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Loss of periostin ameliorates adipose tissue inflammation and fibrosis *in vivo*

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Recent evidence suggests that the accumulation of macrophages as a result of obesity-induced adipose tissue hypoxia is crucial for the regulation of tissue fibrosis, but the molecular mechanisms underlying adipose tissue fibrosis are still unknown. In this study, we revealed that periostin (*Postn*) is produced at extraordinary levels by adipose tissue after feeding with a high-fat diet (HFD). *Postn* was secreted at least from macrophages in visceral adipose tissue during the development of obesity, possibly due to hypoxia. *Postn*^{-/-} mice had lower levels of crown-like structure formation and fibrosis in adipose tissue and were protected from liver steatosis. These mice also showed amelioration in systemic insulin resistance compared with HFD-fed WT littermates. Mice deficient in *Postn* in their hematopoietic compartment also had lower levels of inflammation in adipose tissue, in parallel with a reduction in ectopic lipid accumulation compared with the controls. Our data indicated that the regulation of *Postn* in visceral fat could be beneficial for the maintenance of healthy adipose tissue in obesity.

A gradual and healthy expansion of adipose tissue in response to caloric intake is associated with appropriate vascular and extracellular matrix (ECM) remodeling¹. In contrast, increased adiposity is likely to involve a state of chronic inflammation in adipose tissue, thereby inducing systemic insulin resistance and ectopic lipid accumulation.

Previous studies showed that macrophages are crucial in shifting adipose tissue inflammation and ECM remodeling towards a pathological state versus an adaptive response. Macrophages in lean adipose tissue secrete anti-inflammatory cytokines, which contribute to the maintenance of tissue homeostasis^{2–5}. However, M1-like macrophages, which produce pro-inflammatory mediators such as tumor necrosis factor α (TNF- α), accumulate in adipose tissue and promote insulin resistance in conditions of obesity^{2,3}. It is conceivable that increased adipose tissue inflammation stimulates adipocyte lipolysis and tissue fibrosis, and thus enhances the release of free fatty acids, which may accumulate in non-adipose tissues as ectopic fat^{6,7}. Although recent evidence suggests that the accumulation of macrophages resulting from obesity-induced adipose tissue hypoxia are crucial for the regulation of tissue fibrosis^{8–10}, the molecular mechanisms underlying adipose tissue fibrosis are still unknown.

Periostin (*Postn*) is a secreted ECM/matricellular protein that contains 4 fasciclin domains homologous to the insect protein fasciclin I, which is involved in cell adhesion^{11,12}. At baseline, low levels of *Postn* were detected in various adult tissues. On the other hand, *Postn* was induced by various cytokines and can sustain or amplify inflammation. It is also known that *Postn* is secreted as a latent consequence of inflammatory responses rather than regulating its processes¹³. In addition, recent work has shown that hypoxia-inducible *Postn* can act as a chemoattractant for tumor-associated macrophage recruitment in a mouse model of glioblastoma^{14,15}. Another role of *Postn* has been defined in many tissues, where the loss of *Postn* results in decreased collagen cross-linking and degradation of the ECM^{16–18}. *Postn* has been shown to be linked to several inflammatory diseases, including atherosclerosis, asthma, and skin inflammation as well as malignant diseases such as breast, colon, head and neck, pancreatic, lung, gastric, and hepatocellular carcinoma¹⁹. In the case of metabolic disease, there is experimental

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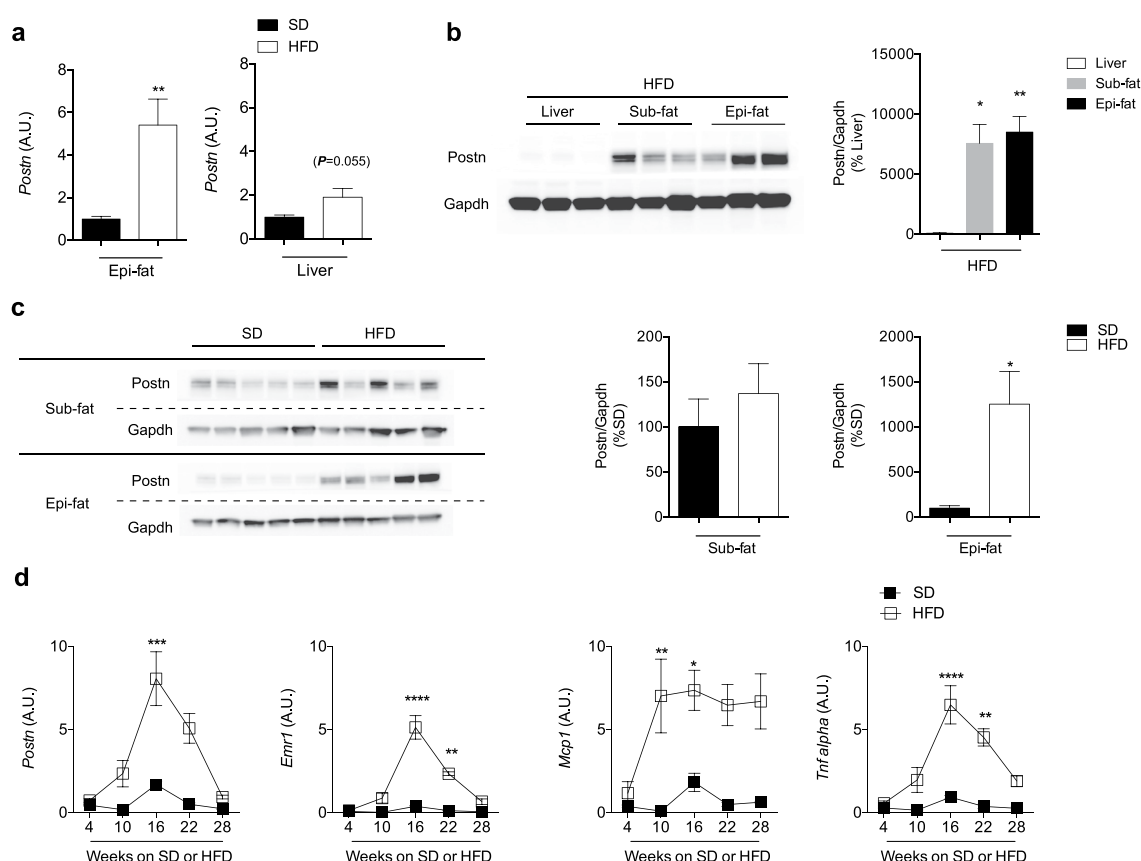


Figure 1. Periostin (*Postn*) is dramatically upregulated in adipose tissue during obesity. (a) mRNA expression levels of *Postn* in the liver and adipose tissue (epididymal fat [Epi-fat]) of C57BL/6J male mice after SD or HFD feeding for 16 weeks. $n = 5$ in each group. $**p < 0.01$ by unpaired t-test. (b) Protein levels of *Postn* in the liver and adipose tissue (subcutaneous fat [Sub-fat]) from diet-induced obese mice after 16 weeks of HFD feeding. $n = 5$, $*p < 0.05$, $**p < 0.01$ by one-way ANOVA. (c) Protein levels of *Postn* in adipose tissue from lean and obese mice after 16 weeks of SD or HFD feeding, respectively. $n = 5$, $*p < 0.05$ by unpaired t-test. (d) Time course of *Postn* and pro-inflammatory gene profiling in Epi-fat of WT SD-fed or HFD-fed mice for up to 36 weeks. $n = 3$ to 6, $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, $****p < 0.0001$ versus SD at each time point. Statistical comparisons were made by unpaired t-test.

evidence implicating *Postn* in hepatic steatosis, inflammation, and fibrosis²⁰. It was also reported that hepatic *Postn* expression increased eight fold in mice fed a high-fat diet (HFD) compared with a standard diet (SD)²⁰.

Here, we revealed that *Postn* is produced in extraordinary levels from adipose tissue after HFD feeding, and the protein level of *Postn* was ~80 times higher than that observed in the liver. The source of *Postn* was at least from macrophages in visceral adipose tissue during the development of obesity. Hypoxia increased *Postn* expression in macrophages, resulting in further recruitment of macrophages. *Postn*-deficient (*Postn*^{-/-}) mice were resistant to obesity-induced crown-like structure (CLS) formation and adipose tissue fibrosis. Less fibrosis was accompanied with the expansion of adipocyte size and reduced ectopic lipid accumulation compared to wild-type (WT) mice. Thus, *Postn*^{-/-} mice showed amelioration in systemic insulin resistance compared with HFD-fed WT littermates (*Postn*^{+/+}). Mice deficient in *Postn* in their hematopoietic compartment were also resistant to ectopic lipid accumulation, in parallel with a marked reduction in inflammation in adipose tissue.

Results

Periostin expression in adipose tissues during obesity. To explore the role of *Postn* during obesity, we analyzed *Postn* expression in the liver and adipose tissue (subcutaneous fat [Sub-fat] and epididymal fat [Epi-fat]) from C57BL/6J male mice. Eight-week-old mice were fed either SD or HFD for up to 30 weeks. Body and liver weights of mice gradually increased over the duration of HFD feeding, but adipose tissue weights peaked after 10–16 weeks of HFD feeding (Supplementary Fig. 1a). Histological examination revealed that liver steatosis and Epi-fat inflammation were evident after HFD feeding for 16 weeks. However, there was little change in Sub-fat tissues (Supplementary Fig. 1b). After 16 weeks of HFD feeding, *Postn* mRNA expression was higher in Epi-fat tissue than in the liver (Fig. 1a). *Postn* protein levels in Epi-fat were approximately 80 times higher than those observed in the liver (Fig. 1b). A marked induction of *Postn* in Epi-fat compared with Sub-fat by HFD feeding suggested a specific role of *Postn* in visceral fat during obesity (Fig. 1c).

Next, we examined the time course of *Postn* expression in Epi-fat, and it was found that *Postn* mRNA levels changed roughly in parallel with inflammatory gene expression, including expression of *Emr1*, *Mcp1*, and *Tnf- α* (Fig. 1d). Of note, interstitial fibrosis in Epi-fat decreased gradually after 16 weeks of HFD feeding (Supplementary Fig. 1c) in parallel with down-regulation of *Postn* and inflammatory gene expression levels (Fig. 1d).

Altered lipid distribution in *Postn*^{-/-} mice. Eight-week-old male C57BL/6J mice that had undergone systemic genetic ablation of *Postn* and age- and sex-matched WT littermates were fed either SD or HFD for 16 weeks. The weight of SD-fed *Postn*^{-/-} mice tended to be reduced throughout the experimental period. Although the body weight of HFD-fed *Postn*^{-/-} mice was lower in the early phase (Fig. 2a), it caught up with that of WT littermates after 10 weeks of HFD feeding (18 weeks of age). There was no difference in HFD intake for a week (Supplementary Fig. 2a). After 16 weeks of HFD feeding, the Epi-fat weight normalized to body weight was increased with a reciprocal reduction in liver weight in *Postn*^{-/-} mice relative to WT littermates after HFD feeding (Fig. 2b). Epi-fat *Postn* levels were increased both at the mRNA and protein levels (Supplementary Fig. 2b and c). Histological examination revealed that hepatic steatosis and interstitial fibrosis in Epi-fat were markedly attenuated in *Postn*^{-/-} mice compared with WT littermates (Figs 2c, 3a–c). Consistent with this, hepatic triglyceride content and serum aspartate aminotransferase (AST) and alanine aminotransferase (ALT) levels were significantly reduced in *Postn*^{-/-} mice compared with WT littermates (Fig. 2c, Supplementary Fig. 2c). On the other hand, there were no apparent differences in T-cho, LDL-C, HDL-C, TG or NEFA after 16 weeks of HFD feeding between these two genotypes of mice (Supplementary Fig. 2d). However, when we observed serial changes in organ weight after HFD feeding, Epi-fat weight peaked at 10 weeks after HFD feeding and liver weight is increased gradually until 22 weeks on a HFD (Supplementary Fig. 1a). Similar changes were also observed in the histology of the liver and adipose tissue (Supplementary Fig. 1c). These results also suggested that changes in adipose tissue can affect lipid distribution in the liver. It is known that *Postn*^{-/-} mice showed specifically impaired pancreatic regeneration in islet β -cells²¹. However, pancreatic regeneration was not observed after HFD feeding and it was not related with our findings in *Postn*^{-/-} mice. Actually, the islet size in *Postn*^{-/-} mice was relatively smaller than that of WT mice (Supplementary Fig. 2e), which may indicate the amelioration of insulin resistance in *Postn*^{-/-} mice compared with WT mice. All of these results demonstrated that the loss of *Postn* affected lipid distribution between liver and Epi-fat tissues. Concerning to the *Postn* receptors, we measured the expression levels of *Itgav*, *Itga6*, and *Itgb3* in Epi-fat and liver. As shown in Supplementary Fig. 2f, *Itgav* and *Itgb3* were reduced in *Postn*^{-/-} mice Epi-fat, which may indicate the reduced *Postn* signaling.

Reduced CLS formation and adipose tissue fibrosis in *Postn*^{-/-} mice. Histological examination revealed that adipocyte cell size from Epi-fat and Sub-fat in HFD-fed mice was larger in *Postn*^{-/-} mice than in WT littermates (Fig. 3a,b, Supplementary Fig. 3a,b). In addition, Masson's trichrome staining showed extensive interstitial fibrosis in Epi-fat of WT littermates, which was markedly suppressed in *Postn*^{-/-} mice (Fig. 3c).

