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DOI: [10.1161/ATVBAHA.117.309852](https://doi.org/10.1161/ATVBAHA.117.309852)

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Citation for published version (APA):

Xie, Y., Potter, C. M. F., Le Bras, A., Nowak, W. N., Gu, W., Bhaloo, S. I., Zhang, Z., Hu, Y., Zhang, L., & Xu, Q. (2017). Leptin Induces Sca-1+ Progenitor Cell Migration Enhancing Neointimal Lesions in Vessel-Injury Mouse Models. Arteriosclerosis, Thrombosis, and Vascular Biology, 37(11), 2114. <https://doi.org/10.1161/ATVBAHA.117.309852>

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Leptin Induces Sca-1⁺ Progenitor Cell Migration Enhancing **Neointimal Lesions in Vessel-Injury Mouse Models**

Yao Xie, Claire M.F. Potter, Alexandra Le Bras, Witold N. Nowak, Wenduo Gu, Shirin Issa Bhaloo, Zhongyi Zhang, Yanhua Hu, Li Zhang, Qingbo Xu

- *Objective—Leptin is an adipokine initially thought to be a metabolic factor. Recent publications have shown its roles in* inflammation and vascular disease, to which Sca-1⁺ vascular progenitor cells within the vessel wall may contribute. We sought to elucidate the effects of leptin on Sca-1⁺ progenitor cells migration and neointimal formation and to understand the underlying mechanisms.
- Approach and Results—Sca-1⁺ progenitor cells from the vessel wall of Lepr^{+/+} and Lepr^{-/−} mice were cultured and purified. The migration of Lepr^{+/+} Sca-1⁺ progenitor cells in vitro was markedly induced by leptin. Western blotting and kinase assays revealed that leptin induced the activation of phosphorylated signal transducer and activator of transcription 3, phosphorylated extracellular signal–regulated kinases 1/2, pFAK (phosphorylated focal adhesion kinase), and Rac1 (ras-related C3 botulinum toxin substrate 1)/Cdc42 (cell division control protein 42 homolog). In a mouse femoral artery guidewire injury model, an increased expression of leptin in both injured vessels and serum was observed 24 hours post-surgery. RFP (red fluorescent protein)-Sca-1⁺ progenitor cells in Matrigel were applied to the adventitia of the injured femoral artery. RFP⁺ cells were observed in the intima 24 hours post-surgery, subsequently increasing neointimal lesions at 2 weeks when compared with the arteries without seeded cells. This increase was reduced by pre-treatment of Sca-1+ cells with a leptin antagonist. Guidewire injury could only induce minor neointima in Lepr^{-/−} mice 2 weeks post-surgery. However, transplantation of Lepr^{+/+} Sca-1⁺ progenitor cells into the adventitial side of injured artery in Lepr^{→−} mice significantly enhanced neointimal formation.
- *Conclusions*—Upregulation of leptin levels in both the vessel wall and the circulation after vessel injury promoted the migration of Sca-1+ progenitor cells via leptin receptor–dependent signal transducer and activator of transcription 3- Rac1/Cdc42-ERK (extracellular signal–regulated kinase)-FAK pathways, which enhanced neointimal formation.
- *Visual Overview*—An online visual overview is available for this article. **(***Arterioscler Thromb Vasc Biol***. 2017;37: 2114-2127. DOI: 10.1161/ATVBAHA.117.309852.)**

Key Words: adipokines ■ adventitia ■ leptin ■ mice ■ neointima

besity is associated with a significantly higher risk of cardiovascular disease.¹ The expansion of adipose tissue in obese individuals is closely linked to the secretion of plasma adipokines, which were originally thought only to be related to energy homeostasis.² Among all adipokines, including adiponectin, visfatin, and resistin, leptin was the first to be discovered in 1994.³ Obesity level of individuals strongly correlates with higher levels of plasma leptin, a peptide hormone, mainly secreted into the circulation by white adipose tissue.⁴ Leptin has long been known to play a role in the regulation of food intake and energy expenditure, but recent studies have demonstrated its additional effects on the cardiovascular system, where widespread distribution of OBR (leptin receptor) has been identified.⁵ Leptin

may contribute to atherosclerosis through activation of various mechanisms, including endothelial dysfunction,⁶ lipid metabolism,^{7,8} proinflammatory effect,⁹ and proliferation of smooth muscle cells (SMCs).^{10,11} Shan et al¹² discovered that leptin stimulates proliferation of murine SMCs via the mTOR (mammalian target of rapamycin)-signaling pathway, which may contribute to enhancing neointimal hyperplasia in obese humans. Deletion of either leptin or OBR in leptin-deficient (ob/ob) or leptin receptor–deficient (db/db) mice significantly mitigated the formation of neointima.¹³ The mechanism of leptin-induced neointimal formation after guidewire injury in the femoral artery is thought independent of blood pressure and energy balance.¹⁴ Heart and vascular SMCs are capable of secreting leptin,¹⁵ which can

Received on: June 23, 2017; final version accepted on: September 8, 2017.

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subsequently enhance coronary vasoconstriction and smooth muscle proliferation via the Rho kinase pathway.16 Recent research has demonstrated that leptin induces activation, migration, and proliferation of both endothelial cells and vascular SMCs.17 Leptin may also participate in vascular remodeling and increasing stiffness by altering extracellular matrix production in vascular SMC through the PI3K/Akt (phosphoinositide 3-kinase/protein kinase B [PKB]) pathway.18 Although a significant amount of research has focused on the effect of leptin on SMCs or endothelial cells, its influence on adventitial progenitor cells (APCs) remains unknown.

