

Figure 6. Blockade of nuclear factor- κ B (NF- κ B) signaling prevented vascular senescence and prolonged life span. **A**, Control (○; n=6) and endothelial dominant-negative I κ B α transgenic (E-DN1 κ B; ■; n=8) mice received glucose tolerance tests with a peritoneal glucose load (1.5 g/kg body weight) after a 10-hour fast at 50 weeks of age. **B**, Insulin tolerance tests were performed in an ad libitum-fed state. Data are expressed as percentages of blood glucose levels immediately before intraperitoneal insulin loading (0.75 U/kg body weight) at 50 weeks of age. **C**, Aortas from aged control and E-DN1 κ B mice at 90 weeks of age were stained for senescence-associated β -galactosidase (SA β -gal). SA β -gal-positive areas of aortas from aged control (white bars, n=5) and E-DN1 κ B (black bars, n=5) mice were measured (right). Data are presented as mean \pm SEM. * P <0.05, ** P <0.01 vs control littermate group by 1-way and repeated-measures ANOVA. WT indicates wild type. **D**, Cumulative survival of control (○; n=24) and E-DN1 κ B (■; n=21) mice on a normal chow diet was analyzed by the Kaplan-Meier method. ** P <0.01 by the log-rank test.

be lower in E-DN1 κ B mice, although the differences were not statistically significant (Figure 6A). Insulin tolerance tests revealed better insulin sensitivity in E-DN1 κ B mice (Figure 6B). Thus, blockade of endothelial NF- κ B signaling inhibited the development of age-related insulin resistance.

Furthermore, systolic blood pressure was significantly lower in E-DN1 κ B mice than in wild-type controls of the same age (Figure VIIIA in the online-only Data Supplement).

To determine whether eNOS mediates age-related blood pressure elevation via endothelial NF- κ B signaling, we next crossed E-DN1 κ B mice with eNOS-deficient (Nos3^{-/-}) mice. Whereas endothelial DN1 κ B expression suppressed blood pressure elevation in Nos3^{+/+} mice, eNOS deficiency blunted the inhibitory effects of DN1 κ B on blood pressure elevation (Figure VIIIA in the online-only Data Supplement). These findings indicate involvement of eNOS in protection from age-related hypertension in E-DN1 κ B mice.