We next examined the number of CLSs in Epi-fat by double-immunofluorescence staining with anti-F4/80 (red) and anti-Perilipin (green) antibodies. The number of F4/80-positive cells and the density of CLSs in the Epi-fat of HFD-fed *Postn*^{-/-} mice were markedly reduced compared with those of HFD-fed WT littermates (Fig. 3d). Consistent with these staining results, flow cytometric analysis revealed that the number of F4/80⁺ cells isolated from Epi-fat was lower in HFD-fed *Postn*^{-/-} mice than in HFD-fed WT littermates (Supplementary Fig. 3c,d). Moreover, there was a significant decrease in pro-inflammatory gene expression levels, such as *Emr1*, *Mcp1*, and *Tnf- α* in Epi-fat of *Postn*^{-/-} mice than in those of WT littermates (Fig. 3e). In addition, *Pnpla2* and *Hsl* expressions were significantly reduced in *Postn*^{-/-} mice compared with WT littermates, which indicated that reduced lipolysis may have attenuated the altered lipid distribution in the liver of *Postn*^{-/-} mice (Fig. 3f).

***Postn*^{-/-} mice show ameliorated insulin resistance *in vivo*.** We next examined the effect of *Postn* deficiency on glucose homeostasis in HFD-fed mice. We performed intraperitoneal (IP) glucose tolerance tests (GTTs) after 12 weeks of SD or HFD feeding and found that *Postn*^{-/-} mice showed better glucose tolerance than HFD-fed WT littermates (Fig. 4a). The insulin response during the GTTs was significantly lower in HFD-fed *Postn*^{-/-} mice versus WT littermates (Fig. 4b). ITTs after 14 weeks of HFD feeding showed that HFD-fed *Postn*^{-/-} mice were more insulin sensitive than WT littermates (Fig. 4c). There was no significant difference between these mice when they were fed SD (Fig. 4d). Moreover, the insulin-induced phosphorylation of AKT was significantly increased in the Epi-fat of *Postn*^{-/-} mice compared with that of WT littermates, and the same tendency was observed in the liver in *Postn*^{-/-} mice compared with that of WT littermates (Fig. 4e).

***Postn* is induced in Epi-fat tissue macrophages by obesity.** To identify the cellular sources of *Postn*, we isolated adipocytes and the stromal vascular fraction (SVF), which was further divided into F4/80⁻ cells and F4/80⁺ cells, from the Epi-fat of lean and obese mice (Supplementary Fig. 4a). *Postn* mRNA levels were significantly higher in SVF than that in adipocytes (Fig. 5a). When cells were divided into subfractions, *Postn* mRNA levels were significantly higher in F4/80⁻ cells than in other fractions both with SD and HFD feeding. Although *Postn* mRNA levels were lower in F4/80⁺ cells than in any of the other cell fractions from the Epi-fat of SD-fed mice, it was remarkably increased in HFD-fed mice (Fig. 5b), suggesting a requirement for HFD feeding for the activation of *Postn* production in macrophages. In addition, myofibroblast marker (α SMA) and preadipocyte marker (*Pdgfr- α* and *Ppar- γ*) were activated in F4/80⁻ cells of HFD-fed mice (Supplementary Fig. 4b).

To confirm which cells produce *Postn*, we utilized *Rosa26-loxP-stop-loxP-tdTomato* transgenic mice crossed with *Periostin-Cre* mice. *tdTomato* is activated by *Postn*-specific promoter-regulated *cre* expression (Supplementary Fig. 4c). After HFD feeding for 24 weeks, when obesity-induced histological changes such as enlargement of adipocyte sizes were prominent, *tdTomato*-positive cells were both positive and negative for

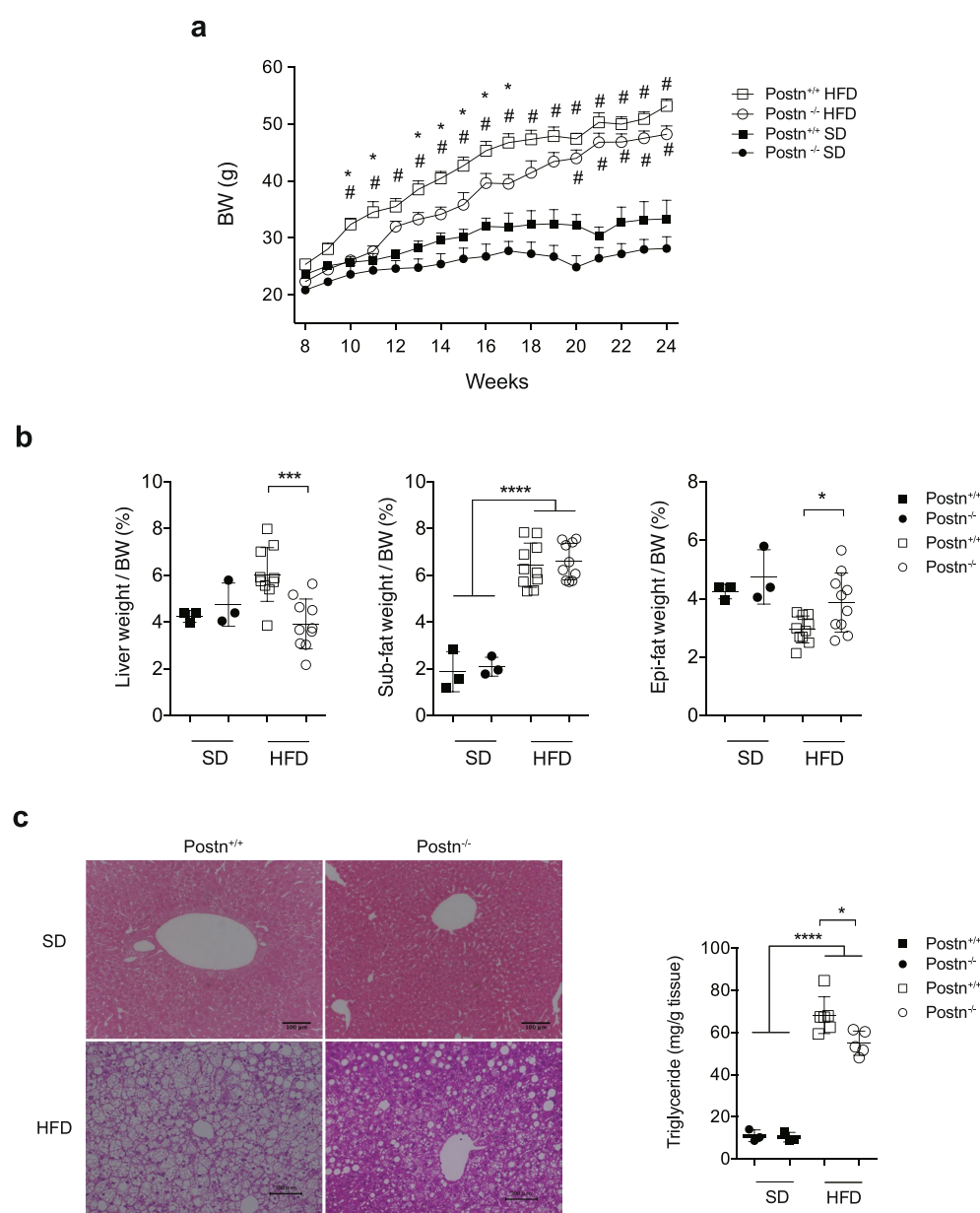


Figure 2. Altered lipid distribution in $Postn^{-/-}$ mice. **(a)** Development of body weight of $Postn^{+/+}$ and $Postn^{-/-}$ male SD-fed or HFD-fed mice. $n = 5$ in the SD groups, $n = 11-13$ in the HFD groups, * $p < 0.05$ versus HFD-fed $Postn^{+/+}$ mice, # $p < 0.05$ versus SD-fed $Postn^{+/+}$ mice by one-way ANOVA. **(b)** Liver and adipose tissue weights normalized to whole body weight of $Postn^{+/+}$ and $Postn^{-/-}$ SD-fed or HFD-fed mice. $n = 3$ for the SD groups, $n = 10$ for the HFD groups. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$ by one-way ANOVA. **(c)** Left, representative microscope images of the livers of $Postn^{+/+}$ and $Postn^{-/-}$ SD-fed or HFD-fed mice. Scale bars, 100 μ m. Right, triglyceride levels in the livers of $Postn^{+/+}$ and $Postn^{-/-}$ mice. $n = 3$ for the SD groups, $n = 5$ for the HFD groups. * $p < 0.05$, **** $p < 0.0001$ by one-way ANOVA.

anti-F4/80 antibody immunofluorescent staining (Fig. 5c). tdTomato expression did not overlap with perlipin staining, which indicates that a portion of $Postn$ -positive cells are interstitial macrophages (Supplementary Fig. 4d).

Loss of $Postn$ reduces macrophage migration *in vitro*. A previous study showed that $Postn$ can act as a chemoattractant for macrophages in a model of glioblastoma¹⁴. Actually, the migration of J774.1 macrophages towards recombinant POSTN (rPOSTN) was significantly enhanced in accordance with the increase in rPOSTN protein concentration (Fig. 6a,b). Moreover, the migration ability of $Postn^{-/-}$ macrophages was significantly

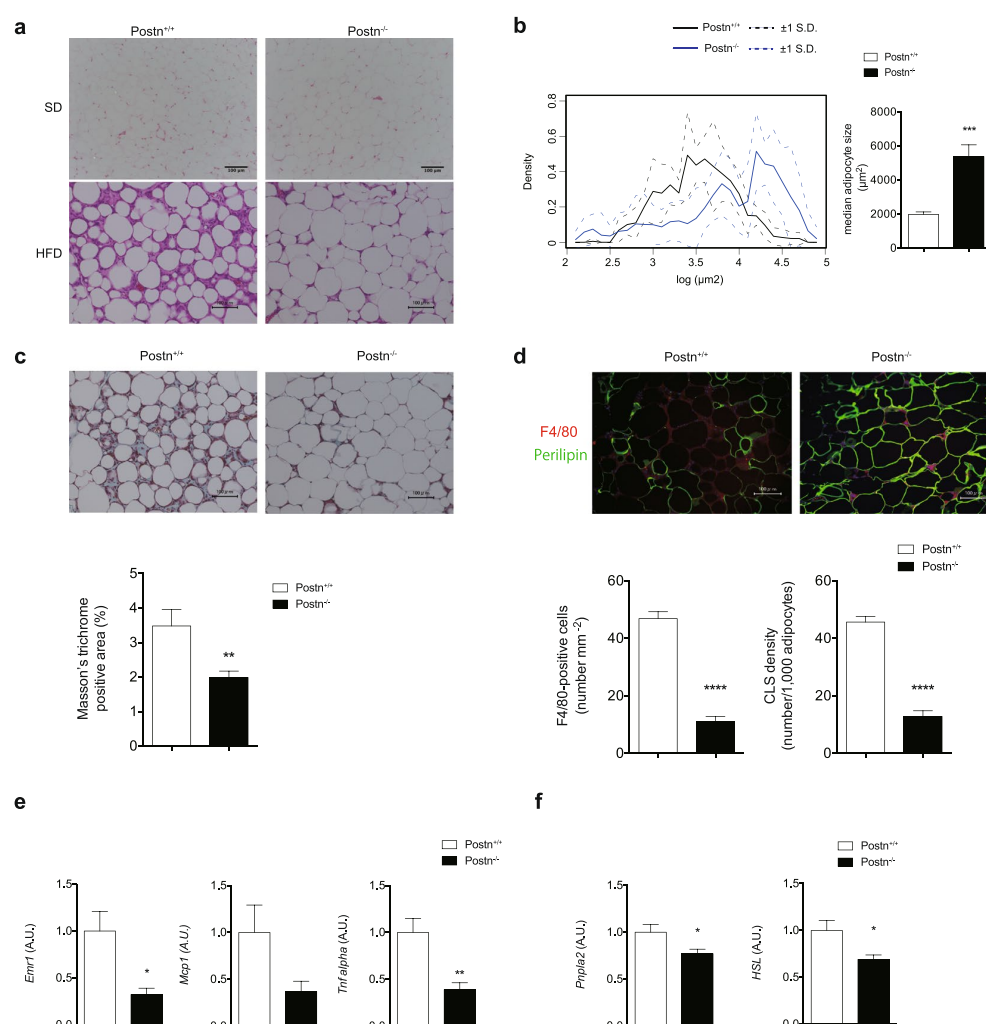


Figure 3. Ameliorated inflammation of adipose tissue in *Postn*^{-/-} mice. **(a)** Representative H&E staining images of adipose tissue (Epi-fat) of *Postn*^{+/+} and *Postn*^{-/-} SD-fed or HFD-fed mice. **(b)** Left, density curve of adipocyte surface areas in Epi-fat. Right, median adipocyte cell size in Epi-fat of *Postn*^{+/+} and *Postn*^{-/-} HFD-fed mice. ****p* < 0.001 by unpaired t-test. *n* = 3 images per mouse, 10 mice each. **(c)** Top, representative Masson's trichrome staining images of adipose tissue (Epi-fat) of *Postn*^{+/+} and *Postn*^{-/-} HFD-fed mice. Bottom, quantification of Masson's trichrome-positive area (identical with interstitial fibrosis). ***p* < 0.01 by unpaired t-test. **(d)** Top, representative immunofluorescent staining of F4/80 (red) and perilipin (green). Bottom, the number of F4/80-positive cells and crown-like structure (CLS) density in Epi-fat of *Postn*^{+/+} and *Postn*^{-/-} mice after HFD feeding for 16 weeks. Scale bars, 50 μm. *****p* < 0.0001 by unpaired t-test. **(e)** Pro-inflammatory (*Emr1*, *Mcp1* and *Tnf-α*) mRNA levels in Epi-fat from HFD-fed *Postn*^{+/+} and *Postn*^{-/-} mice. *n* = 6 each, **p* < 0.05, ***p* < 0.01 by unpaired t-test. **(f)** Lipolysis associated mRNA levels (*Pnpla2* and *Hsl*) mRNA levels in Epi-fat from HFD-fed *Postn*^{+/+} and *Postn*^{-/-} mice. *n* = 6 each, **p* < 0.05 by unpaired t-test.

impaired compared with that of WT macrophages. In addition, both TGF-β1 and rPOSTN caused a significant increase in the migration of WT macrophages but not in *Postn*^{-/-} macrophages. (Fig. 6c,d). These observations indicated a fundamental defect in the ability of macrophages to migrate in *Postn*^{-/-} mice.