Accumulating studies have shown that a range of multipotent stem/progenitor cells exist in the adventitia of the vascular wall.19–21 Previous studies in our laboratory have identified the presence of APCs, which are positive for Sca-1 (stem cells antigen-1) and CD34 (hematopoietic progenitor cell antigen) expression.²² This heterogeneous population of cells can give rise to different cell lineages, including $SMCs$ ^{23,24} endothelial cells,^{25,26} and macrophages,^{27,28} which may contribute to neointimal formation.21 Considering the positive correlation between plasma leptin and cardiovascular disease, several laboratories have investigated the biological effects of leptin on the cardiovascular system. However, little is known about whether leptin exerts an effect on APC. We hypothesize that leptin induces the migration of Sca-1⁺ progenitor cells, consequently enhancing neointimal formation. In the present study, we aim to address the role of leptin on Sca-1+ progenitor cell chemotaxis both in vitro and in vivo. We demonstrate that the effect of leptin on induction of progenitor cell migration is mediated by the signal transducer and activator of transcription 3 (STAT3) signaling pathway via OBR. Importantly, our data suggest a clear and novel relationship between plasma leptin, the OBR, and APCs in vascular remodeling after endovascular injury.

Materials and Methods

Materials and Methods are available in the online-only Data Supplement.

Results

Leptin Enhances the Migration of Sca-1+ Progenitor Cells

To investigate whether leptin is involved in the migration of Sca-1+ APCs, we performed an in vitro transwell assay. Cell numbers on the underside of the transwell membrane were counted after staining with crystal violet. We found that treatment with 100 ng/mL of leptin markedly induced the chemotaxis of Sca-1⁺ progenitor cells in a time-dependent way, with a peak of migratory difference after a 16-hour incubation (Figure 1A and 1B). In addition, leptin could markedly induce the chemotaxis of Sca-1+ progenitors in a dose-dependent manner (Figure 1C and 1E), with maximal chemotaxis using 100 ng/mL of leptin after a 16-hour incubation. Similarly, we confirmed that leptin could also induce the migration of Sca-1+ progenitor cells in wound-scratch assays (Figure 1D and 1F). BrdU (bromodeoxyuridine) can be incorporated into the newly synthesized DNA when living cells are proliferating. To exclude the potential effect of cell proliferation in migratory assays, cell proliferation assay using a BrdU kit was performed. Data demonstrated that leptin did not enhance the proliferative ability of Sca-1⁺ progenitor cells (Figure 1G). In summary, leptin can increase the chemotaxis and migration of Sca-1+ progenitor cells in vitro, particularly at a high concentration of 100 ng/mL. Subsequent experiments including G-LISA, quantitative polymerase chain reaction (qPCR), Western blotting, and immunofluorescent staining were therefore performed using a concentration of 100 ng/mL of leptin.

Sca-1+ Progenitor Cells Express OBR Both In Vitro and In Vivo

OBRb is considered to play a role in functionally transducing downstream signaling pathways. As leptin induced the migration of Sca-1+ progenitor cells, the presence of OBR was analyzed in Sca-1+ progenitor cells isolated from the adventitia of the aortas of C57BL/6J mice. We confirmed the expression of OBR in Sca-1+ progenitor cells by PCR (Figure 2A and 2B) and Western blotting analysis (Figure 2C and 2D). The expression of OBR was not upregulated in response to 100 ng/mL of leptin at either RNA or protein level. We also performed immunostaining using CD29 (integrin β-1), CD34 (hematopoietic progenitor cell antigen), CD45 (protein tyrosine phosphatase, receptor type, C), Sca-1, and OBR primary antibodies on the Sca-1+ progenitor cells, which had migrated to the lower side of the membrane after transwell assays. The presence of OBR and Sca-1 (Figure IA in the online-only Data Supplement) was confirmed. Additionally, we discovered that Sca-1⁺ progenitor cells that migrated in response to leptin could express CD29 (Figure IB in the online-only Data Supplement) and CD34 (Figure IC in the online-only Data Supplement) but not CD45 (Figure ID in the online-only Data Supplement). Further immunostaining was performed on mouse tissue. When using OBR and Sca-1 primary antibodies to stain both the aortic (Figure IIA in the online-only

Figure 1. Leptin can induce the migration of Sca-1⁺ adventitial progenitor cells. A, Chemotaxis of Sca-1⁺ progenitor cells at different time points with or without the treatment of 100 ng/mL recombinant leptin was assessed by using a 8.0-µm transwell assay (n=4). **B**, The ratio between control and leptin-treated group was analyzed at each time point (n=4). **C** and **E**, Chemotaxis of Sca-1+ progenitor cells in response to an increasing gradient of leptin in an 8.0-µm transwell system was identified by applying 1% crystal violet staining after 16-h incubation (scale bars, 50 µm; n=6). **D** and **F**, Migration of Sca-1+ progenitor cells in response to an increasing gradient of leptin after 16-h incubation was evaluated in a scratch assay by following crystal violet staining (scale bars, 100 µm; n=6). **G**, Proliferation of Sca-1+ progenitor cells was examined by cell proliferation ELISA, BrdU (bromodeoxyuridine; colorimetric) after 16-h incubation with leptin (n=6). Serum-free cultured medium without leptin treatment was used as a control for the migration assays above. Migration index for transwell and wound-healing assays was defined as the mean ratio of treatment to control of cell numbers counted per 5 random fields at ×20 magnification. Fold change represented the ratio of the number of cells in experimental groups compared with control. All graphs are shown as mean±SEM. **P*<0.05, ***P*<0.01, ****P*<0.001.