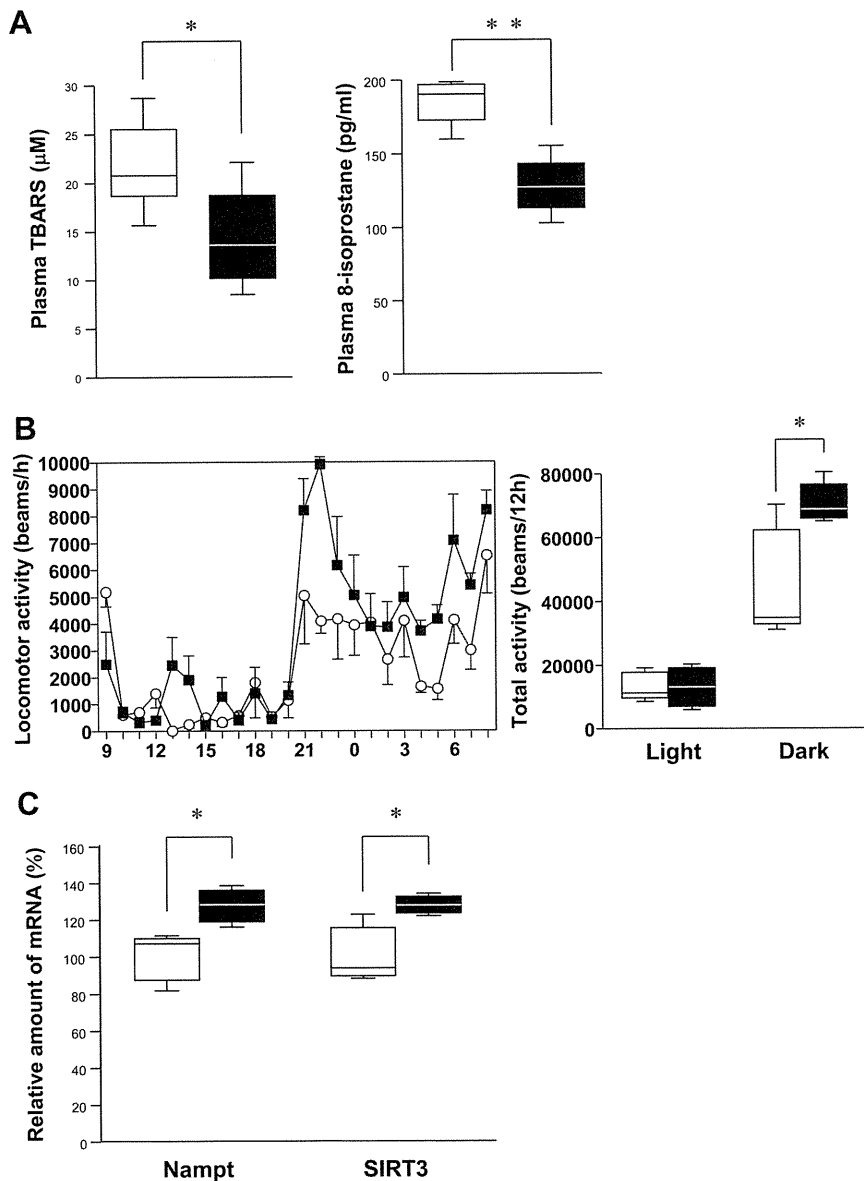


Figure 7. Increments in locomotor activity and upregulation of mitochondrial prosurvival genes in endothelial dominant-negative $\text{I}\kappa\text{B}\alpha$ transgenic (E-DN $\text{I}\kappa\text{B}$) mice. **A**, Plasma thiobarbituric acid-reactive substance (TBARS) and 8-isoprostane concentrations were measured in aged control (white bars, $n=6$) and E-DN $\text{I}\kappa\text{B}$ (black bars, $n=8$) mice at 50 weeks of age. **B**, Locomotor activities of aged control (\circ ; $n=5$) and E-DN $\text{I}\kappa\text{B}$ (\blacksquare ; $n=5$) mice at 50 weeks of age (left). Estimated activities of aged control and E-DN $\text{I}\kappa\text{B}$ mice in 12-hour light and dark periods (right) were calculated. **C**, Aortic gene expression of mitochondrial prosurvival genes in 50-week-old aged control and E-DN $\text{I}\kappa\text{B}$ mice was analyzed by reverse-transcriptase polymerase chain reaction. The relative amounts of mRNA were calculated with β -actin mRNA as the invariant control. Data are presented as means \pm SEM. Namp1 indicates nicotinamide phosphoribosyltransferase $n=5$ in each group. * $P<0.05$, ** $P<0.01$ by 1-way ANOVA.

Blockade of NF- κ B Signaling Prevented Vascular Senescence and Prolonged Life Span in Mice

Next, we analyzed vascular senescence by β -galactosidase staining of the aortas of aged E-DN $\text{I}\kappa\text{B}$ mice at 90 weeks of age because endogenous β -galactosidase activity is reportedly increased in senescent states.²⁰ As shown in Figure 6C, β -galactosidase staining was much weaker in the aortas of E-DN $\text{I}\kappa\text{B}$ mice, suggesting prevention of vascular senescence by endothelial blockade of NF- κ B signaling. These findings prompted us to hypothesize that endothelial NF- κ B signaling affects longevity. Therefore, we analyzed the life spans of these mice. E-DN $\text{I}\kappa\text{B}$ mice and control littermates were fed a standard chow diet ad libitum and maintained in regular housing until death. Intriguingly, E-DN $\text{I}\kappa\text{B}$ mice exhibited significantly prolonged life spans compared with control littermates ($P=0.0095$ by log-rank test; Figure 6D). Thus, blockade of endothelial NF- κ B signaling may prevent age-related metabolic deterioration and vascular senescence and thereby increase longevity.

Similarly in the obesity model, plasma levels of oxidative stress markers were significantly lower in aged E-DN $\text{I}\kappa\text{B}$ mice at 50 weeks of age than in wild-type controls of the same age (Figure 7A). Blood flow in muscles was increased in aged E-DN $\text{I}\kappa\text{B}$ mice compared with wild-type littermates, whereas eNOS deficiency blunted the effects of increased blood flow in aged E-DN $\text{I}\kappa\text{B};\text{Nos}3^{-/-}$ mice (Figure VIII B in the online-only Data Supplement). Furthermore, blockade of endothelial NF- κ B signaling enhanced locomotor activity during the 12-hour dark phase with no significant alterations during the 12-hour light phase (Figure 7B).

Oxidative damage to mitochondria reportedly contributes to aging and various age-related disorders.³⁰ In human subjects, endurance exercise reportedly enhances mitochondrial SIRT3 expression.³¹ Upregulation of nicotinamide phosphoribosyltransferase (Namp1) and SIRT3, which are highly expressed in mitochondria, is linked to life-span extension in the context of caloric restriction.³² Therefore, we next examined the expression of Namp1 and a mitochondrial sirtuin,

SIRT3, in the aortas of 50-week-old E-DN1 κ B and control mice. In E-DN1 κ B mice, both Nampt and SIRT3 were actually upregulated (Figure 7C) despite no food intake differences (Figure VIIA in the online-only Data Supplement). Thus, decreased oxidative stress, enhanced active-phase locomotor activity, and upregulation of mitochondrial prosurvival genes might contribute to the life-span prolongation resulting from NF- κ B signaling blockade in the endothelium.

Discussion

The endothelium forms an interface between vascular structures and blood. Endothelial cells produce and react to a wide variety of mediators, including cytokines, growth factors, vasoactive substances, and chemokines, as well as adhesion molecules. Therefore, in this study, we focused on proinflammatory responses in the endothelium in an effort to elucidate the role of endothelial NF- κ B signaling. We blocked this signaling in vivo by expressing the dominant-negative form of I κ B α in endothelial cells using the transgenic procedure. Although E-DN1 κ B mice displayed no overt phenotypic changes when young and lean, they were protected from the development of both obesity-induced and age-related insulin resistance. Furthermore, intriguingly, E-DN1 κ B mice were also protected from vascular senescence and showed extended longevity.