Hypoxia promotes *Postn* secretion in macrophages. Recent reports suggested that hypoxia is a new potential risk factor for chronic inflammation in obesity^{22,23}. Infiltrated macrophages were predominantly found in hypoxic areas in the adipose tissue of obese mice²⁴. To determine whether hypoxia induced an increase in the expression of *Postn* in macrophages, we first treated the J774.1 macrophage cell line with deferoxamine mesylate, which mimics hypoxia, for 12 or 48 hours. We found that *Postn* expression was elevated in hypoxia mimetic conditions in parallel with genes downstream of hypoxia-inducible factor (HIF), such as *Glut1* and *Pgk1* (Fig. 6e, Supplementary Fig. 5a). Similar to this observation, *Postn* expression was enhanced in both J774.1 cells and peritoneal macrophages exposed to hypoxic conditions (5% O₂) (Fig. 6f, Supplementary Fig. 5b).

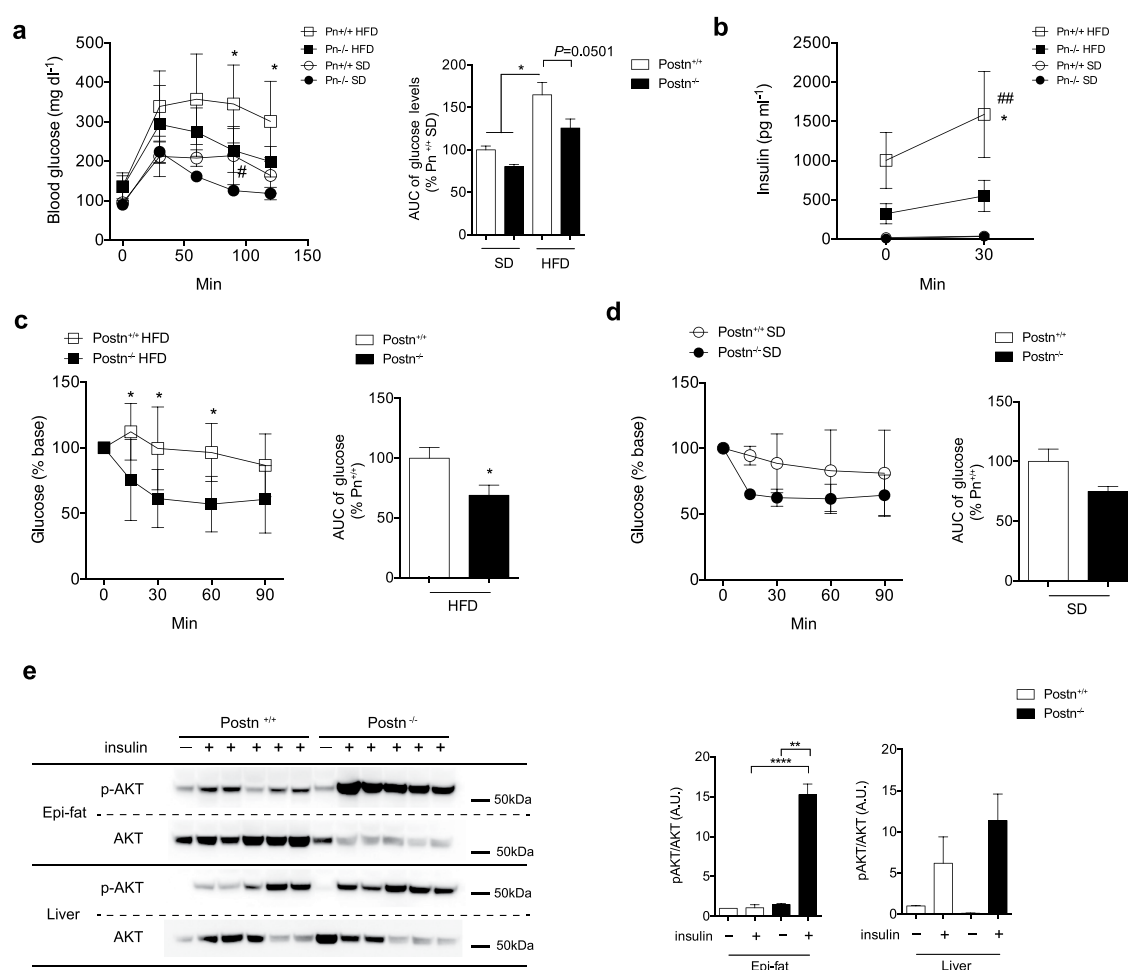


Figure 4. Ameliorated glucose metabolism in *Postn*^{-/-} mice. (a) Serial changes in glucose levels and area under the curve (AUC) of glucose levels after intraperitoneal injection of glucose in *Postn*^{+/+} and *Postn*^{-/-} SD-fed or HFD-fed mice at 12 weeks for HFD. n = 3 for the SD groups, n = 8–9 for the HFD groups. *p < 0.05 versus *Postn*^{-/-} mice fed HFD by one-way ANOVA. (b) Serial changes in insulin levels after intraperitoneal injection of glucose in *Postn*^{+/+} and *Postn*^{-/-} SD-fed or HFD-fed mice at 14 weeks for HFD. n = 4 for the SD groups, n = 6 for the HFD groups. ##p < 0.01 versus *Postn*^{+/+} SD-fed mice. *p < 0.05 versus *Postn*^{-/-} HFD-fed mice by one-way ANOVA. (c) Serial changes in glucose levels and AUC of glucose levels after intraperitoneal injection of insulin in *Postn*^{+/+} and *Postn*^{-/-} HFD-fed mice at 16 weeks for HFD. n = 5–6 each, analyzed by unpaired t-test. (d) Serial changes in glucose levels and AUC of glucose levels after intraperitoneal injection of insulin in *Postn*^{+/+} and *Postn*^{-/-} SD-fed mice. n = 4 each, analyzed by unpaired t-test. (e) Protein levels of phosphorylated Akt (Thr308) in Epi-fat and liver. *Postn*^{+/+} and *Postn*^{-/-} HFD-fed mice were injected with saline or insulin via the postcaval vein. n = 3 for saline, n = 5 for insulin. *p < 0.05, ***p < 0.0001 versus *Postn*^{+/+} insulin by unpaired t-test.

Because *Postn* is a cell-secreted protein, we incubated J774.1 cells in hypoxia mimetic conditions for 12 or 48 hours to obtain hypoxia-stimulated J774.1 cell supernatants. We found that *Postn* protein levels in the supernatants increased in hypoxia mimetic conditions (Fig. 6g,h). In contrast, protein levels of *Postn* inside the cells did not change (Supplementary Fig. 5c).

***Postn* deletion in hematopoietic cells reduced obesity-induced adipose tissue inflammation.** Considering the role of macrophages in the propagation of inflammatory signals in adipose tissue, we hypothesized that deletion of *Postn* in hematopoietic-derived cells, which include macrophages, could reduce obesity-related increases in macrophage infiltration and inflammation and subsequently prevent *in vivo* ectopic lipid accumulation and insulin resistance. To address this hypothesis, we generated mice with *Postn* deleted exclusively in hematopoietic cells. Mice transplanted with bone marrow (BM) from *Postn*^{-/-} mice (BM-*Postn*^{-/-}) were deficient in *Postn* in hematopoietic-derived cells. Mice transplanted with bone marrow from WT littermates

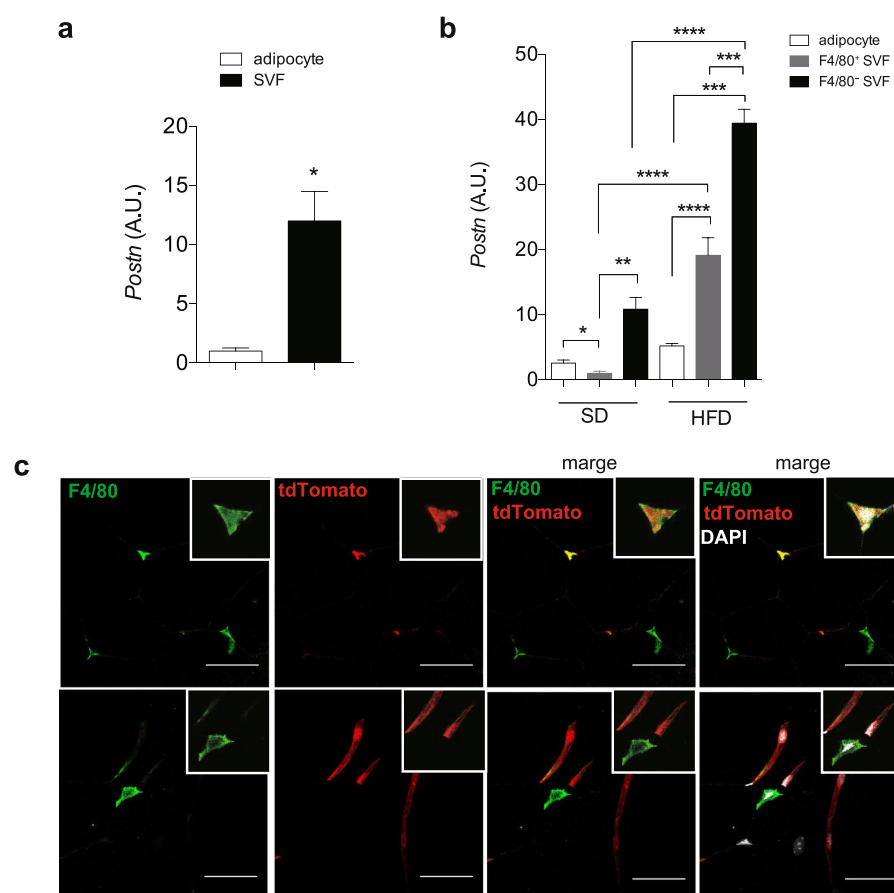


Figure 5. *Postn* is secreted by the stromal vascular fraction (SVF). (a) mRNA expression of *Postn* in adipose tissue (Epi-fat) from C57BL/6J male HFD-fed mice. $n = 3$, $*p < 0.05$ by unpaired t-test. (b) mRNA expression of *Postn* in adipocytes, F4/80⁺ and F4/80⁺ cells, isolated from SVF. $n = 3$ each, $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, $****p < 0.0001$ by one-way ANOVA. (c) Representative immunofluorescent staining of F4/80 (green) in Epi-fat from obese *tdTomato* fluorescent protein reporter mice demonstrated *Postn* expressing cells. Scale bars, 50 μ m.

(BM-*Postn*^{+/+}) displayed normal *Postn* expression in all hematopoietic and nonhematopoietic cells/tissues (Supplementary Fig. 6a).

As expected, body weight gain in HFD-fed mice significantly outpaced SD-fed mice. There were no significant differences in body weight between BM-*Postn*^{+/+} and BM-*Postn*^{-/-} mice (Fig. 7a). We did not observe differences in absolute body weight or adipose tissue weight between these two groups. However, HFD-fed BM-*Postn*^{-/-} mice had lower liver weight per body weight compared with BM-*Postn*^{+/+} mice (Fig. 7b). *Postn* mRNA and protein levels were lower in Epi-fat from HFD-fed BM-*Postn*^{-/-} mice than in those from BM-*Postn*^{+/+} mice, although *Postn* mRNA levels in the liver were not different between the two groups (Supplementary Fig. 6b,c).

Histological examination revealed that hepatic steatosis was markedly attenuated in BM-*Postn*^{-/-} mice relative to WT littermates (Fig. 7c). Consistent with this, hepatic triglyceride content and serum AST and ALT levels were significantly reduced in BM-*Postn*^{-/-} mice compared with those in BM-*Postn*^{+/+} mice (Fig. 7d,e).

Epi-fat from BM-*Postn*^{+/+} mice was characterized by higher interstitial fibrosis compared with those from BM-*Postn*^{-/-} mice, as determined using Masson's trichrome staining (Fig. 8a,b). By double-immunofluorescent staining of F4/80 (red) and perilipin (green), we observed that the number of F4/80-positive cells in BM-*Postn*^{-/-} mice was reduced compared with BM-*Postn*^{+/+} mice (Fig. 8c). We also observed higher mRNA expression levels of pro-fibrotic collagen types I, III, and VI (*Col1a1*, *Col3a1*, and *Col6a1*) and inflammatory genes (*Emr1*, *Mcp1*, and *Tnf- α*) in HFD-fed BM-*Postn*^{+/+} mice than in BM-*Postn*^{-/-} mice (Fig. 8d,e). In addition, adipocyte size from the Epi-fat was larger in HFD-fed BM-*Postn*^{-/-} mice than in BM-*Postn*^{+/+} mice (Supplementary Fig. 6d). These results were the same as those of the whole-body *Postn*^{-/-} phenotype.

Finally, we examined the effect of *Postn* deficiency in hematopoietic cells on glucose homeostasis in HFD-fed mice. Fasting glycemia, fasting insulin concentration, and insulin secretion during IPGTTs were not different between BM-*Postn*^{+/+} and BM-*Postn*^{-/-} mice. There was no change between hematopoietic genotypes in either IPGTTs or ITTs (Supplementary Fig. 7a-h).