Data Supplement) and femoral (Figure IIB in the online-only Data Supplement) arteries of wild-type mice OBR+Sca-1+ cells were observed to be mainly located in the adventitia. These data suggested that Sca-1⁺ progenitor cells expressed OBR. Further phenotyping of Sca-1⁺ adventitial cells by flow cytometry revealed that Sca-1⁺ progenitor cells were positive for CD29, CD34, CD105 (endoglin), CD140a (PDGFR-α [platelet-derived growth factor receptor A]), c-kit, and kinase insert domain receptor, but negative for CD11b (integrin alpha M), CD31 (cluster of differentiation 31), CD45, and CD146 (melanoma cell adhesion molecule; Figures IIIA and IVA in the online-only Data Supplement).

Absence of OBR Does Not Affect the Nature of Sca-1+ Progenitor Cells

To further investigate the role of OBR, we isolated primary aortic adventitial Sca-1⁺ progenitors from Lepr^{-/−} mice (db/ db). PCR was first performed to confirm the genotype of db/ db mice (Figure IVD in the online-only Data Supplement). Immunostaining of femoral arteries from wild-type (Figure VA in the online-only Data Supplement) and db/db (Figure VB in the online-only Data Supplement) mice confirmed that the femoral artery from db/db mice showed no expression of OBR and that Sca-1⁺ cells were similarly distributed in the vascular walls of both db/db and wild-type mice. Although a higher expression of Sca-1 was observed in the adventitia of the arteries from db/db mice, semiquantitative analysis of the numbers of Sca-1⁺ cells showed that there was no significant difference between wild-type and db/db mice (Figure VC in the online-only Data Supplement). The aorta of confirmed db/db mice was harvested for the isolation of Lepr−/− Sca-1+ progenitor cells. To investigate whether the absence of OBR affected the phenotype of Sca-1⁺ progenitor cells, flow cytometry with various hematopoietic and progenitor markers was performed. Data revealed that there was no difference in the expression of any markers between $Lepr^{+/+}$ (Figures IIIA and IVA in the online-only Data Supplement) or Lepr−/− (Figures IIIB and IVB in the online-only Data Supplement)

Figure 2. Leptin induces the activation of phosphorylated signal transducer and activator of transcription 3 (pSTAT3), pERK1/2 (phosphorylated extracellular signal–regulated kinases 1/2), pFAK (phosphorylated focal adhesion kinase), and vinculin in Sca-1+ adventitial progenitor cells. **A** and **B**, The presence of leptin receptor b (OBRb) in Sca-1+ progenitor cells was confirmed by performing conventional polymerase chain reaction (n=4). **C** and **D**, The presence of OBRb in Sca-1+ progenitor cells was confirmed by Western blotting (n=7). **E** and **F**, The activation of pSTAT3 was detected by performing Western blotting on Sca-1⁺ progenitor cells in response to 100 ng/mL of leptin (n=6). **G** and **H**, The activation of pERK1/2 was detected by performing Western blotting on Sca-1+ progenitor cells in response to 100 ng/mL of leptin (n=10). **I** and **J**, Immunofluorescence was performed on Sca-1+ progenitor cells in response to 100 ng/mL of leptin for the detection of pFAK (n=5) and vinculin (n=5). Dashed boxes represent magnified fields. Red arrows indicate the activated proteins (scale bars, 3 and 1 µm). Untreated cells served as a control. Images shown are representative of at least 3 independent experiments.

APC. Both Sca-1+ progenitor cell populations were negative for the hematopoietic markers CD45 and CD11b and endothelial marker CD31 and showed similar staining for CD29, CD34, CD105, CD140a, CD146, c-Kit, and kinase insert domain receptor. Collectively, our results suggest that the absence of OBR does not affect Sca-1⁺ progenitor cells phenotypic characteristics.

OBR-STAT3-MAPK Pathways and Rho GTPase Are Activated in Sca-1+ Cells in Response to Leptin

To understand the underlying mechanism behind Sca-1+ progenitor cell migration in response to leptin, we next sought to elucidate whether OBR and its downstream signaling pathway were involved in this process. Western blot analysis demonstrated that stimulation with 100 ng/mL of leptin led to the activation of pMEK1/2 (phosphorylated mitogen-activated protein kinase kinase 1/2; Figure VI in the online-only Data Supplement), pSTAT3 (Figure 2E and 2F), and pERK1/2 (phosphorylated extracellular signal–regulated kinases 1/2; Figure 2G and 2H), as indicated by phosphorylation at early time points within 5 minutes after the treatment. Rho GTPase family members have been reported to be critical factors for cell migration.29 To elucidate whether RhoA (Ras homolog gene family, member A), Cdc42 (cell division control protein 42 homolog), or Rac1 (ras-related C3 botulinum toxin substrate 1) were involved in Sca-1⁺ progenitor cell leptininduced migration, G-LISA activity assay was performed. Treatment with 100 ng/mL of leptin led to the early activation of Cdc42 (Figure VIIA in the online-only Data Supplement) and Rac1 (Figure VIIB in the online-only Data Supplement)

within 5 minutes, but not of RhoA. In addition, in response to leptin, a relocation of cytoskeleton-related proteins such as phosphorylated FAK (pFAK [phosphorylated focal adhesion kinase]; Figure 2I) and vinculin (Figure 2J) was observed 15 minutes after the treatment, which is indicative of the final stages of cell migration. Taken together, these results implied that leptin could induce the activation of pSTAT3, pMEK, pERK1/2, Cdc42, Rac1, pFAK, and vinculin.