The mechanisms underlying protection from obesity-induced insulin resistance in E-DN1 κ B mice are likely to involve suppression of macrophage infiltration into adipose tissue. Recent studies have demonstrated that macrophage infiltration into white adipose tissues is increased in obesity, raising levels of proinflammatory cytokines such as TNF- α .^{22,23} Such macrophage infiltration into adipose tissue may be triggered by the interaction between endothelial cells and macrophages via adhesion molecules. The expression of adhesion molecules is reportedly regulated by NF- κ B in the endothelium.³³ In the present study, the expression of vascular adhesion molecules was actually decreased in E-DN1 κ B mice. A series of recent reports have indicated angiogenic factors to be involved in the development of obesity in adipose tissue.³⁴ However, adipose expression of Tie2 and vascular endothelial growth factor did not differ between E-DN1 κ B;Ay/+ mice and control Ay/+ littermates, suggesting minimal effects of endothelial NF- κ B signaling on angiogenesis in adipose tissue. In addition, proinflammatory cytokines secreted by macrophages in adipose tissue may further activate the endothelial NF- κ B pathway, producing a deleterious cycle. Indeed, epididymal fat weight was significantly lower and plasma adiponectin was higher in E-DN1 κ B;Ay/+ mice than in Ay/+ littermates. Therefore, interruption of this deleterious cycle by blockade of endothelial NF- κ B signaling is speculated to contribute to protection from obesity-related chronic inflammation and increased oxidative stress in E-DN1 κ B mice.

Decreased eNOS production may lead to a reduction in microcirculatory blood flow and elevated blood pressure. Endothelium-derived NO is a major vasodilator,³⁵ and constitutive NO production by endothelial cells reportedly inhibits adhesion molecule expression through stabilization of

I κ B.³⁶ On the other hand, in this study, eNOS expression was enhanced in E-DN1 κ B mice, suggesting that endothelial NF- κ B signaling per se negatively regulates eNOS expression. Furthermore, in NOS3^{-/-} mice, endothelial DN1 κ B expression did not significantly suppress age-related hypertension or increase blood flow in muscle, indicating the involvement of eNOS in the antihypertensive effects of endothelial NF- κ B blocking. NO production and NF- κ B activation affect each other in endothelial cells. NF- κ B activation decreases eNOS expression, resulting in decreased NO production and thus further NF- κ B activation, producing a vicious cycle that further decreases microcirculation and increases proinflammatory responses and oxidative stress. The attenuation of microcirculatory blood flow observed in eNOS-deficient mice decreases transcapillary passage of insulin to metabolically active tissues such as muscle, thereby contributing to impairment of insulin action.¹⁸ Negative regulation of eNOS expression by NF- κ B in endothelial cells is thus another important mechanism underlying insulin resistance and hypertension.

Insulin resistance, elevated blood pressure, and increased oxidative stress are commonly observed not only in obese but also in aged states. These age-related metabolic deteriorations were also prevented in E-DN1 κ B mice, in association with increased muscle blood flow and decreased oxidative stress markers. Furthermore, it is noteworthy that E-DN1 κ B mice were protected from vascular senescence and lived longer even under normal chow-fed conditions. At the cellular level, NF- κ B has been implicated in age-dependent induction of cellular senescence.³⁷ However, in this study, blockade of NF- κ B signaling selectively in endothelial cells affected vascular senescence of the whole aorta and the life spans of the model animals. Therefore, intracellular events in the endothelium alone cannot explain these antiaging phenotypes manifesting in the whole body. Relatively early deaths of E-DN1 κ B mice (60–100 weeks of age) were decreased and maximum life span was longer in E-DN1 κ B mice, suggesting that endothelial NF- κ B blockade prevents both fatal morbidities and senescence. Amelioration of insulin resistance and decreased oxidative stress likely contribute to these systemic antiaging phenotypes.

In addition, mitochondrial function may be involved in the underlying mechanism. Mitochondrial dysfunction in muscle promotes the development of insulin resistance in obese²⁹ and aged³⁸ human subjects and contributes to aging and various age-related disorders.³⁰ Mitochondrial dysfunction is also linked to decreased muscle blood flow with aging.³⁹ Because eNOS reportedly modulates mitochondrial biogenesis,²⁶ eNOS upregulation may contribute to enhanced mitochondrial contents in muscles of E-DN1 κ B mice. Furthermore, endurance exercise, which is considered to confer life-span-extending effects, enhances the expression of a mitochondrial sirtuin, SIRT3, in human subjects.³¹ SIRT3 was initially reported to be linked to caloric restriction-induced cell survival.^{40,41} Nampt provides mitochondrial NAD⁺ as the cosubstrate for SIRT3, and enhanced Nampt and SIRT3 expression maintains mitochondrial viability and promote cell survival.⁴² In the present study, aortic expression of Nampt and SIRT3 was significantly upregulated in aged

E-DN1κB mice, suggesting contributions of the sirtuin pathway to the antisenescence and prolonged longevity phenotypes. Thus, various mechanisms derived from endothelial NF-κB signaling, including systemic locomotion, systemic oxidative stress, peripheral blood flow, and mitochondrial sirtuins, may influence longevity in a complex manner (Figure IX in the online-only Data Supplement).

