 ^b
Tnfrsf1b	−2.14 ^a	1.87 ^b
Casp3	−1.62 ^a	nc
Casp6	−1.74 ^a	nc
Birc5	nc	−1.41 ^a
Birc4	nc	−1.52 ^a
Adipogenesis		
Cebpb	−3.48 ^a	1.74 ^b
Srebf1	nc	1.74 ^b
Adipoq	3.73 ^b	−6.96 ^a
Retn	nc	−6.06 ^a
Adn	7.46 ^b	−13 ^a
Fabp4	1.62 ^b	−4.92 ^a
Lpl	3.03 ^b	nc

nc no change

^a Down-regulated genes^b Up-regulated genes

Tnfrsf1b (tumor necrosis factor receptor superfamily, member 1b)] was increased, while some inhibitors of apoptosis [Birc5 (baculoviral IAP repeat-containing 5), Birc4 (baculoviral IAP repeat-containing 4)] were down-regulated. In contrast, high-fat diet affected the expression of the above pro-apoptotic genes in the Matrigel-populating cells from the wild-type (WT) mice in the opposite way to the mutant cells. Expression of the genes encoding caspases (*Casp3*, *Casp6*), TNF receptors (*Tnfrsf21*, *Tnfrsf1b*) NF κ B pathway (*Nfkbie*, *Nfkbl*), *Bcl2l11*, perforin 1 (*Prf1*) and granzyme B (*Gzmb*) was uniformly decreased by 1.32 (*Nfkbl*) to 5.66 (*Gzmb*)-fold (Table 1).

HF diet also promoted expression of transcription factors important in the early steps of adipogenesis in the Matrigel-infiltrating cells from hRXR α -deficient mice. Genes encoding transcription factors recognized as

essential for adipogenesis [5], such as *Cebpb* (CCAAT/enhancer binding protein β) or *Srebf1* (sterol regulatory element binding factor 1), were up-regulated, whereas markers of differentiated adipocytes that included *Adipoq* (adiponectin), *Retn* (resistin), *Adn* (adipsin) and *Fabp4* (fatty acid binding protein 4) were down-regulated. Surprisingly, in WT control mice, HF diet resulted in decrease of gene expression of *Cebpb* that regulates pre-adipocyte differentiation, while genes associated with adipocyte differentiation, *Adipoq*, *Adn*, *Fabp4*, and *Lpl* (lipoprotein lipase), were up-regulated (Table 1).

Discussion

We demonstrated that mice with hepatocyte RXR alpha deficiency manifest tendency to decreased angiogenic response in the Matrigel model. The number of the blood vessels with lumen was smaller in the Matrigel plugs removed from the mutant mice compared to the wild-type control ones under a high-fat diet. This histopathological observation was confirmed by the gene expression analysis of the cells harvested from the Matrigel plug. Genes critical for angiogenesis, such as *Kdr*, the receptor for VEGF, and *Nos3*, endothelial constitutive nitric oxide synthase important for vascular network formation [1], were clearly down-regulated. Our notion that new vessel formation in the Matrigel plugs from hRXR α -deficient animals was reduced due to inhibition of angiogenic response was also supported by decreased gene expression of integrins, gap junction proteins and matrix remodeling enzymes such as metalloproteinases (data not shown).

We propose that impaired angiogenesis observed in hRXR α -deficient mice is related on one hand to lipotoxicity of FFA, and on the other to activation of adipogenesis. Deficiency of RXR α in liver could lead to increased level of circulating FFA in the blood, which could result in lipotoxicity, endothelial dysfunction and apoptosis of peripheral tissues [3]. It was shown that excess of FFA leads to lipotoxicity of endothelium by increased de novo synthesis of diacylglycerol and activation of protein kinase C (PKC) that initiates a cascade of events leading to stimulation of endothelial superoxide production, inhibition of NOS3 activity and activation of nuclear factor NF- κ B [6]. In microarray study, we observed that HF diet in the matrigel plug of hRXR α -deficient mice resulted in activation of PKC gene expression (data not shown), inhibition of *Nos3* and up-regulation of proinflammatory pathways: NF- κ B and TNF α receptors. Our observation that pro-apoptotic genes are up-regulated under HF diet in the absence of hepatocyte RXR α reinforces the hypothesis that FFA-induced lipotoxicity is the cause of the impaired angiogenic response. Increased caspase activity promoted

by *Prfl*, *Gzmb*, up-regulation of pro-apoptotic protein *Bcl2L11* and inhibition of antiapoptotic genes (*Birc4*, *Birc5*) confirm the induction of apoptosis in the Matrigel infiltrating cells. On the contrary, in control WT mice, HF diet resulted in down-regulation of the expression of pro-apoptotic genes such as: caspases and their activators. Beside down-regulation of the listed proangiogenic factors, HF diet activated early modulators of adipocyte differentiation such as transcription factors: (*Srebf1*, *Cebpb*) in the cells from hRXR α -deficient mice.

In summary, the presented study demonstrates that HF diet leads to biochemical changes that result in impaired angiogenic response in the hepatocyte RXR α -deficient mice. The observed decline in the formation of new blood vessels is due to the parallel activation of the expression of proadipogenic and proapoptotic genes in the cells involved in angiogenesis.

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Conflict of interest statement There is no conflict of interest.

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