Involvement of OBR, ERK1/2, and STAT3 in Sca-1+ Cell Migration

Our results suggested that OBR, STAT3, and ERK1/2 were involved in leptin-induced signaling in Sca-1+ progenitor cells, but whether this signaling pathway was involved in cell migration and the interactions between these proteins were still unknown. We, therefore, performed transwell, wound-healing, and Western blotting analyses using a knockout cell model and pathway inhibitors. Leptin antagonist (Figure 3A and 3C) and inhibition of STAT3 (Figure 3B and 3D) led to a considerable reduction of Sca1+ cell migration in transwell assays. Consistent with the results above, inhibition of the ERK pathway substantially attenuated cell migration, both in wound-healing (Figure VIIIA and VIIIB in the online-only Data Supplement) and in transwell assays (Figure VIIIE and VIIIG in the onlineonly Data Supplement). Cell migration of Lepr−/− Sca-1+ progenitor cells in both wound-healing (Figure VIIIC and VIIID in the online-only Data Supplement) and transwell assay (Figure VIIIF and VIIIH in the online-only Data Supplement) was significantly reduced in response to 100 ng/mL of leptin when compared with Lepr^{+/+} cells. Western blotting revealed that inhibition of the ERK pathway did not affect the activation of pSTAT3 in response to leptin at an early stage (Figure IXA through IXC in the online-only Data Supplement). The inhibition of the STAT3 pathway markedly decreased the expression of pERK1/2 at an early stage, indicating that pERK1/2 is activated downstream of pSTAT3 in the OBR-mediated response to leptin (Figure IXD through IXF in the online-only Data Supplement). The late activation of pSTAT3 and pERK at 4 hours (Figure IXD through IXF in the online-only Data Supplement) after treatment may be caused by the depletion of STAT3 inhibitor. OBR deficiency prevented the increase of pSTAT3 with the treatment of 100 ng/mL of leptin (Figure IXG and IXH in the online-only Data Supplement), but pERK1/2 was still activated at the late stage (Figure IXG and IXI in the onlineonly Data Supplement), indicating that other receptors or signaling pathway independent of OBR may be involved in leptin-induced cell response.

OBR Is Required for the Migration of Sca-1+ Progenitor Cells In Vivo

We investigated the role of OBR in Sca-1⁺ cell migration in vivo, after a guidewire injury of the femoral artery in mice. Sca-1+ progenitor cells were transfected with RFP (red fluorescent protein)-lentivirus to trace live cell migration in vivo (Figure 3E). The efficiency of transfection was identified using fluorescence-activated cell sorting with an RFP primary

antibody (Figure X in the online-only Data Supplement). Male mice were randomized into 2 groups: one group underwent guidewire injury of femoral artery, followed by transplantation of RFP Sca-1⁺ progenitor cells in Matrigel on the adventitial side of the injured vessel. The other group underwent the same procedure, but CYT-354 (leptin inhibitor) was mixed to the Matrigel. Our data from en face staining demonstrated that Sca-1⁺ progenitor cells could migrate from the adventitia to intima at a very early stage (Figure 3Fa, 3Fb, 3G through 3I). Three days after the endovascular injury, Sca-1⁺ progenitor cells were discovered in the media layer (Figure 4A). When the leptin inhibitor CYT-354 was added in Matrigel, Sca-1+ progenitor cells migration was significantly reduced, indicating the substantial effect of leptin and OBR on cell migration in vivo (Figure 3Fc, 3Fd, 3G, 3H, and 3J). We also transplanted Sca-1+ progenitor cells to the adventitial side of femoral arteries in db/db mice. Surprisingly, the migratory cells were also observed in injured arteries 3 days post-surgery (Figure 4B, 4D, and 4F) although the number of migratory cells was reduced compared to what we observed in wild-type mice (Figure 4F). Additionally, a population of migratory RFP Sca-1+ progenitor cells acquired SMC markers (Figure 4D) and lost Sca-1 marker expression (Figure 4E) during the migration, revealing that these cells may undergo differentiation in vivo before they reach intima. We next applied Lepr−/− Sca-1+ progenitor cells on the injured arteries of wild-type mice and found that very few RFP⁺ cells had migrated inside the artery (Figure 4C and 4F). These results demonstrated that the OBR on Sca-1+ progenitor cells plays a key role in cell migration from adventitia to intima after endothelial injury. Indeed, lack of OBR substantially diminished cell migration.

Expression of Leptin in the Femoral Artery and Serum Leptin Levels Were Upregulated Post-Surgery

Data presented to date indicated that Sca-1⁺ progenitor cells can migrate into the injured artery and that the absence of OBR can mitigate this effect. However, the factor responsible for Sca-1+ progenitor migration into the intima was still unknown. Our data revealed that serum leptin reached a high level 1 day post-surgery (Figure 5A). Db/db mice showed a much higher concentration of serum leptin (Figure 5B), which may explain the Sca-1⁺ cell migration we observed in db/db mice (Figure 4B). In addition, immunofluorescence also indicated a higher expression of leptin in cells of the vessel wall (Figure 5F) 1 day post-surgery compared with the expression in an uninjured artery (Figure 5E). Western blotting for the whole artery revealed that expression of leptin was upregulated 1 day after the guidewire injury (Figure 5C and 5D). We also confirmed that SMCs could express leptin in vitro under normal culture conditions (Figure 5G). These data, taken together, demonstrate that the expression of leptin was enhanced both systemically and locally after the endovascular injury.