It was recently reported that global disruption of the angiotensin II type 1 receptor promotes longevity.⁴³ In these knockout mice as well, Nampt and SIRT3 were upregulated in the kidney. The angiotensin II type 1 receptor pathway activates a variety of intracellular signaling pathways, including NF-κB signaling.⁴⁴ However, although a growing body of evidence for angiotensin II signaling in smooth muscle cells has accumulated, much less is known about endothelial angiotensin II signal transduction and function.⁴⁵ The present study suggests the specific importance of NF-κB signaling in endothelial cells for the mechanism underlying the life-span extension observed in globally angiotensin II type 1 receptor-deficient mice. Although further intensive studies are necessary to elucidate the precise mechanisms, the endothelium apparently plays important roles in determining life span.

Adhesion molecule expression in the endothelium is well known to promote atherosclerotic plaque formation.⁴⁶ Indeed, blockade of endothelial NF-κB signaling suppresses hypercholesterolemia-induced atherosclerosis caused by apolipoprotein E deficiency.⁴⁷ Therefore, endothelial NF-κB signaling is apparently involved in the development of obesity-related disorders, including insulin resistance, hypertension, and atherosclerosis, via a variety of processes in different tissues. Furthermore, in this study, selective blockade of endothelial NF-κB signaling not only prevented age-related insulin resistance but also inhibited senescence and increased longevity with normal chow feeding. Thus, endothelial NF-κB signaling is a potential target for treatment of the metabolic syndrome and for antiaging strategies.

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Disclosures

None.

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CLINICAL PERSPECTIVE

Insulin resistance is an important mechanism underlying obesity-related disorders, eg, diabetes, hyperlipidemia, and hypertension, collectively called the metabolic syndrome. In particular, inflammation and oxidative stress are well known to play important roles in the pathogenesis of this systemic syndrome and the resultant development of atherosclerosis. A transcription factor, nuclear factor- κ B (NF- κ B), has been considered to mediate the responses to both inflammation and oxidative stress intracellularly. However, the site(s) at which the NF- κ B signaling pathway plays critical roles in these pathological processes remains to be elucidated. This study focused on the roles of endothelial NF- κ B signaling. By expressing the dominant-negative I κ B mutant in the endothelium using the transgenic procedure, the NF- κ B signaling pathway was blocked selectively in endothelial cells. These mice were protected from the development of both obesity- and age-related insulin resistance. Furthermore, importantly, these mice exhibited prolonged lifespans. In addition to the decrease in relatively early deaths, maximum lifespan was shown to be longer in these transgenic mice. Vascular senescence was markedly inhibited by blockade of endothelial NF- κ B. Thus, the endothelium plays important roles in obesity- and age-related disorders through intracellular NF- κ B signaling, ultimately impacting lifespan. Blockade of endothelial NF- κ B signaling apparently protects the whole body from both fatal morbidities at earlier ages and the development of senescence. Amelioration of insulin resistance and decreased oxidative stress are likely to contribute to these beneficial phenotypes. Therefore, endothelial NF- κ B signaling is a potential target not only for treating the metabolic syndrome, but also for anti-aging strategies.

SUPPLEMENTAL MATERIAL

Supplemental Methods

Animals

Animal studies were conducted in accordance with the institutional guidelines for animal experiments at Tohoku University. The animals were housed in an air-conditioned environment, with a 12-h light-dark cycle, and fed a regular unrestricted diet or a high-fat diet consisting of 15.3% (wt/wt) fat (Quick Fat; Nippon CLEA, Shizuoka, Japan) starting at 8 weeks of age. The mutant cDNA for human I κ B α , with alanine substitutions of two serine residues (32 and 36), was cloned into a transgenic vector, pSPTg.T2FpAXK, provided by Thomas N. Sato. This vector contains the Tie2 promoter, SV40 polyA signal and Tie2 minimum enhancer fragment. To generate transgenic mice, the construct cDNA was linearized with Sall1 digestion and microinjected into fertilized oocytes by Oriental Yeast Co. Genotyping was performed by PCR of tail DNA using the primers 5'-CCATGCGAGCGGGAAGTC-3' and 3'-CGGAGCTCAGGATCACA-5'. The two lines of transgenic mice used in the experiment had similar phenotypes. Founder mice were backcrossed for at least 6 generations with C57BL/6J mice (The Jackson Laboratory, ME, USA). E-DN I κ B;A^{y/+} mice were obtained by mating male KK A^{y/+} (A^{y/+}) mice (Nippon CLEA, Shizuoka, Japan), a genetic model for obesity-diabetes syndrome, and female E-DN I κ B mice. Male E-DN I κ B Tg/+; A^{y/+} and littermate control male A^{y/+} mice were used in the experiment. E-DN I κ B mice were crossed with endothelial nitric oxide synthase (eNOS)-deficient (Nos3^{-/-}) mice with the C57BL/6J background¹ (The Jackson Laboratory, ME, USA) to generate E-DN I κ B;Nos3^{-/-} mice. Littermate Nos3^{-/-} mice were used as controls in these experiments.

Blood analysis

Blood glucose was determined as described previously.² Plasma TBARS and 8-isoprostane levels were measured with Assay Kits (Cayman Chemical Co, MN, USA).