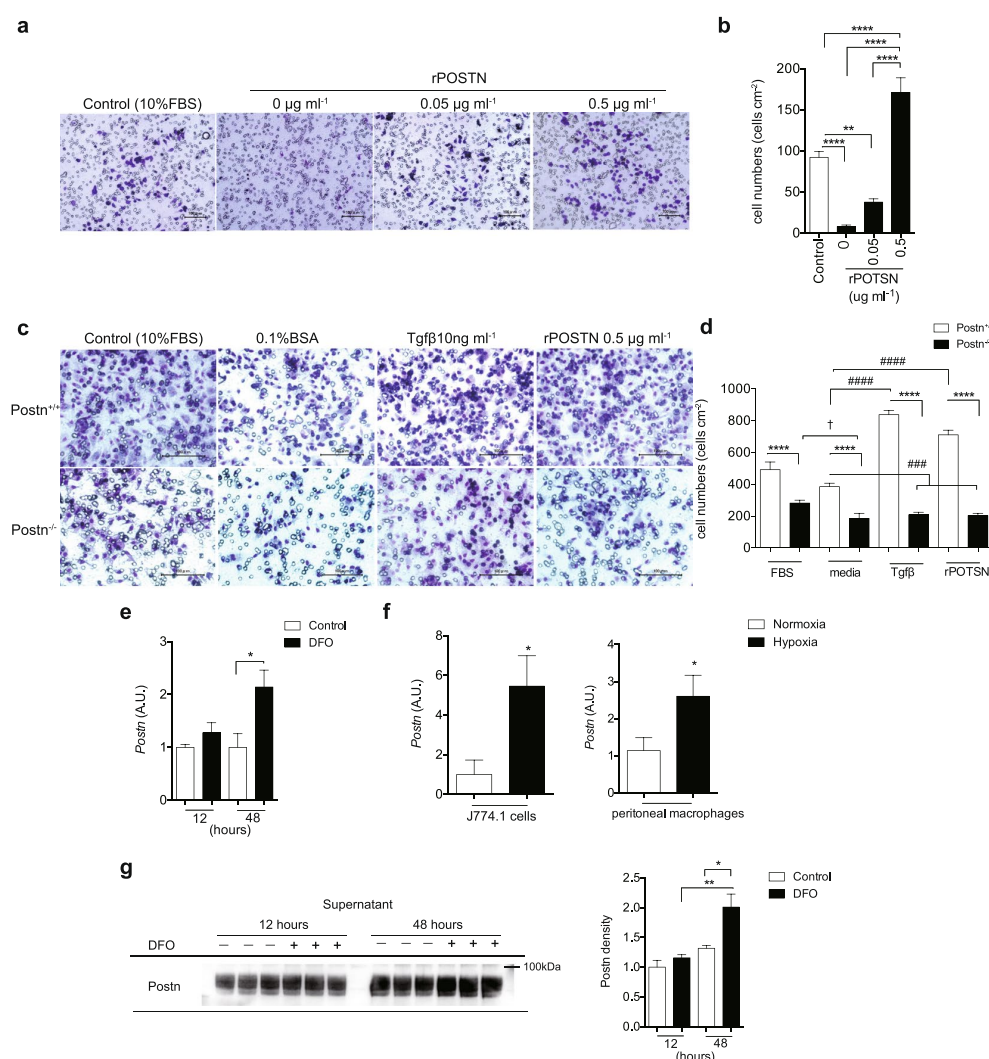


Figure 6. *Postn* regulates macrophage migration. **(a)** Representative images of J774.1 cells migrating toward rPOSTN in Transwell assays. Scale bar, 100 μm . **(b)** Graphical analysis of **a** showing the migration of J774.1. $n = 10$ –15 fields, $^{*}p < 0.01$, $^{***}p < 0.0001$ by one-way ANOVA. **(c)** Representative images of *Postn*^{+/+} and *Postn*^{-/-} peritoneal macrophages migrating toward transforming growth factor- β (TGF- β) or rPOSTN in transwell assays. Scale bar, 100 μm . **(d)** Graphical analysis of **c** showing the migration of peritoneal macrophages. $n = 8$ –12 fields for *Postn*^{+/+}, $n = 21$ –22 fields for *Postn*^{-/-}, $^{***}p < 0.001$ versus *Postn*^{+/+} at each condition, $^{***}p < 0.01$, $^{****}p < 0.0001$ versus *Postn*^{+/+} media, $^{\dagger}p < 0.05$ versus *Postn*^{-/-} media by one-way ANOVA. **(e)** mRNA expression of *Postn* in J774.1 cells with or without deferoxamine mesylate (DFO) treatment over the time course. $n = 6$, $^{*}p < 0.05$ by one-way ANOVA. **(f)** mRNA expression of *Postn* in J774.1 cells and peritoneal macrophages that were exposed to hypoxia. $n = 5$ for J774.1, $n = 5$ –7 for peritoneal macrophages, $^{*}p < 0.05$ by unpaired t-test. **(g)** Protein levels of *Postn* in J774.1 cells supernatants with or without DFO treatments over the time-course. $n = 3$ each, $^{*}p < 0.05$, $^{**}p < 0.01$ by one-way ANOVA.

Discussion

In this study, we revealed that *Postn* is produced at extraordinary levels from adipose tissue after HFD feeding for the first time. Macrophages in visceral adipose tissue were responsible for the *Postn* production and secretion during the development of obesity, possibly due to hypoxia. *Postn*^{-/-} mice fed HFD had less CLS formation and fibrosis in adipose tissue and were protected from liver steatosis and systemic insulin resistance compared with WT littermates. Mice deficient of *Postn* in their hematopoietic compartment also had lower inflammation in adipose tissue, in parallel with a reduction in ectopic lipid accumulation compared with controls.

Adipose tissue expansion is required to store surplus energy efficiently and prevent ectopic lipid deposition in metabolically sensitive tissues, thus preventing metabolic complications. Acute and local inflammation within adipose tissue plays an essential role in healthy adipose tissue expansion, remodeling, and overall homeostasis by stimulating ECM degradation and angiogenesis⁴. However, during the development of obesity, chronic,

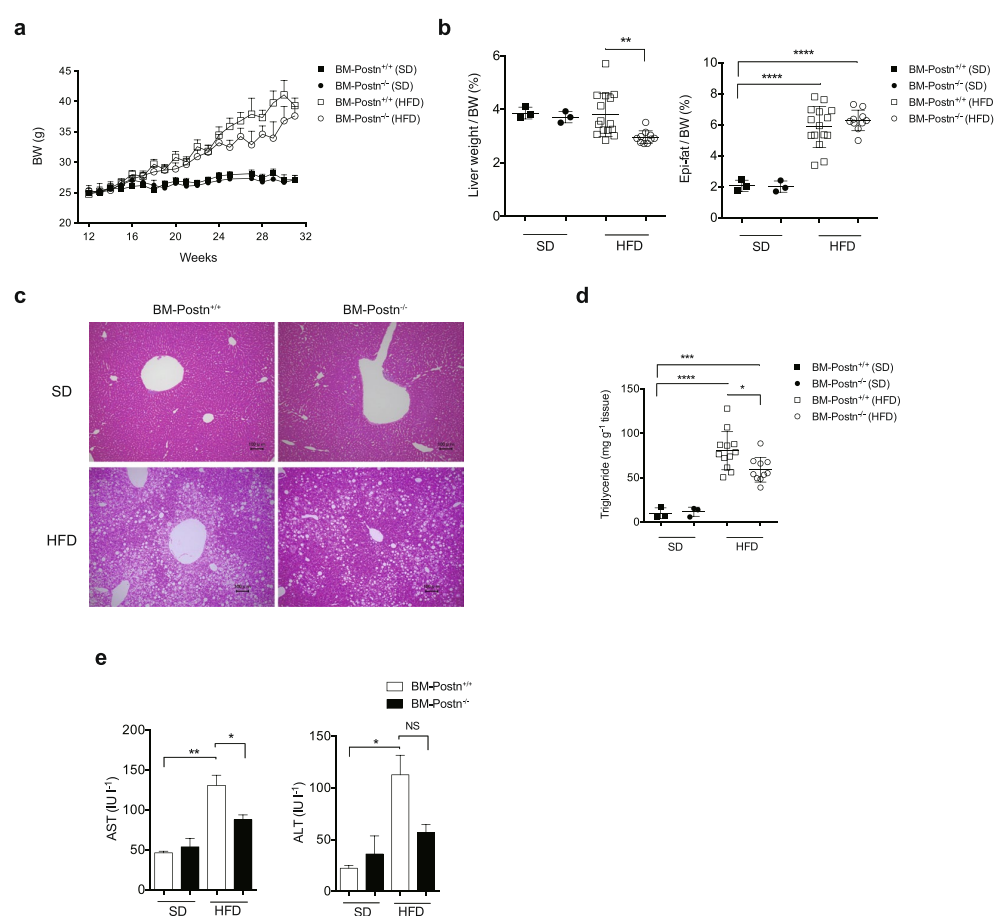


Figure 7. Improved liver steatosis in BM-*Postn*^{-/-} mice. **(a)** Development of body weight of SD-fed or HFD-fed mice over 30 weeks after BM transplantation. $n = 3$ for the SD groups, $n = 11-15$ for the HFD groups, analyzed by one-way ANOVA at each time point. **(b)** Liver and adipose tissue weights of BM transplantation recipient *Postn*^{+/+} SD-fed or HFD-fed mice transplanted with *Postn* WT hematopoietic cells (BM-*Postn*^{+/+}) and *Postn* deletion mutant hematopoietic cells (BM-*Postn*^{-/-}). $n = 3$ for the SD groups, $n = 11-15$ for the HFD groups. * $p < 0.05$, ** $p < 0.01$ by one-way ANOVA. **(c)** Representative microscope images of the livers of BM transplanted SD-fed or HFD-fed mice. Scale bars, 100 μ m. **(d)** Triglyceride levels in the livers of BM transplanted mice. $n = 3$ for the SD groups, $n = 9$ for the HFD groups. * $p < 0.05$, *** $p < 0.001$, **** $p < 0.0001$ by one-way ANOVA. **(e)** Serum profile of BM transplanted SD-fed or HFD-fed mice. $n = 3$ for the SD groups, $n = 11-15$ for the HFD groups, * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ analyzed by unpaired t-test.

low-grade inflammation is induced in adipose tissue^{25,26}. Increased adipose tissue inflammation stimulates adipocyte lipolysis and tissue fibrosis, and thus led to increases in metabolic abnormalities, such as type 2 diabetes and coronary artery disease²⁷. It has been accepted that abnormal collagen deposition is a key feature of adipose tissue dysfunction during obesity²⁸. However, how ECM dynamics reflect states of pathological fibrosis in response to over-nutrition is poorly understood. In this study, we showed that Periostin (*Postn*) is a novel regulator of adipose tissue fibrosis.

Postn, which is secreted as a latent consequence of inflammatory responses, sustains or amplifies the inflammation response^{13,29}. In fact, *Postn* expression is quite low in adipose tissue in healthy lean mice. We observed that *Postn* expression in Epi-fat changed in parallel with inflammatory gene expression during the development of obesity. It is conceivable that *Postn* is involved in pathological tissue remodeling in adipose tissue. Because *Postn* is required for the regulation of ECM production and maturation, the loss of *Postn* resulted in decreased collagen cross-linking and the degradation of ECM³⁰. Targeted deletion of matrix-involved protein such as collagen VI or matrix metalloproteinase 12 in obese mice resulted in the healthy expansion of individual adipocytes^{31,32}. In contrast, abnormal collagen accumulation in association with developmental fibrosis limited adipose tissue expansion^{33,34}. Consistent with these studies, we observed that systemic *Postn* deficiency resulted not only in reduced accumulation of macrophages and abnormal collagen deposition but also in larger adipocyte cell size in Epi-fat. At the same time, Epi-fat weight increase with a reciprocal reduction in liver weight was observed in *Postn*^{-/-} mice, which indicated that the adipose tissue fibrosis induced by *Postn* is a detrimental factor for ectopic lipid disposition and insulin resistance in obesity. A recent study demonstrated that aberrant expression of *Postn* in the