Migration of Sca-1+ Progenitor Cells Contribute to Neointimal Formation

To explore the long-term role of Sca-1+ progenitor cells in neointimal formation, guidewire injury was performed in

Figure 3. Leptin-induced migration can be abolished by CYT-354 treatment both in vitro and in vivo. **A** and **C**, **B** and **D**, Chemotaxis of Sca-1+ progenitor cells in response to 100 ng/mL of leptin with CYT-354 (**A**; n=5) or WP1066 (**B**; n=5) in an 8.0 µm transwell system was identified using 1% crystal violet staining after 16-h incubation (scale bars, 20 µm). Serum-free medium and dimethyl sulfoxide (DMSO) were used as controls. Migration index of transwell assays was defined as the mean ratio of treatment to control of cell numbers counted per 5 random fields at ×20 magnification. E, Sca-1+ progenitor cells were successfully transfected by RFP (red fluorescent protein) lentivirus (scale bars, 40 µm). **F**, 1×10⁶ RFP Sca-1⁺ with (c and d) or without CTY-354 (a and b) were seeded in the adventitial (*Continued*)

wild-type mice. Wild-type mice in the injury model only developed moderate neointimal formation 2 weeks after the injury, resulting in minor artery narrowing (Figure XIA and XID in the online-only Data Supplement). Neointimal lesions grew significantly 4 weeks post-surgery along with a significant reduction in media area (Figure XIE through XIH in the online-only Data Supplement). The intimal layer was increased markedly in both size and thickness, resulting in a high intima-media ratio 4 weeks after injury (Figure XIG in the online-only Data Supplement). Immunohistochemistry (Figure XIB in the online-only Data Supplement) and immunofluorescence (Figure XIC in the online-only Data Supplement) for $α$ -SMA ($α$ -smooth muscle actin) showed that the main component of neointimal formations was SMCs. Hyperleptinemia has been related to the recruitment of leukocytes and macrophages to the endothelial wall at the early stage of atherosclerosis.30 Immunostaining for F4/80 revealed that macrophages started to infiltrate the injured artery from 5 days postsurgery (Figure XIIC in the online-only Data Supplement). Moreover, macrophages were seen to participate in the formation of neointima 7 days post-surgery in wild-type mice (Figure XIID in the online-only Data Supplement). The data from wild-type mice with guidewire injury only were used as a surgery control for further comparison to other surgical groups. Transplantation of Sca-1⁺ progenitor cells significantly augmented neointimal formation in wild-type mice 2 weeks post-surgery (Figure 6Ad, 6B, 6F through 6I) compared with the mice with injury only (Figure 6I). This phenomenon could be abolished when the cells were transplanted with 1000 ng/mL of CYT-354 (Figure 7Aa, 7Ac, 7F through 7I). There was no difference in the adventitial areas between the groups with or without transplantation of exogenous cells (Figure 6E). Van-Gieson staining demonstrated that transplantation of exogenous Sca-1+ progenitor cells did not affect the extent of fibrosis (Figure XIII in the online-only Data Supplement). In short, transplanted Sca-1+ progenitor cells could significantly enhance neointimal formation 2 weeks after surgery, but the administration of CYT-354 inhibited the formation of neointima (Figure 6I), indicating that leptin and OBR in Sca-1⁺ progenitor cells are crucial in neointimal formation.

Wire Injury Induces Neointimal Formation in Wild-type But Not in db/db Mice

We performed guidewire injury of femoral arteries in db/db mice. The injured arteries were collected 2 or 4 weeks after surgery. The size of neointimal formation measured in db/db mice was greatly reduced 2 weeks after the injury (Figure 7A) although they had a higher concentration of leptin in the circulation (Figure 5B), implying that the absence of the OBR may serve as a protective role in vascular remodeling.

Neointimal Accumulation of Sca-1+ Progenitor Cells Is Mediated by OBR

To better understand the specific role of OBR expressed on Sca-1⁺ progenitor cells in neointimal formation, the guidewire injury mouse model was performed in wild-type and db/ db mice. Briefly, experiments were performed on 2 groups of animals: one group of wild-type mice underwent guidewire injury, followed by a transplantation of 1×10^6 Lepr^{-/-} Sca-1⁺ progenitor cells on the adventitial side of the injured artery and another group of db/db mice underwent the same procedure but received 1×10^6 Lepr^{+/+} Sca-1⁺ progenitor cells. The injured arteries were harvested 2 weeks after the surgery. Our data revealed that in db/db mice receiving Lepr^{+/+} Sca-1⁺, there was a significant increase in neointimal formation (Figure 7B) compared with the group with injury only (Figure 7A and 7E). Lepr−/− Sca-1+ cells (Figure 7D) could not enhance neointimal lesion in wild-type mice compared with the group with injury only and to the group with $Lepr^{+/+}$ Sca-1⁺ cells (Figure 7C and 7F). Immunofluorescence for SMC and endothelial cell markers showed the composition of Sca-1+-induced neointimal formations (Figure 7B and 7C). Interestingly, there was a thick layer of CD31+ cells in db/db mice 2 weeks after the vessel injury, which may serve as a protective role against neointimal formation (Figure 7A). Taken together, our data indicate that the expression of OBR on Sca-1+ progenitor cells is crucial for the formation of neointima.