Glucose and insulin tolerance tests

Glucose and insulin tolerance tests were performed as described previously.³ Glucose tolerance tests were performed on fasted (10 h) mice. Mice were injected with glucose into the intraperitoneal space, and blood glucose was assayed immediately before and at 15, 30, 60, 90 and 120 min postadministration. Insulin tolerance tests were performed on fed mice. Mice were injected with human regular insulin (Eli Lilly, Kobe, Japan), and blood glucose was assayed immediately before and at 15, 30, 45 and 60 min postinjection.

Hyperinsulinemic-euglycemic clamp

Hyperinsulinemic-euglycemic clamp studies were performed as described previously.⁴ Chronically cannulated, conscious and unrestrained mice were fasted for 6 h before the study. Insulin ($500 \text{ mU} \cdot \text{kg}^{-1} \cdot \text{min}^{-1}$) was infused throughout the clamp study. Blood glucose was monitored every 5 min via carotid arterial catheter samples. Glucose was infused at a variable rate to maintain blood glucose at 120 mg/dl. The glucose infusion rate was calculated as described previously.⁴

Histological analysis

Tissues sections were removed, fixed with 10% formalin, and embedded in paraffin. The streptavidin–biotin (SAB) method was performed using a Histofine SAB-PO kit

(Nichirei, Tokyo, Japan) for immunostaining with antibodies against NF κ B p65 (C20) (Santa Cruz Biotechnology, CA, USA) and MOMA2 (Serotec Immunological Excellence, Oxford, UK). Slides were next incubated with the biotinylated IgG for 1h and then with peroxidase-conjugated streptavidin for 30 min at room temperature. Finally, immunoreactivity was visualized by incubation with a substrate solution containing 3,3'-diaminobenzidine tetrahydrochloride (DAB). For experiments involving TNF- α induced p65 translocation, mice were injected with recombinant murine TNF- α (R&D Systems Inc, MN, USA), and killed 30 minutes later.

Immunoblotting

Lung samples obtained from mice were homogenized and subjected to immunoblotting as described previously.⁵ Antibodies to I κ B- α (C-15, 21) (Santa Cruz Biotechnology, CA, USA) were commercially obtained.

Quantitative RT-PCR–based gene expression

Total RNA was isolated from mouse tissues with Isogen (Wako Pure Chemical, Osaka, Japan), and cDNA was synthesized with a Cloned AMV First Strand Synthesis Kit (Invitrogen, MD, USA) using 5 μ g of total RNA. cDNA synthesized from total RNA was evaluated using real-time quantitative PCR (Light Cycler Quick System 350S; Roche Diagnostics). The relative amount of mRNA was calculated with β -actin as the invariant control. Samples from skeletal muscle were calculated with α -actin as the invariant control. The oligonucleotide primers are described in Table S1. The mitochondrial DNA content was quantified using a sequence detection system (QIAGEN Inc., CA, USA). Total DNA was extracted from gastrocnemius muscles. The reactions were performed as follows: initial denaturing step at 95°C for 10 minutes and 40 cycles of 95°C for 15

seconds and 60°C for 1 minute. A melting curve was analyzed to check the specificity of the PCR product. The primer sequences were: 5'-GCC TTT CAG GAA TAC CAC GA-3' and 5'-CCA ATT TTA GGG GGT TCG AT-3' (GenBank NC 005089). The relative amounts of mitochondrial DNA were calculated with α -actin mRNA as the invariant control.

Oxygen consumption

Oxygen consumption was measured with an O₂/CO₂ metabolism measuring system (model MK-5000RQ; Muromachi Kikai, Tokyo, Japan) as described previously.²

Locomotor activity

Spontaneous locomotor activities of mice were analyzed with an infrared activity monitor (Supermex; Muromachi Kikai, Tokyo, Japan), as described previously.⁶ In this system a sensor monitors motion in multiple zones of the cage and movement of animal in the X, Y and Z axis can be determined. Mice were first acclimatized to the cages and housed individually for 4 days before measurement were taken. Food and water were provided *ad libitum*. All counts were automatically recorded at 60-min intervals and totalled for both the 12-hour-light and 12-hour-dark period. Data were averaged over the 3-day period of measurement.

Muscle blood flow measurement

A catheter was inserted in the carotid artery for injection of fluorescent microspheres, a second one was inserted in the femoral artery for reference blood sample withdrawal. After stabilization of hemodynamic parameters, 200 μ l of yellow-green fluorescent

microspheres (Triton, CA, USA) were injected into the carotid artery with an injection syringe over 10 s followed by 0.1 ml of saline. A reference blood sample was withdrawn from the femoral artery at a rate of 0.25 ml/min, into a pre-weighed heparinated syringe, starting 10 s before microsphere injection and lasting for a total of 60 s. The syringe containing the blood sample was weighed and the blood was digested with 250 μ l of 16N KOH. Mice were sacrificed and skeletal muscles were weighed and digested in 4 ml of 4N KOH with 20% Tween 80. After 24 h, the digested tissues were filtered individually and processed for fluorescence quantification.⁷

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Supplemental Figure Legends

Supplemental Figure 1. Body compositions of $A^y/+$ control and E-DN1κB; $A^y/+$ mice

(A) Body weights of $A^y/+$ control (white bars) and E-DN1κB; $A^y/+$ (black bars) mice at 20 weeks of age. (B) Epididymal fat weights of $A^y/+$ control and E-DN1κB; $A^y/+$ mice at 20 weeks of age. (C) Liver weights of $A^y/+$ control and E-DN1κB; $A^y/+$ mice at 20 weeks of age. Data are presented as means \pm SEM. * $P < 0.05$ compared with $A^y/+$ control littermate group by one-way ANOVA. $n=5$ in $A^y/+$ control and $n=6$ in E-DN1κB; $A^y/+$ mice.