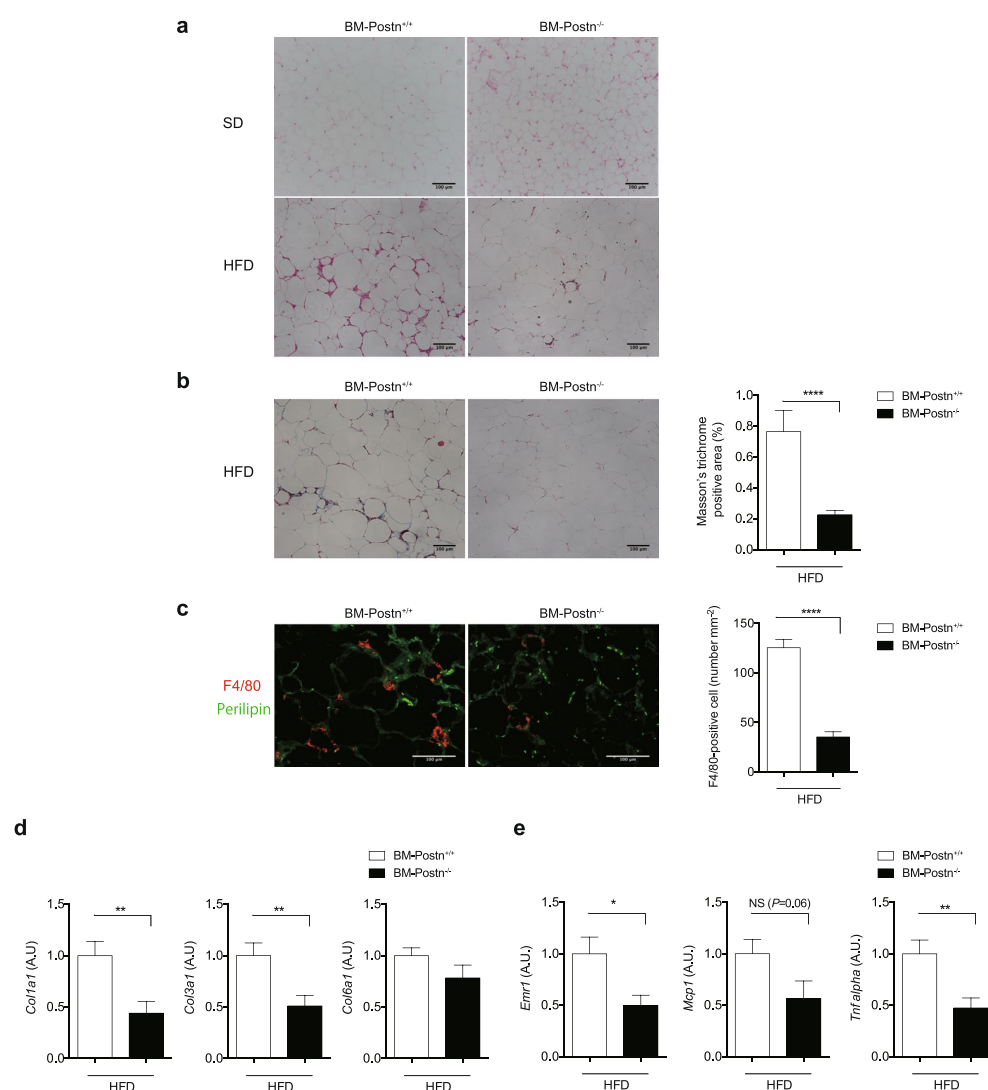


Figure 8. Hematopoietic cell-specific *Postn* deletion altered adipose tissue remodeling. **(a)** Representative H&E staining images of adipose tissue (Epi-fat) of BM transplanted mice. **(b)** Left, representative Masson's trichrome staining images of adipose tissue (Epi-fat) of BM transplanted mice. Right, quantification of Masson's trichrome-positive area (identical with interstitial fibrosis). $n = 3$ images per mouse, 5–10 mice each. **** $p < 0.0001$ by unpaired t-test. **(c)** Left, representative immunofluorescent staining of F4/80 (red) and perilipin (green). Right, the number of F4/80-positive cells in Epi-fat of BM transplanted mice. Scale bars, 100 μ m. **** $p < 0.0001$ by unpaired t-test. **(d)** Collagen (*Col1a1*, *Col3a1* and *Col6a1*) mRNA levels in Epi-fat from BM transplanted mice. $n = 9$ –15, * $p < 0.05$ by unpaired t-test. **(e)** Pro-inflammatory (*Emr1*, *Mcp1* and *Tnf- α*) mRNA levels in Epi-fat from BM transplanted mice. $n = 9$ –15, * $p < 0.05$ by unpaired t-test.

liver resulted in steatosis and hypertriglyceridemia through JNK-mediated suppression of fatty acid oxidation²⁰. However, the induction of *Postn* in adipose tissue was approximately 80 times higher than those observed in the liver in our experiment, suggesting its potent role in metabolic abnormalities in the obese condition.

ECM dynamics are believed to occur in the pro-inflammatory context of hypertrophic adipocyte, local hypoxia, and pathological immunity. Macrophages, which are known to drive an inflammatory program within obese adipose tissue, are believed to also orchestrate fibrogenesis in several organs^{35,36}. Inflammatory macrophages accumulated in the visceral adipose tissue of obese mice, especially in hypoxic areas. The reasons included: (1) macrophages act to remove dead adipocytes; and (2) macrophages are trapped in hypoxic areas by macrophage migration inhibition factor³⁷. Of note, *Postn* was reported to be increased in hypoxic conditions in non-small cell lung cancer³⁸. In addition, a previous study showed that *Postn* is secreted from infiltrated inflammatory cells and myofibroblasts in valvular heart disease³⁹. Here, we demonstrated that macrophages, both J774.1 cells and peritoneal macrophages, exposed to hypoxia secreted *Postn*. *Postn* expression were remarkably increased in F4/80⁺ cells from Epi-fat in HFD-fed mice, but it was rarely expressed in SD-fed mice or adipocytes.

It is tempting to speculate that adipose tissue macrophages in obese Epi-fat were recruited by hypoxia-inducible *Postn*. Consistent with this, previous studies have shown that hypoxia increased *Postn* expression and promoted the recruitment of monocyte-derived macrophages from peripheral blood through the integrin $\alpha_v\beta_3$ in a mouse model of glioblastoma³⁰.

Postn is expressed both in infiltrated macrophages and other fibroblast like cells in adipose tissue. However, the contribution of macrophages is poorly understood. To address this question, we used BM transplantation to generate chimeric mice with knockout of *Postn* specifically in hematopoietic cells. Our results demonstrated that *Postn* deficiency in hematopoietic cells suppressed *Postn* expression in Epi-fat and decreased macrophage infiltration and fibrosis after HFD feeding, which was accompanied by a reduction in ectopic lipid accumulation in the liver. These findings were compatible with the whole-body *Postn*^{-/-} phenotypes. However, we could not observe amelioration of insulin resistance in BM-*Postn*^{-/-} mice compared with WT littermates. This may be because of the low increase in body weight in this experiment, possibly caused by the effect of whole-body irradiation.

Our studies demonstrated that the F4/80⁺ SVF also secreted *Postn* after HFD feeding. A recent study showed that numbers of α SMA-positive cells or myofibroblasts existed in obese adipose tissue¹⁰. Both myofibroblast marker (α SMA) and preadipocyte marker (*Pdgfr- α* and *Ppar- γ*) levels were increased in the F4/80⁺ SVF from obese Epi-fat compared with those from lean mice, suggesting that α SMA-positive cells may contribute to the expression of *Postn* in the F4/80⁺ SVF in adipose tissue (Supplementary Fig. 4b).

Postn^{-/-} mice tended to have lower body weight than littermate controls. As reported, *Postn*^{-/-} mice showed reduced body weight compared with WT littermates after weaning because of problems with their teeth^{40,41}. In our case, there was no difference in HFD intake (Supplementary Fig. 2a) possibly because HFD is soft. As a result, there was no difference in body weight between genotypes when IPGTTs and ITTs were conducted. Moreover, our chimeric mice had *Postn* deficiency only in bone marrow-derived hematopoietic cells and there was no difference in body weight between chimeric and control mice. Thus, the phenotypic differences observed in our experiments were due entirely to the presence or absence of *Postn*.

In summary, we demonstrated that *Postn* has a critical role in HFD-induced adipose tissue inflammation and fibrosis, which may reduce lipid storage capacity and enhance ectopic lipid accumulation. Our data indicated that the regulation of *Postn* in visceral fat could be beneficial for the maintenance of healthy adipose tissue in obese patients (Supplementary Fig. 8).

Methods

All data generated or analysed during this study are included in this published article (and its Supplementary Information files).

Animals. Periostin (*Postn*) whole-body knockout mice were generated by Dr. Kudo⁴². We crossed these mice and C57BL/6J mice (purchased from Japan SLC) to generate the *Postn*^{-/-} and *Postn*^{+/+} littermate controls in a C57BL/6J genetic background. HFD (Research Diet, 60% fat) was started at 8 weeks of age and continued for 16 weeks with ad libitum access to food. We generated *Rosa26-loxP-stop-loxP-tdTomato* transgenic mice by crossing with *Periostin-Cre* mice, in which *tdTomato* activates the *Postn*-specific promoter to induce *cre* expression. *Periostin-Cre* mice (generated by Dr. Conway and colleagues^{43,44}) were provided by Dr. Manabe (Chiba University). *tdTomato* mice were obtained from The Jackson Laboratory. Mice were maintained in specific pathogen-free conditions at the Institute of Laboratory Animals of Kyoto University Graduate School of Medicine. This study was approved by the Kyoto University Ethics Review Board. All methods were performed in accordance with the relevant guidelines and regulations.

Histological analysis. Adipose fat tissues and liver were fixed in 4% paraformaldehyde and embedded in paraffin. Areas of fibrosis were measured using image analysis software (Axio Observer 7, ZEISS). CLSs in Epi-fat were detected immunohistochemically using an anti-F4/80 antibody (sc-59171, Santa Cruz Biotechnology) and an anti-perilipin antibody (NB100-60554, Novus Biologicals). The CLS density was obtained from the number of CLSs per 1,000 adipocytes in each section⁴⁵. They were counted in more than 5 mm² areas of each section per mouse, and the number of CLSs was expressed as the mean number per mm².

Confocal microscopic analysis. For confocal microscopic analysis, the adipose tissues from fluorescent reporter mice stained with fluorescent antibodies were visualized in whole mounts, as described previously⁴⁶. In brief, Epi-fat cut into small pieces was fixed 1% paraformaldehyde for 30 min and then permeabilized with 0.5% Triton X-100 prior to staining. The fluorescent stains were examined using an SP8 confocal microscope system (Leica).

Western blotting. Western blotting was performed using standard procedures as described previously⁴⁷. Samples were lysed in lysis buffer consisting of 100 mM Tris-HCl, pH 7.4, 75 mM NaCl, and 1% Triton X-100 (Nacalai Tesque). The lysis buffer was supplemented with complete mini protease inhibitor (Roche), 0.5 mM NaF and 10 mM Na₂VO₄ just before use. The protein concentration was determined using a bicinchoninic acid (BCA) protein assay kit (Bio-Rad). All samples (10 μ g of protein) were suspended in lysis buffer, fractionated using NuPAGE 4–12% Bis-Tris (Invitrogen) gels, and transferred to a Protran nitrocellulose transfer membrane (Whatman). The membrane was blocked using 1 $\times$ phosphate-buffered saline (PBS) containing 5% non-fat milk for 1 h and incubated with the primary antibodies against Periostin/OSF-2 (NBP1-30042, Novus Biologicals), HIF-1 (NB100-479, Novus Biologicals), or GAPDH (14C10, cell signaling) overnight at 4 °C. After a washing step in PBS with 0.05% Tween 20 (0.05% T-PBS), the membrane was incubated with a secondary antibody (anti-rabbit, anti-mouse, or anti-goat IgG horseradish peroxidase (HRP)-linked; 1:2,000) for 1 hour at room temperature. The membrane was then washed in 0.05% T-PBS, and signals were detected using ECL Western Blotting Detection Reagent (GE Healthcare) with an LAS-4000 system (GE Healthcare Life Science).

Western blotting for AKT *in vivo*. After an 18-h fast, mice were injected with insulin (1 U kg^{-1}) or saline via the inferior vena cava. Liver and Epi-fat were removed 1 and 3 min after the injection, respectively. The following antibodies were used: anti-phospho-AKT (Thr308) (#4056, Cell Signaling) and anti-AKT (#9227, Cell Signaling).

Triglyceride content in the liver. Livers were homogenized in ice-cold chloroform/methanol (2:1 v/v). Lipid extracts were prepared by the Folch method⁴⁸. Triglyceride content in the liver was measured using a Triglyceride E-test Wako (Wako Pure Chemical Industries).

Glucose and insulin tolerance tests. For GTTs, after overnight fasting, male mice, after feeding with SD or HFD for 12 weeks and HFD after transplantation for 16 weeks, were injected with 1.0 g kg^{-1} glucose intraperitoneally. For ITTs, after a 4-h fast, male mice at 10 and 16 or 20 weeks of age were injected intraperitoneally with insulin (1 U kg^{-1} for SD-fed and HFD-fed mice after BM transplantation, and 2.0 U kg^{-1} for HFD-fed mice; Humulin R; Eli Lilly Japan KK). Blood was obtained from the orbital vein, and glucose levels were measured using a glucose sensor.

Measurement of serum insulin levels. We quantified serum levels of insulin in male mice after 14 weeks of SD or HFD feeding and after BM transplantation at 18 weeks for HFD feeding using an ELISA assay kit for mouse insulin in accordance with the manufacturer's instructions (Shigayagi Co. Ltd, Shibukawa, Japan).

Biochemical analysis of serum. After mice were fasted for 4–6 h, blood was obtained from the inferior vena cava of anaesthetized mice, and serum was separated by centrifugation at 4°C and stored at -80°C . Biochemical data were measured by standard methods using a Hitachi 7180 Auto Analyzer (Nagahama Life Science Laboratory, Nagahama, Japan).

Stromal vascular fractionation and flow cytometry analysis. We isolated adipocytes and stromal vascular cells as described previously^{49,50}. In brief, we removed Epi-fat tissue and minced it into small pieces in Krebs Ringer phosphate buffer and stirred using a shaker at 100 rpm for 20–30 min in collagenase solution with 3 mg mL^{-1} collagenase type 2 (Worthington). After filtering through nylon mesh, we centrifuged at $300 \times g$ for 10 min and collected the floating adipocyte layer. We resuspended the resultant pellet containing stromal vascular cells and filtered it through a $70\text{-}\mu\text{m}$ mesh. We washed the stromal vascular cells twice with PBS, and finally resuspended them in PBS supplemented with 2% fetal bovine serum (FBS). We added 50 ng of Fcblock antibody per 1×10^6 cells and incubated these isolated cells with either labeled anti-F4/80 (#123107, Biolegend) antibody or isotype control antibody (#400110, Biolegend) and analyzed by flow cytometry using a FACSAria II (BD Biosciences) and FlowJo software (Tomy Digital Biology).