Neointima in Db/Db Mice Partly Originated From Exogenous Sca-1+ Progenitors

When Lepr^{+/+} Sca-1⁺ progenitors were transplanted to db/db mice, an apparent neointima formation could be detected. Because db/db arteries were unable to develop any neointimal lesions after guidewire injury only, the origin of neointima in the db/db mice after cell transplantation was of interest. Immunostaining did not show any positive expression of OBR in the femoral artery (Figure IIC in the online-only Data Supplement). When L epr^{+/+} Sca-1⁺ cells were applied in db/db mice, immunostaining of OBR revealed a significant number of Lepr+ cells in the neointima, indicating that they originated from transplanted Lepr+/+ Sca-1+ progenitor cells (Figure 7G). It is worth noting that not all neointimal cells were OBR positive and most of the OBR-positive cells did not coexpress SMC markers. Further immunostaining with a CD45 primary antibody showed positive staining in the neointimal lesion but did not costain with OBR, suggesting that both adventitial and hematopoietic cells contribute to vascular remodeling (Figure XIV in the online-only Data Supplement). The origins of other OBR negative cells and the roles of OBR-positive cells in neointima in db/db mice will need further investigation. When Lepr−/− Sca-1+ cells were applied in wild-type mice, immunostaining revealed that OBR was expressed in all 3 vessel layers,

Figure 3 Continued. side of injured femoral arteries (n=6). En face staining showed that the RFP cells migrated from the adventitia to the intima on 1 or 3 d post-surgery (scale bar, 100 µm). **G**, Quantification of migratory cells from the adventitia to intima with or without CYT-354 inhibitor on 1 d post-surgery. **H**, Quantification of migratory cells from the adventitia to intima with or without CYT-354 inhibitor on 3 d post-surgery. **I**, Quantification of migratory cells from the adventitia to intima without CYT-354 inhibitor on 1 or 3 d post-surgery. **J**, Quantification of migratory cells from the adventitia to intima with CYT-354 inhibitor on 1 or 3 d post-surgery. Cell migration assay in vivo was performed in wild-type mice. All graphs are shown as mean±SEM. **P*<0.05, ***P*<0.01, ****P*<0.001.

Figure 4. Lack of leptin receptor in Sca-1+ progenitor cells abolished the leptin-induced cell migration in vivo. **A**, Cell migration assay in vivo was performed in wild-type mice. 1×10⁶ RFP (red fluorescent protein) Sca-1⁺ were seeded in the adventitial side of injured femoral artery in wild-type mice. En face staining showed that the RFP cells migrated toward the intima at 72 h after the surgery (*Continued*)

which was considered as an intrinsic expression in wild-type mice and neointima could not be increased (Figure 7H).

Discussion

In the present study, we investigated the relationship between leptin and resident stem cells in vascular remodeling. We demonstrate that leptin is a chemokine for stem/progenitor cell migration mediating neointimal formation in damaged vessels, which can be abrogated by inhibition or knockout of the OBR. Modulation of the signaling pathways involving OBR-STAT3-Rac/Cdc42-FAK influences cell motility in vitro and vascular remodeling in vessel-injury models. Importantly, we showed that significant neointimal formation could develop in db/db mice transplanted with exogenous Sca-1+ cells, which may indicate a fundamental role for OBRb and Sca-1+ progenitors in vascular remodeling. Our novel findings suggest an interaction between leptin and vascular resident stem cells in the pathogenesis of vascular disease.

Obesity leads to changes in the levels of plasma adipokines such as adiponectin, leptin, visfatin, and resistin, all of which may influence the cardiovascular system. To determine whether Sca-1⁺ progenitors could respond to these adipokines, migratory assays including transwell (Figure XVA in the online-only Data Supplement) and wound-healing (Figure XVB in the online-only Data Supplement) in response to different adipokines were performed. We confirmed that only resistin and leptin could induce the migration of Sca-1⁺ progenitor cells. OBRb is considered to be the functional receptor that mediates most of the biological leptin-induced effects.³¹ OBR is expressed in multiple cell types.32 Single-cell analysis from previous study in our laboratory demonstrated that Sca-1⁺ progenitor cells express long-form OBR,³³ followed by confirmation with PCR, Western blotting, and immunostaining assays. The action of OBR is commonly induced by JAK (Janus kinase)/STAT,³⁴ AMPK (5' AMP-activated protein kinase)³⁵ PI3/Akt, and MAPK (mitogen-activated protein kinase)³⁶ pathways. Consistent with previous studies, leptin could induce the activation of phosphorylated STAT3, MEK1/2, and ERK1/2 at an early stage. Either inhibition of OBR, STAT3, or ERK1/2 led to a reduction in cell migration in vitro. However, a late activation of pERK in Lepr−/− cells indicated that other signaling pathways independent of OBR may also be involved. We hence performed a qPCR for the expression of cell migration-related receptors such as C-C chemokine receptor type 1, 2, 7, 9 and CXCR (C-X-C chemokine receptor type) 3, 4, 5 as it was shown previously that they were upregulated in Sca-1⁺ progenitor cells during migration.24 Surprisingly, the gene expression of CXCR5 (Figure XVI in the online-only Data Supplement) was upregulated 24 hours after the treatment of leptin, indicating its potential

role in leptin-induced signaling pathways. We also performed MAPK protein arrays on leptin-stimulated Sca-1⁺ progenitors. Various phosphorylated MAPK proteins were activated (Figure VIIC in the online-only Data Supplement). However, the potential interactions between CXCR5, MAPK pathway, and leptin still need further investigation. The Rho GTPase family including Rac1, Cdc42, and RhoA is known to regulate the formation of lamellipodia, filopodia, and focal adhesions.³⁷ In our study, Rac1 and Cdc42, commonly considered to act upstream of MAPK pathways, were activated by 100 ng/mL leptin, followed by an enhanced rearrangement of cytoskeleton-related proteins such as phosphorylated FAK and vinculin. Taken together, we provide robust data to identify OBR-STAT3- Rac1/Cdc42-ERK-FAK as the signaling pathway (Figure XX in the online-only Data Supplement) involved in Sca-1⁺ progenitor cell migration in response to leptin, implicating new potential therapeutic targets for vascular disease.