Supplemental Figure 2. Hepatic expressions of gluconeogenic genes from $A^y/+$ control and E-DN1κB; $A^y/+$ mice

Hepatic expressions of gluconeogenic genes from $A^y/+$ control (white bars, $n=4$) and E-DN1κB; $A^y/+$ (black bars, $n=6$) mice at 20 weeks of age were analyzed by RT-PCR. The relative amounts of mRNA were calculated with β -actin mRNA as the invariant control. Data are presented as means \pm SEM.

Supplemental Figure 3. Adipose expressions of adhesion molecules and angiogenesis markers from $A^y/+$ control and E-DN1κB; $A^y/+$ mice

Expressions of adhesion molecules and angiogenesis markers in epididymal fat tissues from $A^y/+$ control (white bars, $n=5$) and E-DN1κB; $A^y/+$ (black bars, $n=4$) mice at 20 weeks of age were analyzed by RT-PCR. The relative amounts of mRNA were calculated with β -actin mRNA as the invariant control. Data are presented as means \pm SEM. * $P < 0.05$ compared with control littermate group by one-way ANOVA.

Supplemental Figure 4. Blockade of endothelial NF- κ B signaling prevented insulin resistance in genetically obese ($A^y/+$) mice

Plasma levels of adipokines, inflammatory-related cytokines and insulin in $A^y/+$ control (white bars, n=5) and E-DN κ B; $A^y/+$ (black bars, n=6) mice were measured. Data are presented as means \pm SEM. * P <0.05 compared with $A^y/+$ control littermate group by one-way ANOVA.

Supplemental Figure 5. Aortic expressions of anti-oxidant enzymes were suppressed in E-DN κ B; $A^y/+$ mice

Aortic gene expressions of anti-oxidant enzymes from $A^y/+$ control (white bars, n=5) and E-DN κ B; $A^y/+$ (black bars, n=6) mice at 20 weeks of age were analyzed by RT-PCR. The relative amounts of mRNA were calculated with β -actin mRNA as the invariant control. Data are presented as means \pm SEM. * P <0.05, ** P <0.01 compared with $A^y/+$ control littermate group by one-way ANOVA.

Supplemental Figure 6. Oxygen consumption during the dark phase was increased in E-DN κ B; $A^y/+$ mice

Oxygen consumption of $A^y/+$ control littermates (white bars, n=4) and E-DN κ B; $A^y/+$ mice (black bars, n=4) were measured at 16 weeks of age. Data are presented as means \pm SEM. * P <0.05 compared with $A^y/+$ control littermate group by one-way ANOVA.

Supplemental Figure 7. Protection from age-related body weight gain in E-DN κ B mice

(A) Food intakes of control (white bars, n=5) and E-DN κ B (black bars, n=6) of

50-week-old mice maintained on a normal chow diet. (B) Body weights of aged control and E-DN κ B mice at 50 weeks of age. Data are presented as means \pm SEM. * P <0.05 compared with control littermate group by one-way ANOVA.

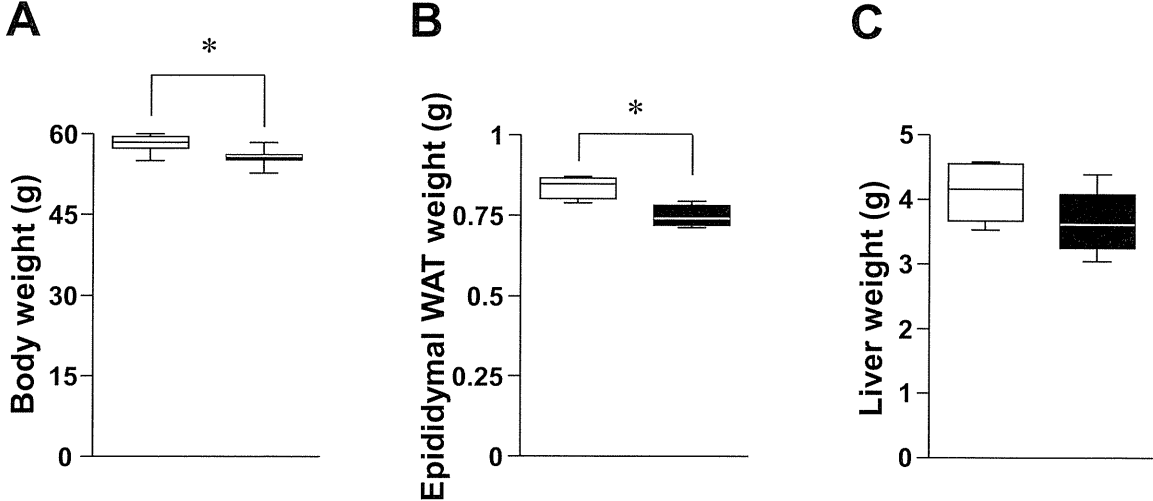
Supplemental Figure 8. eNOS deficiency suppresses the effects of endothelial DN κ B expression on blood pressure and muscle blood flow

(A) Blood pressures and (B) gastrocnemius muscle blood flows of wild-type, E-DN κ B, control Nos3^{-/-} and E-DN κ B;Nos3^{-/-} mice at 50-60 weeks of age. Data are presented as means \pm SEM. * P <0.05 compared with control littermate group by one-way ANOVA. n=4-5 in each group.