Cell culture and Transwell assays. Macrophage cell line J774.1 were seeded at a density of 10,000 cells/mL in the upper chamber and then allowed to migrate for 48 h before fixation for Geimsa staining. Migration assays were performed in media with 10% FBS or the media with 0.1% bovine serum albumin with Recombinant human POSTN protein (R&D system) 0.05 $\mu\text{g/mL}$ and 0.5 $\mu\text{g/mL}$ or vesicles were measured in the lower chamber. Peritoneal macrophages were obtained from the peritoneal cavity of *Postn*^{+/+} or *Postn*^{-/-} mice 4 days after intra-peritoneal injection of 3 mL of 3% thioglycollate. The cells obtained were washed, spun at 1,000 rpm for 5 min. Transwell assays were performed on 24-well plates with inserts (Millicell). *Postn*^{+/+} or *Postn*^{-/-} peritoneal macrophages were seeded at a density of 10,000 cells/mL in the upper chamber and then allowed to migrate for 48 h before fixation for Geimsa staining. Migration assays were performed in media with 10% FBS or the media with 0.1% bovine serum albumin with TGF β (R&D system), Recombinant human POSTN protein (R&D system) or vesicles were measured in the lower chamber.

Bone marrow transplantation. BM transplantation was conducted as described previously^{51,52}. Eight-week-old female mice with genotypes of *Postn*^{+/+} and *Postn*^{-/-} were used as BM donors. BM recipients were 8 weeks old male B57BL/6J (Japan SLC). BM donors were euthanized by cervical dislocation, and BM cells were collected by flushing femurs and tibias with PBS supplemented with 2% FBS. The suspension was passed through $40\text{-}\mu\text{m}$ mesh. Red blood cells were lysed using ACK lysing buffer (Lonza). BM cells were then washed twice with PBS supplemented with 2% FBS. Recipients were irradiated with two doses of 6 Gy within an interval of 3 hours (cesium 137; Gammacell 40 Exactor) and injected intravenously with 2×10^6 BM cells 6 hours after irradiation. Transplanted mice were housed in microisolator housing for 4 weeks prior to challenge with an HFD.

RNA extraction and qRT-PCR. Total RNA was isolated and purified using TriPure Isolation Reagent (Roche), and cDNA was synthesized from 100 ng of total RNA using a Transcriptor First Strand cDNA Synthesis Kit (Roche) in accordance with the manufacturer's instructions. For quantitative RT-PCR, specific genes were amplified by 40 cycles using SYBRTM Green PCR Master Mix (Applied Biosystems). Expression was normalized to the 18S ribosomal RNA housekeeping gene. Gene-specific primers are listed in Supplementary Table S1.

Statistical analysis. Data are presented as means $\pm$ standard error of the mean (SEM). Statistical comparisons were performed using unpaired t-test or one-way analysis of variance (ANOVA) with Sidak's post-hoc test (three or more groups) were used as indicated in the figure legends. A p value of <0.05 was considered as statistically significant. Statistical analyses were performed using GraphPad Prism 6 (GraphPad Software, Inc.).

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Author Contributions

F.N., M.N. and K.O. designed the project; F.N., M.N., T.H., T.N., T.N. and N.S. performed experiments; F.N., M.N., T.H., H.N., Y.N., O.B., Y.K., T.N., T.N., Y.I., S.K., M.K., S.T., S.Y., S.J.C., M.Y., T.K. and K.O. analysed and interpreted data; S.J.C. and S.Y. contributed materials; and F.N., M.N., and K.O. prepared the manuscript.

Additional Information

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Competing Interests: The authors declare no competing interests.

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Loss of periostin ameliorates adipose tissue inflammation and fibrosis *in vivo*

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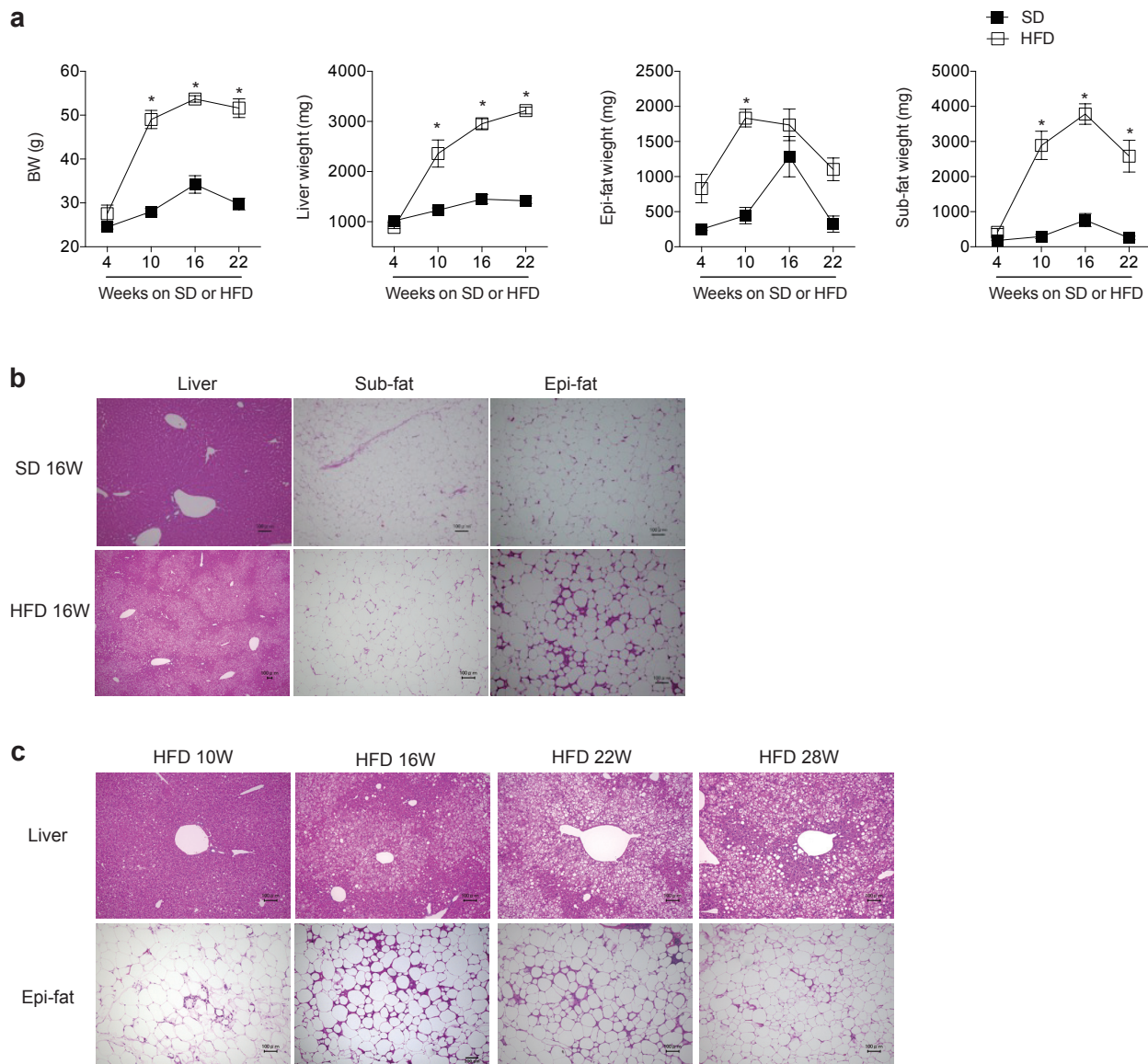
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Supplementary Table S1

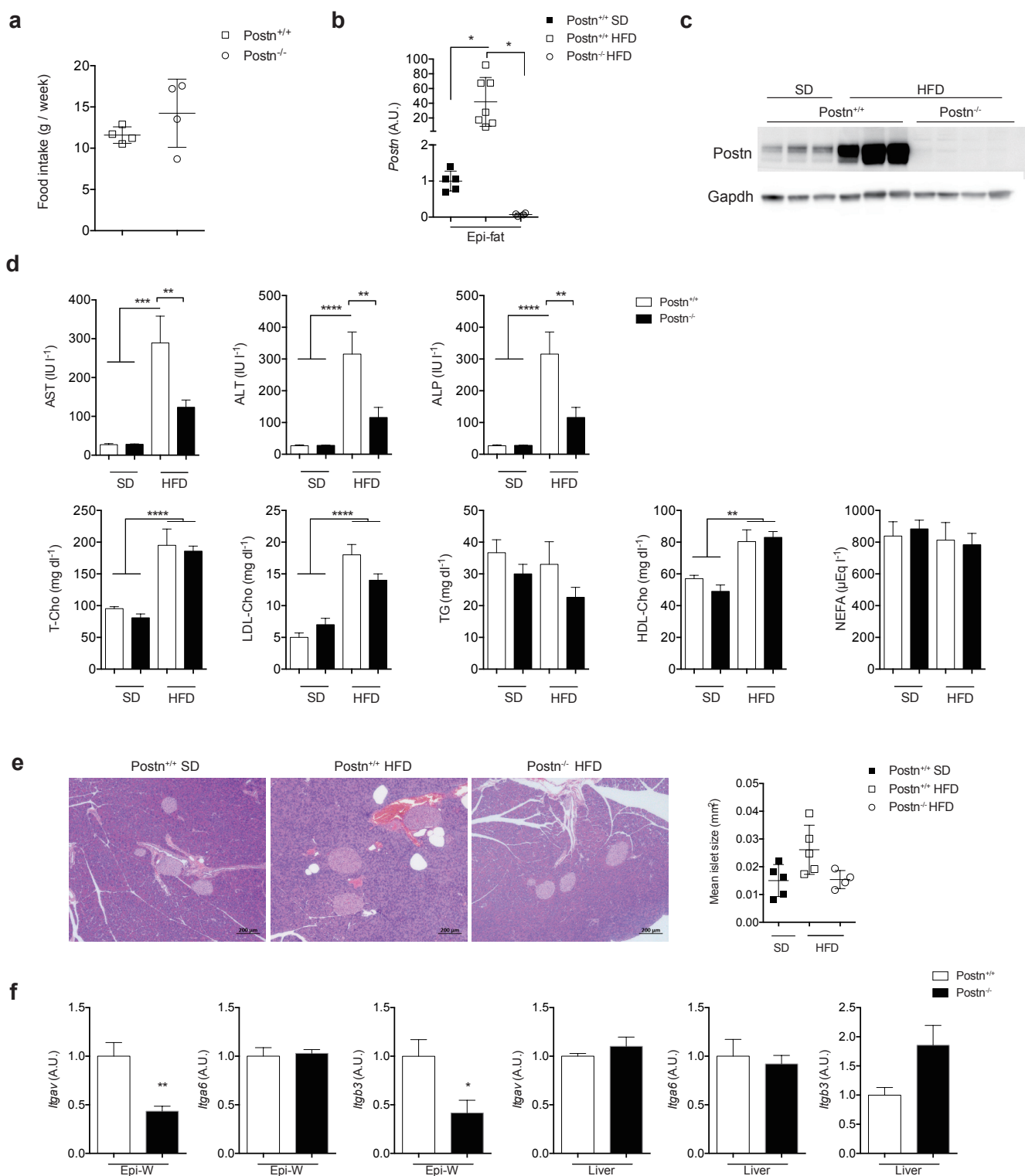
Used gene-specific oligonucleotide primer sequences

Gene	Forward	Reverse
<i>Postn</i>	AACCAAGGACCTGAAACACG	GTGTCAGGACACGGTCAATG
<i>Emr1</i>	TCCAGAAGGCTCCCAAGGATA	GGGCACTTTTGTCTCACAGGTA
<i>Mcp1</i>	AGGTCCCTGTCATGCTTCTG	GCTGCTGGTGATCCTCTTGT
<i>Tnf-α</i>	CCAGACCCTCACACTCAGATC	CACTTGGTGGTTTGCTACGAC
<i>Col1a1</i>	GCCAAGAAGACATCCCTGAAG	TCATTGCATTGCACGTCATC
<i>Col3a1</i>	ACCAAAAGGTGATGCTGGAC	GACCTCGTGCTCCAGTTAGC
<i>Col6a1</i>	TGCAGGCATTGAGATCTTTG	CAGGGCCTGGTAGTTTGTA
<i>Pgk1</i>	TGTCGCTTTCCAACAAGCTG	GGTGGCTCATAAGGACAACG
<i>Glut1</i>	GGGCATGTGCTTCCAGTATGT	ACGAGGAGCACCGTGAAGAT
<i>Vimentin</i>	AGATCGATGTGGACGTTTCC	TCCGGTACTCGTTTGACTCC
<i>α-Sma</i>	TCCCTGGAGAAGAGCTACGAACT	GATGCCCCTGACTCCAT
<i>Pdgfr-α</i>	TCCTTCTACCACCTCAGCGAG	CCGGATGGTCACTCTTTAGGAAG
<i>C/EBP-α</i>	CTGCGAGCACGAGACGTCTA	CTGTCGGCTGTGCTGGAA
<i>Vegf-α</i>	CTGCCGTCCGATTGAGACC	CCCCTCCTTGTACCACTGTC
<i>18S RNA</i>	CGCGGTTCTATTTTGTGGT	AGTCGGCATCGTTTATGGTC