After the endovascular injury, an inflammatory response takes place in the vessel wall, with the release of chemokines secreted by SMCs and mononuclear cells.³⁸ We showed that the expression of leptin in both serum and vessel wall was upregulated 1 day post-surgery, which might induce the migration of Sca-1+ progenitor cells from the adventitia to the intima. In addition, we also counted the numbers of Sca-1+ APCs before and 2 weeks after the vessel injury. Interestingly, db/db mice showed a reduced ability to maintain the adventitial progenitors after the injury (Figure XVIII in the online-only Data Supplement). Using a RFP labeling/ tracing technique, migratory Sca-1+ cells could be traced on the inner side of the media layer of SMCs. On the other hand, Lepr−/− cells or Lepr+/+ treated with CYT-354 lost their migratory abilities. However, when Lepr^{+/+}Sca-1⁺ progenitor cells were transplanted on injured arteries of db/db mice, the migration of the progenitor cells was not significantly altered. Therefore, progenitor cell migration in vivo is largely dependent on the existence of OBRs on the cell surface. Moreover, Lepr+/+Sca-1+ progenitor cells could acquire SMC markers in vivo 3 days post-surgery. Quantitative RT-PCR in Sca-1+ cells showed an upregulation of SMC markers in response to 100 ng/mL of leptin for 24 hours, suggesting that leptin may have a potential role in the differentiation of the progenitors toward the SMC fate in vivo (Figure XVII in the online-only Data Supplement). Further investigation will be needed to confirm a link between leptin and Sca-1+ progenitor cells' differentiation.

The origin of the proliferative SMCs accumulating during neointimal formation is still under debate. Fibroblasts,³⁹ SMCs, and APCs⁴⁰ are all thought to participate in neointimal formation. Leptin has been reported to promote

Figure 4 Continued. (scale bar, 30 µm; n=6). B, 1×10⁶ RFP Sca-1⁺ were seeded in the adventitial side of injured femoral artery in leptin receptor–deficient (db/db) mice. En face staining showed that the RFP cells migrated toward the intima at 72 h after the surgery (scale bar, 100 µm; n=6). C, 1×10⁶ RFP lepr^{-/-} Sca-1+ were seeded in the adventitial side of injured femoral artery in wild-type mice. En face staining showed that the RFP cells did not migrate toward the intima at 72 h after the surgery (scale bar, 50 µm; n=6). **D**, 1×10⁶ RFP Sca-1+ cells were seeded in the adventitial side of injured femoral artery in db/db mice. En face staining showed that the RFP cells acquired smooth muscle cell marker (Alexa 488; green) at 72 h after the surgery (scale bar, 20 µm; n=6). **E**, 1×10⁶ RFP Sca-1+ cells were seeded in the adventitial side of injured femoral artery in wild-type mice. En face staining showed that some RFP cells lost Sca-1+ marker (Alexa 488; green) at 72 h after the surgery (scale bar, 5 µm; n=6). **F**, Quantification of migratory cells from the adventitia for group A, B and C. All graphs are shown as mean±SEM. **P*<0.05, ***P*<0.01, ****P*<0.001.

Figure 5. The expression of leptin was upregulated after surgery both in the circulation and in the injured artery. **A**, Serum leptin in wild-type mice 1 (n=10), 7 (n=5) and 14 (n=12) d after injury were quantified by using leptin ELISA kit. **B**, Serum leptin in wild-type and db/db mice was quantified by using leptin ELISA kit. **C**, Expression of leptin in the injured or noninjured arteries was documented by performing Western blotting 1 d post-surgery (n=4). **D**, Quantification of leptin expression in noninjured and injured vessels. **E** and **F**, Difference in expression of leptin (Alexa 488; green) in noninjured (**E**) artery (n=6) and injured (**F**) artery (n=10) was analyzed on 1 d post-surgery by immunofluorescence (scale bars, 30 µm). **G**, Expression of leptin in smooth muscle cell (Alexa 594; red) in vitro was detected by immunofluorescence (scale bars, 8 µm). DAPI indicates 4',6-diamidino-2-phenylindole for nucleus staining; and EC, endothelium. All graphs are shown as mean±SEM. **P*<0.05, ***P*<0.01, ****P*<0.001.

integrin-mediated adhesion, recruiting endothelial progenitor cells into neointima after vessel injury.⁴¹ Db/db mice displayed a phenotype of obesity, higher systemic arterial blood pressure, depressed heart rates, hyperlipidemia, severe

hyperglycemia, hyperinsulinemia, and pancreatic dysfunction.42,43 We also confirmed that db/db mice had a higher serum leptin. Surprisingly, no neointimal formation could be observed in db/db mice at 2 or 4 weeks post-operation.

Figure 6. CYT-354 can significantly inhibit neointimal formation 2 wk after endovascular injury. Sca-1+ progenitor cells (1×10^e) were seeded on the adventitia of previously injured vessels and allowed to migrate with or without CYT-354 in the Matrigel. **A**, Paraffin sections of injured femoral arteries were observed by performing H&E staining (scale bars, 50 and 10 µm; n=6). **B**, Quantification of neointimal area of injured arteries (n=6). **C**, Quantification of media area of injured arteries (n=6). **D**, Quantification of adventitial area of injured arteries (n=10). **E**, Quantification of adventitial area of injured arteries of the groups with or without receiving exogenous cells (n=10 and 6). **F**, Quantification of ratio of media to adventitia of controls and injured arteries (n=10). **B**, Quantification of ratio of neointima to media of injured arteries (n=6). **G**, Quantification of ratio of neointima to adventitia of injured arteries (n=6). **H**, Quantification of ratio of media to adventitia of injured arteries (n=6). **I**, Quantification of ratio of neointima to media of injured arteries, injured arteries with transplanted cells, and injured arteries with CYT-354 (n=6). APC indicates adventitial progenitor cell; M, media; and Neo, neointima. Graphs are shown as mean±SEM. Dashed boxes represented the magnified field. **P*<0.05, ***P*<0.01, ****P*<0.001.