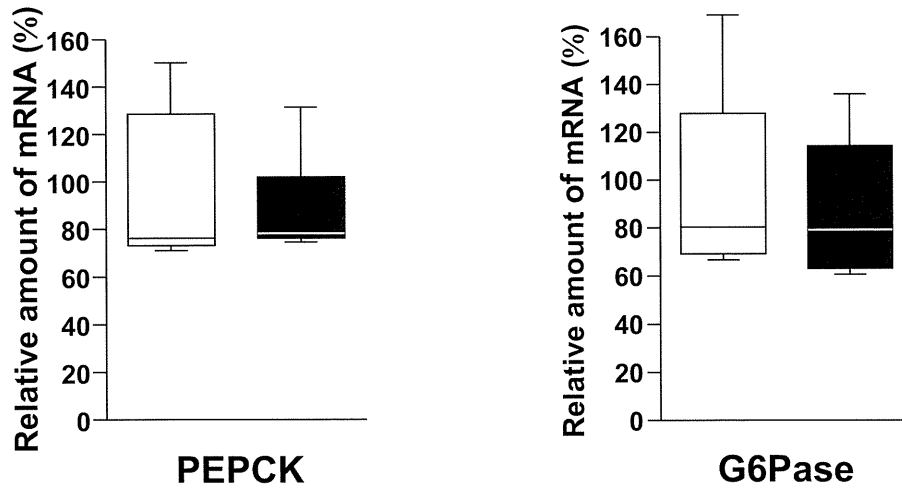
Supplemental Figure 9. Schematic diagram illustrating the proposed functions of endothelial NF- κ B signaling