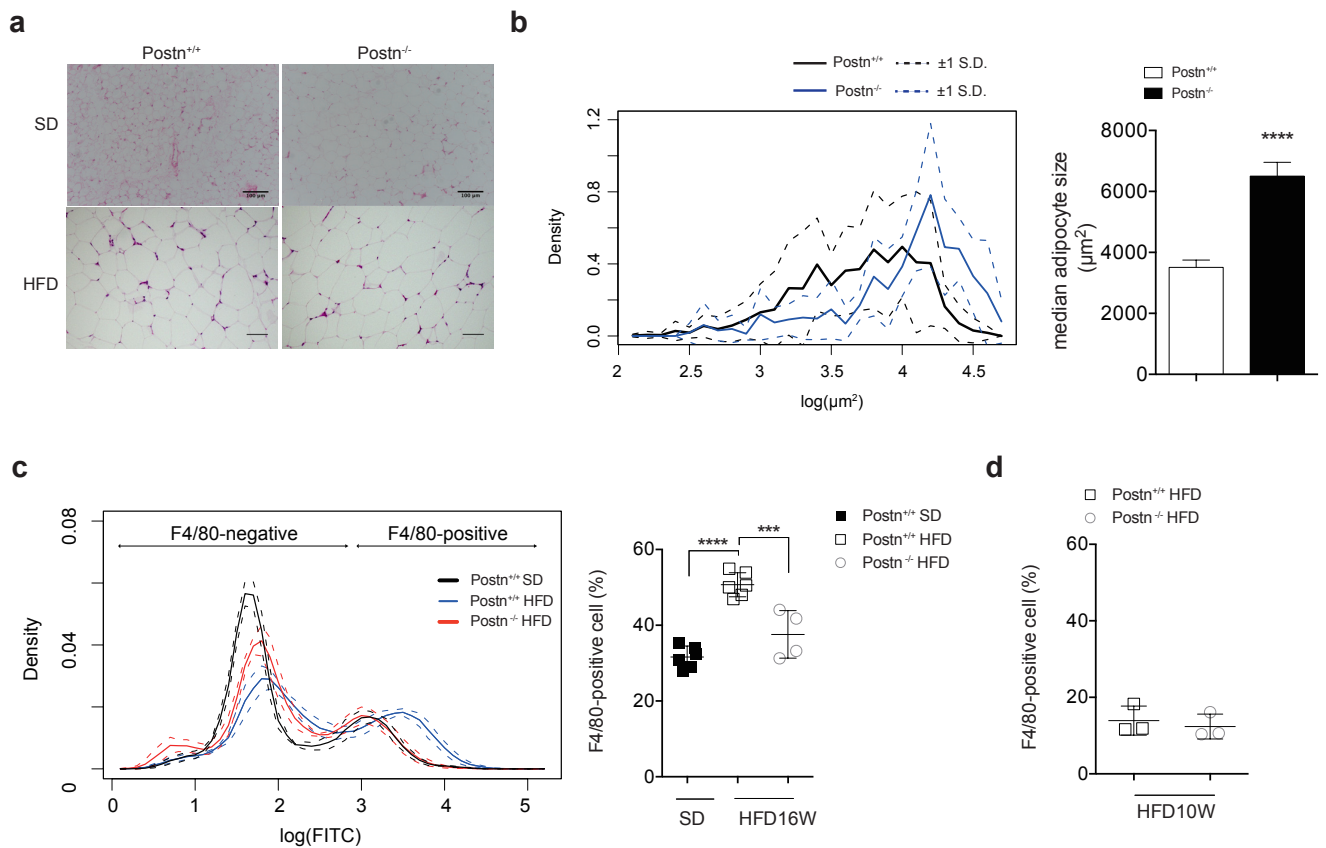
Supplementary Fig. 1. Phenotypes of SD-fed or HFD-fed WT mice.

(a) Time course of body, liver, and adipose tissue weights of WT mice after SD or HFD feeding for up to 30 weeks. $n=3-6$, * $P<0.05$, ** $P<0.01$, *** $P<0.001$, **** $P<0.0001$ versus SD feeding at each time point. Statistical comparisons were made by unpaired t-test. (b) Representative H&E staining images of tissues of WT mice after SD or HFD feeding for 16 weeks. Scale bars, 100 μ m. (c) Representative H&E staining images of tissues of WT mice after HFD feeding for 10, 16, 22 and 28 weeks. Scale bars, 100 μ m.



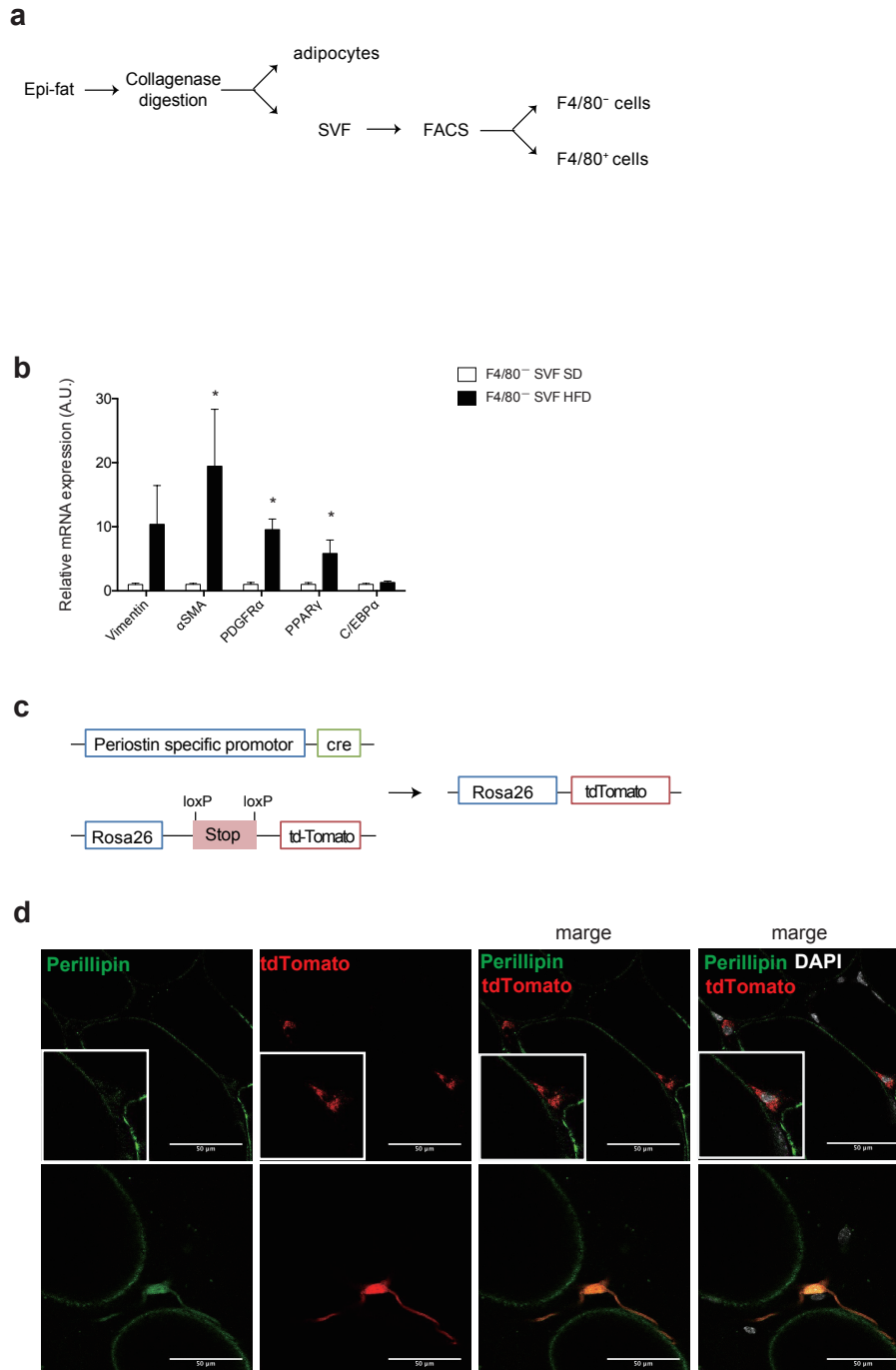
Supplementary Fig. 2. Analysis of *Postn*^{+/+} and *Postn*^{-/-} mice fed a SD or a HFD.

(a) Food intake of *Postn*^{+/+} and *Postn*^{-/-} mice fed HFD in metabolic cages. n=4 each, by unpaired t-test. (b) mRNA levels of *Postn* in adipose tissue (Epi-fat) from *Postn*^{+/+} and *Postn*^{-/-} mice. n=5-7, *P<0.05 by one-way ANOVA. (c) Protein levels of *Postn* in adipose tissue (Epi-fat) from *Postn*^{+/+} and *Postn*^{-/-} mice. (d) Serum profile of *Postn*^{+/+} and *Postn*^{-/-} mice. n=3 for the SD groups, n=8-10 for the HFD groups, **P<0.01, ***P<0.001, ****P<0.0001 by one-way ANOVA. (e) Left, representative microscope images of pancreas from *Postn*^{+/+} and *Postn*^{-/-} mice. Right, the mean islet size in pancreas of *Postn*^{+/+} and *Postn*^{-/-} mice. n=4-5, by one-way ANOVA. (f) mRNA levels of *Ilg* in adipose tissue (Epi-fat) and liver from *Postn*^{+/+} and *Postn*^{-/-} mice. n=5-7, *P<0.05, **P<0.01 by unpaired t-test.



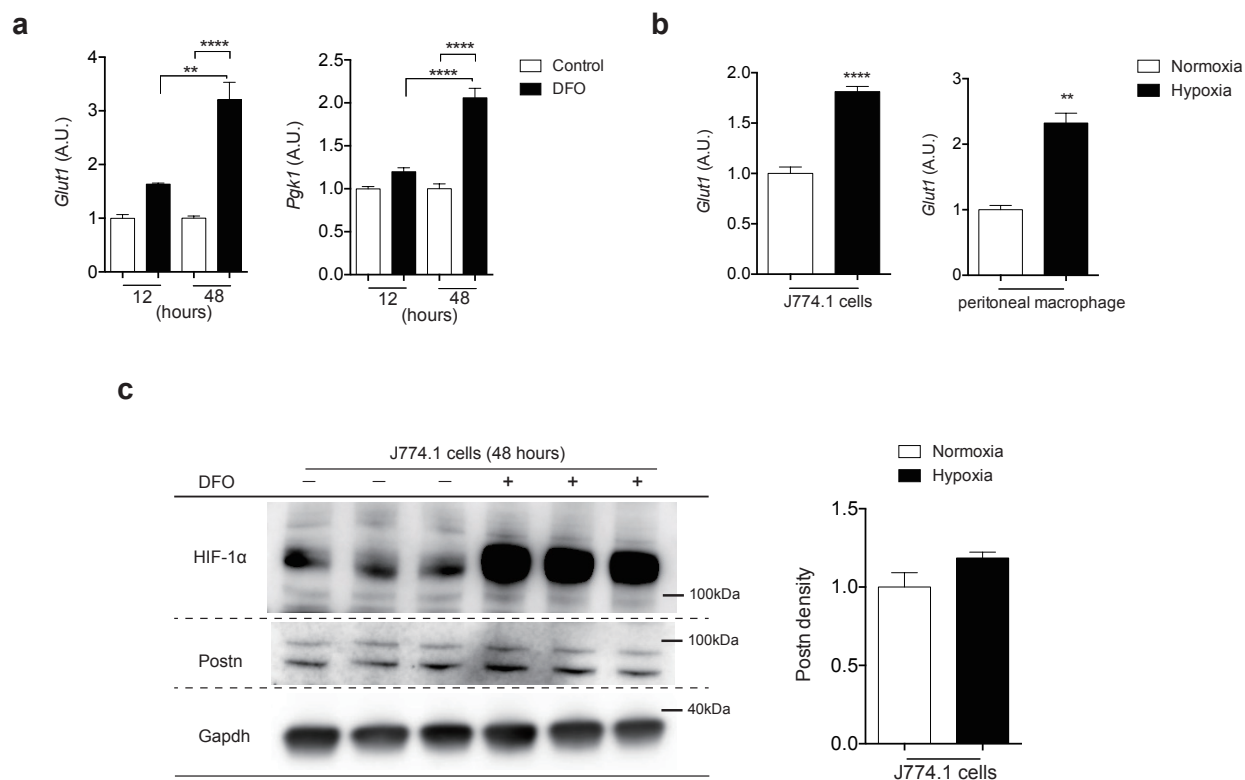
Supplementary Fig. 3. Analysis of the adipose tissues from *Postn*^{+/+} and *Postn*^{-/-} mice.

(a) Representative H&E staining images of the adipose tissue (Sub-fat) of SD-fed and HFD-fed *Postn*^{+/+} and *Postn*^{-/-} mice. (b) Left, density curve of adipocyte surface areas in Sub-fat. Right, the median of adipocyte cell size in Sub-fat of *Postn*^{+/+} and *Postn*^{-/-} HFD-fed mice. ****p*<0.001 by unpaired t-test. *n*=3 images per mouse, 10 mice each. (c) Left, representative flow cytometry plots showing density curves of F4/80⁺ cells isolated from the Epi-fat. Right, quantification of stromal F4/80⁺ cells isolated from Epi-fat of SD-fed and HFD-fed *Postn*^{+/+} mice and HFD-fed *Postn*^{-/-} mice after 16 weeks. *n*=4 to 6, ****p*<0.001, *****p*<0.0001 by one-way ANOVA. (d) Quantification of stromal F4/80⁺ cells isolated from Epi-fat of HFD-fed *Postn*^{+/+} and *Postn*^{-/-} mice after 10 weeks. *n*=3 each, by unpaired t-test.



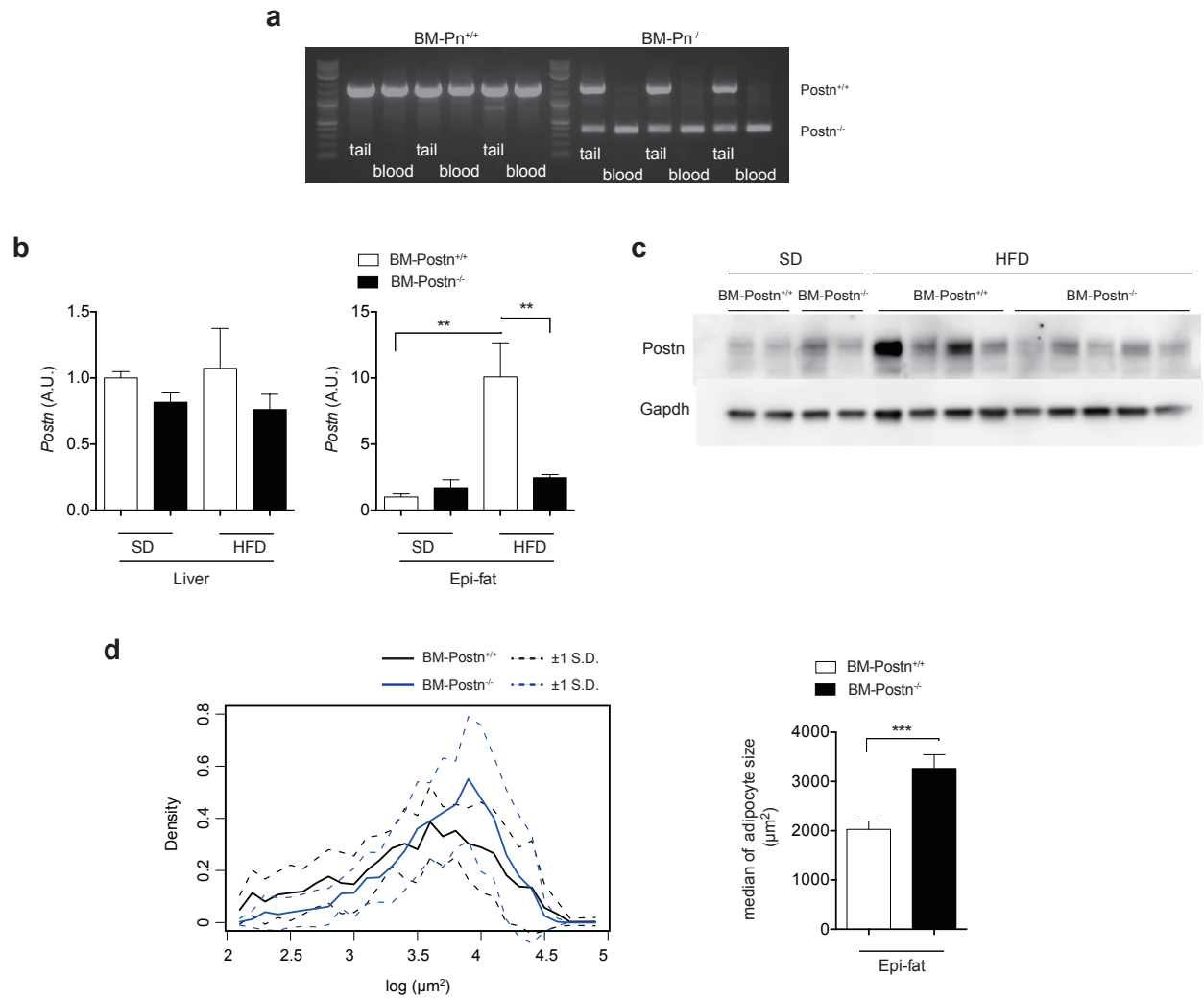
Supplementary Fig. 4. Results of fluorescence-activated cell sorting (FACS) analysis from the SVF.

(a) Illustration of the FACS experiments using SVF. (b) mRNA expression of *Vimentin*, *αSMA*, *Pdgfra*, *Ppar-γ* and *C/EBPα* in F4/80⁻ cells. n=5, *p<0.05 unpaired t-test. (c) Labeling of *Postn*-expressing cells, generated by crossing *Periostin-Cre* with *Rosa26-loxP-stop-loxP-tdTomato* transgenic mice. (d) Representative immunofluorescent staining of perilipin (green) in Epi-fat from obese tdTomato fluorescent protein reporter mice demonstrating *Postn*-expressing cells. Scar bars, 50μm.



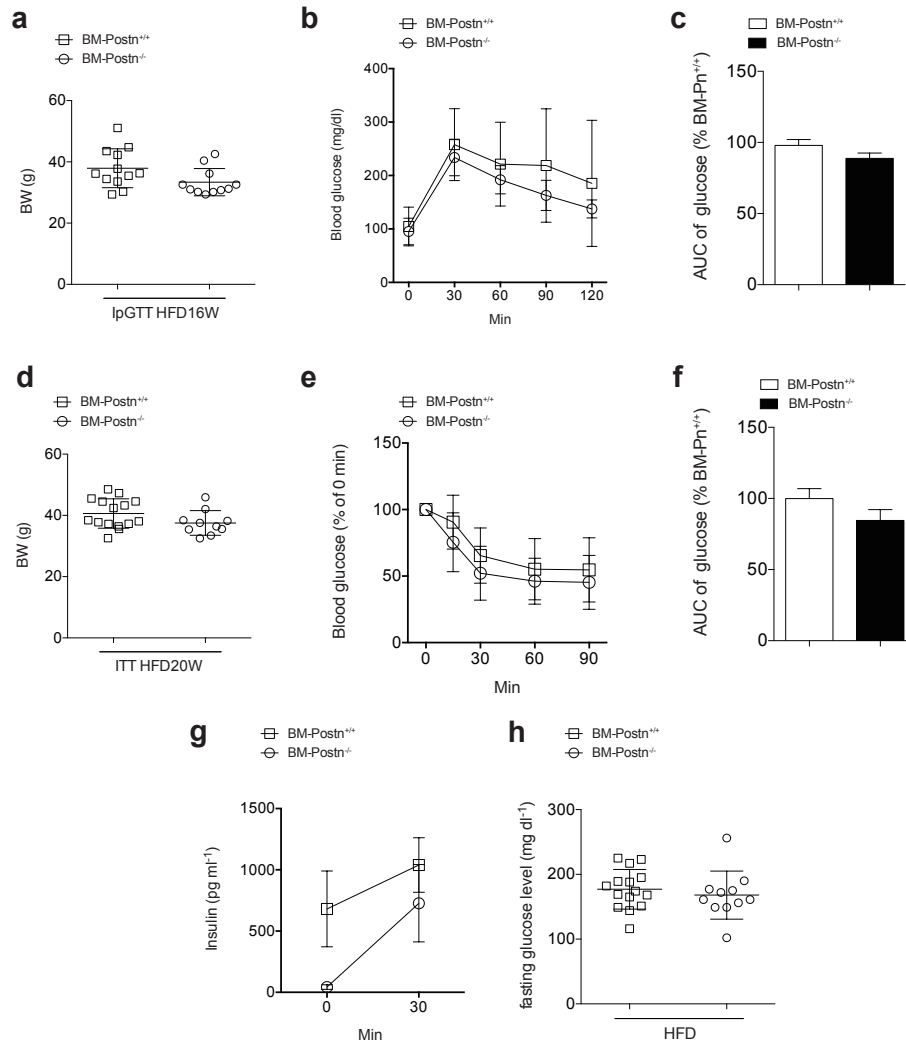
Supplementary Fig. 5. Postn expression under hypoxic condition

(a) mRNA expression of the HIF target genes (*Glut1* and *Pgk1*) in J774.1 cells with or without deferoxamine mesylate (DFO) treatment over the indicated time course. n=6, *p<0.05 by one-way ANOVA. (b) mRNA expression of *Glut1* in J774.1 cells and peritoneal macrophages exposed hypoxia. n=5 for J774.1, n=5-7 for peritoneal macrophages, *p<0.05 by unpaired t-test. (c) Protein levels of Hif-1 α and Postn in J774.1 cells with or without DFO treatments. n=3 each, analyzed by one-way ANOVA.



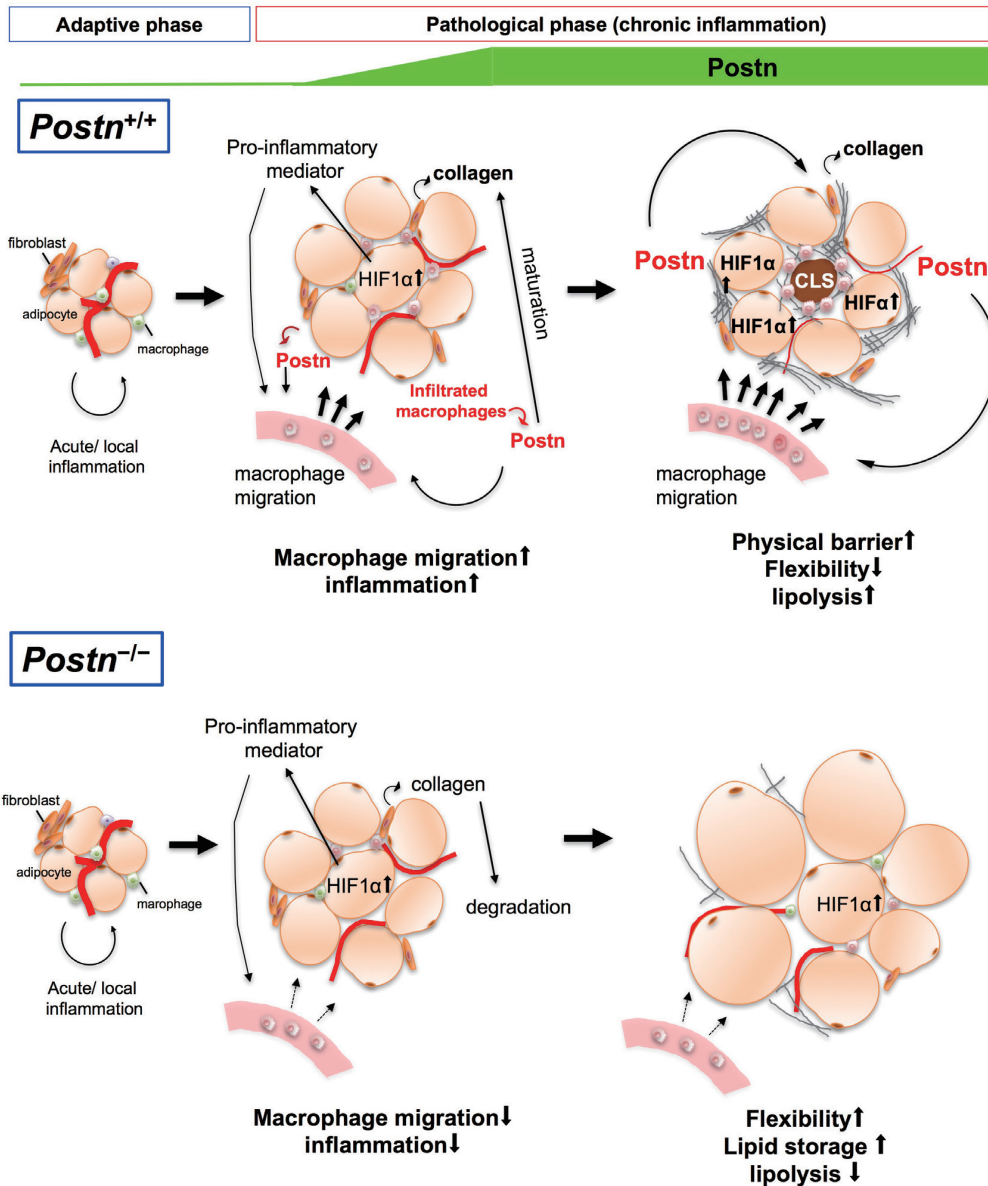
Supplementary Fig. 6. Phenotype of BM transplanted mice.

(a) Gel electrophoresis analysis for the *Postn* genomic locus in tail tissue and peripheral blood cells from BM transplanted mice. (b) mRNA expression of *Postn* in the liver and Epi-fat from BM transplanted mice. n=11-15, **p<0.01 by one-way ANOVA. (c) Protein levels of Postn in Epi-fat of BM transplanted mice. (d) Left, density curve of adipocyte surface areas. Right, the median of adipocyte cell size in Epi-fat of HFD-fed, BM transplanted mice. n=3 images per mouse, 9 mice each, ***p<0.001 by one-way ANOVA.



Supplementary Fig. 7. Analysis of glucose and insulin tolerance in BM transplanted mice.

(a) Body weight of BM transplanted mice used in IPGTTs. $n=11$ each, analyzed by unpaired t-test. (b) Serial changes in glucose levels after intraperitoneal injection of glucose in BM transplanted HFD-fed mice. (c) Area under curve (AUC) of glucose levels after intraperitoneal injection of glucose in BM transplanted HFD-fed mice. $n=11$ each, analyzed by multiple t-test. (d) Body weight of BM transplanted mice in ITTs. $n=11-15$, analyzed by unpaired t-test. (e) Serial changes in glucose levels after intraperitoneal injection of insulin in BM transplanted HFD-fed mice. $n=11-15$, analyzed by unpaired t-test. (f) AUC of glucose levels after intraperitoneal injection of insulin in BM transplanted HFD-fed mice. $n=11-15$, analyzed by unpaired t-test. (g) Serial changes in insulin levels after intraperitoneal injection of glucose in BM transplanted HFD-fed mice. $n=5$ each, analyzed by multiple t-test. (h) Fasting blood concentration of glucose in BM-Postn^{+/+} and BM-Postn^{-/-} HFD-fed mice. $n=11-15$, analyzed by unpaired t-test.



Supplementary Fig. 8. Potential role of Postn in obesity-induced adipose tissue inflammation and fibrosis.

During the development of obesity, Postn is produced in extraordinary amounts by adipose tissue. Postn was secreted at least from macrophages in visceral adipose tissue, possibly in response to hypoxia, and then sustained or amplified inflammation responses. Postn is required for the regulation of ECM production and maturation. Overproduction of ECM may limit the HFD-induced hypertrophy of adipocytes, which has a role in ectopic accumulation in the liver and systemic insulin resistance.