Supplemental Table 1. Sequences of Quantitative RT-PCR primers

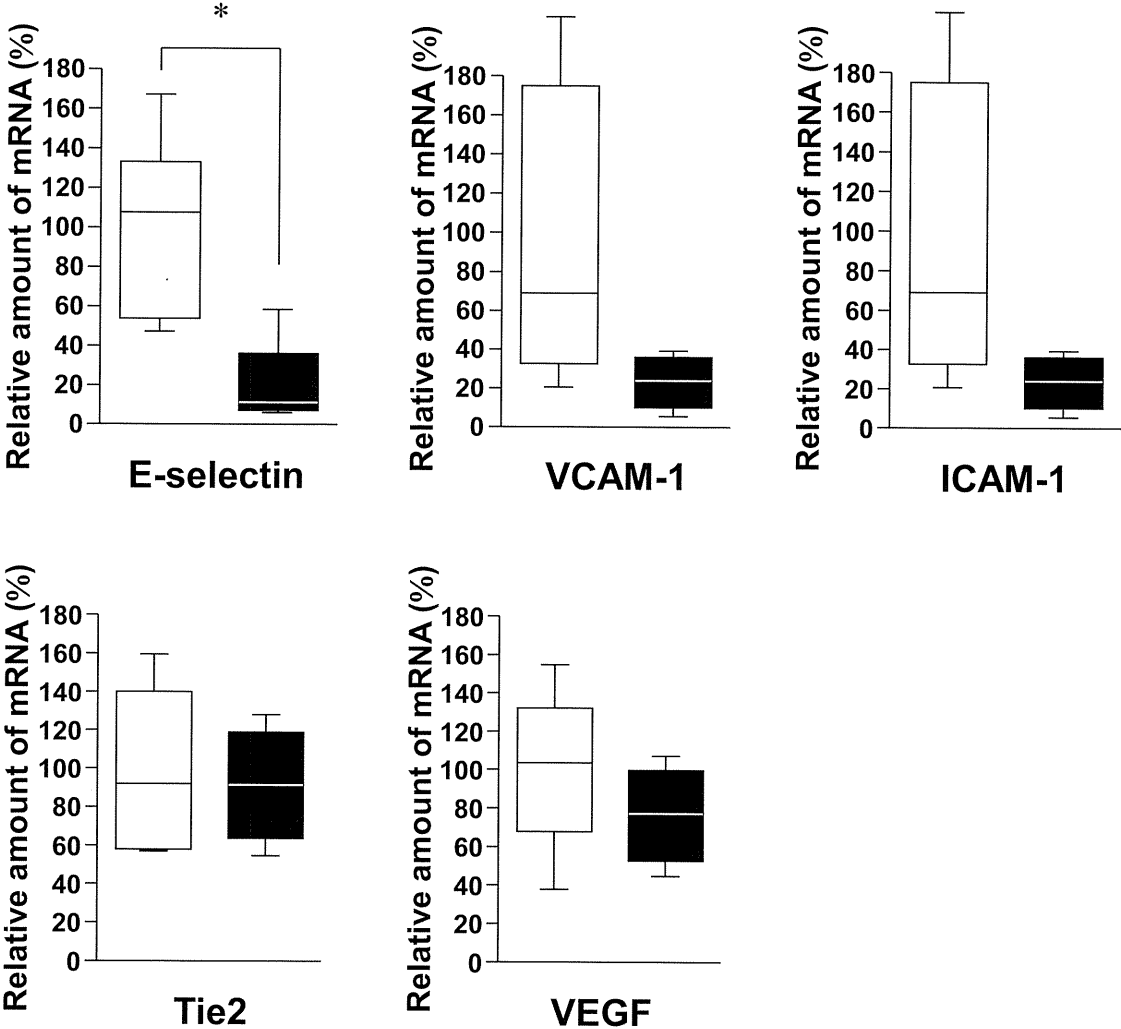
Supplemental Figure 1



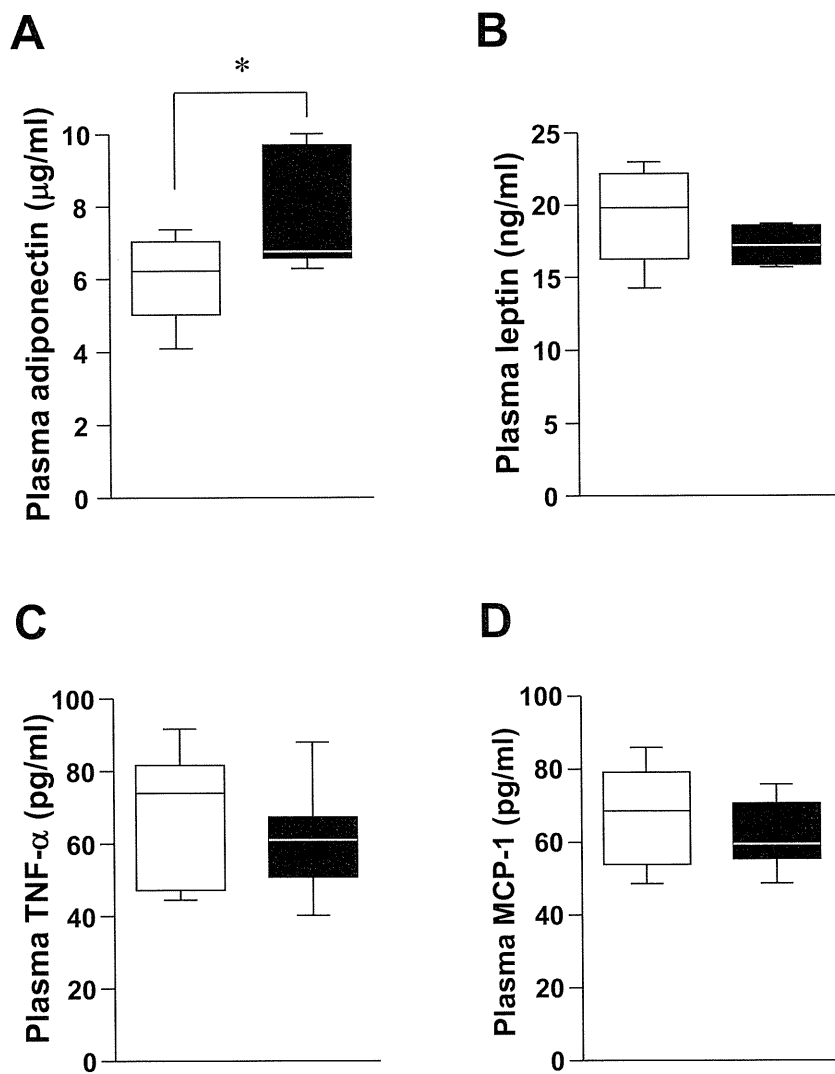
Supplemental Figure 2



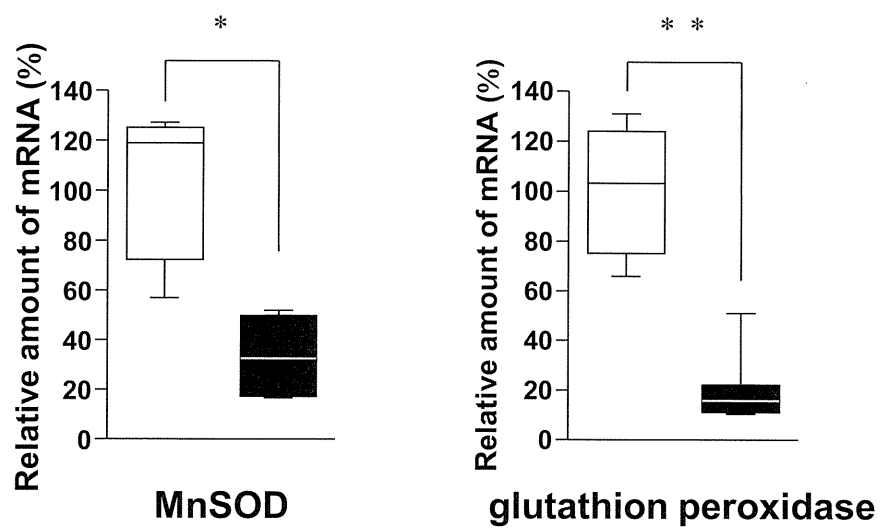
Supplemental Figure 3



Supplemental Figure 4



Supplemental Figure 5



Supplemental Figure 6