Additionally, there was a thick layer of CD31⁺ cells in the intima at 2 weeks post-surgery. Interestingly, application of Lepr^{+/+}Sca-1⁺ progenitors on the adventitial side of the injured artery in db/db mice resulted in marked neointimal formation 2 weeks post-surgery. Previous reports have suggested that the lack of neointimal formation in db/db mice was mainly because of the inhibition of proliferation of SMCs.⁴⁴ We verified this finding by performing BrdU proliferation assays for SMC and progenitor cells with or without OBR. Data showed that the absence of OBR could significantly decrease the proliferative abilities of both SMC and progenitors (Figure XIX in

the online-only Data Supplement). Importantly, we also demonstrated that the inhibition of migration of Sca-1+ progenitor cells is of importance. A significant neointimal formation was detected 2 weeks post-surgery in db/db mice, caused by the transplantation of exogenous Sca-1+ progenitors. However, db/db mice that did not receive the exogenous cells did not develop any neointimal formation. Because of the fluorescence quenching of RFP cells, we were unable to track the cells in vivo 2 weeks post-surgery. Instead, we stained a cross section of db/db mouse tissue with OBR primary antibody after the transplantation of lepr^{+/+} Sca-1⁺ cells. Surprisingly,

Figure 7. Neointimal effect of Sca-1+ progenitor cells is mediated by leptin receptor. **A**, Cross sections of femoral artery from leptin receptor–deficient (db/db) mice at 2 wk after the surgery were analyzed by hematoxylin-eosin staining and immunofluorescence for α-SMA (α-smooth muscle actin) and CD31 (cluster of differentiation 31; scale bars, 50 and 10 µm; n=6). **B**, Lepr+/+ Sca-1+ progenitor cells were seeded on the adventitial side of injured artery in db/db mice. Cross sections of femoral artery from db/db mice at 2 (*Continued*)

Figure 7 Continued. wk after the surgery were analyzed by H&E staining and immunofluorescence for α-SMA and CD31 (scale bars, 50 and 10 um; n=5). C, Lepr^{+/+} Sca-1⁺ progenitor cells were seeded on the adventitial side of injured artery in wild-type mice. Cross sections of femoral artery from wild-type mice at 2 wk after the surgery were analyzed by H&E staining and immunofluorescence for α-SMA and CD31 (scale bars, 50 and 10 µm; n=8). **D**, Lepr−/− Sca-1+ progenitor cells were seeded on the adventitial side of injured artery in wild-type mice. Cross sections of femoral artery from db/db mice at 2 wk after the surgery were analyzed by H&E staining and immunofluorescence for α-SMA and CD31 (scale bars, 50 and 10 µm; n=6). **E,** Quantification of the ratio of media to adventitia of injured arteries for group A and B. **F**, Quantification of the ratio of media to adventitia of injured arteries for group C and D. G, Lepr^{+/+} Sca-1+ progenitor cells were seeded on the adventitial side of injured artery in db/db mice. Cross sections of femoral artery from db/db mice at 2 wk after the surgery were analyzed by immunofluorescence for α-SMA and leptin receptor (OBR; scale bars, 50 and 10 μm; n=5). **H**, Lepr^{-/-} Sca-1⁺ progenitor cells were seeded on the adventitial side of injured artery in wild-type mice. Cross sections of femoral artery from wild-type mice at 2 wk after the surgery were analyzed by immunofluorescence for α-SMA and OBR (scale bars, 50 and 10 µm; n=5). APC indicates adventitial progenitor cell; DAPI, 4',6-diamidino-2-phenylindole for nucleus staining; LEPR, leptin receptor; M, media; and Neo, neointima. Graphs are shown as mean±SEM. Dashed box represented the magnified field. Arrows indicated the smooth muscle cell and endothelial cell, respectively. **P*<0.05, ***P*<0.01, ****P*<0.001.

many neointimal cells were OBR positive, proving that they originated from the transplanted Sca-1+ progenitors. Because OBR⁺ cells only represented part of the neointimal cells, most of which did not coexpress SMC marker, the origin of the remaining cells is unknown and needs further investigation. In contrast, wild-type mice that received Lepr−/− Sca-1+ cells only showed few OBR⁺ cells and smaller neointima. Because of technical limitations, we were unable to track all RFP Sca-1+ progenitors in vivo in a long-term experiment. A future study using Cre-controlled cell linear tracing techniques will be needed for long-term tracing experiments and to study the differentiation of the progenitors.

In summary, obesity and vascular injury result in elevated leptin release in both the blood and the vessel wall and this can serve as a chemokine for Sca-1⁺ progenitor cell migration. During this process, the binding of leptin to its receptor (OBR) leads to the activation of signal pathways, for example, STAT3-Rac1/Cdc42-ERK-FAK, which are abrogated by OBR deficiency in animal models. The migratory response of Sca-1+ progenitor cells to increased leptin levels may be largely responsible for enhanced neointimal formation in injured vessels. These novel findings enhance our understanding of the mechanisms behind obesity-related vascular diseases. The protective role of Lepr^{-/-} Sca-1⁺ progenitors also suggests a possibility to specifically target the receptor in the Sca-1+ progenitor cells to prevent vascular remodeling.

Sources of Funding

This study was supported by grants from the British Heart Foundation (RG/14/6/31144). Y. Xie was supported by the Oak Foundation and the China Scholarship Council.

None.

Disclosures

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Highlights

- Sca-1+ progenitor cells respond to leptin stimulation leading to cell migration, which was mediated by OBR (leptin receptor)-signal transducer and activator of transcription 3-RAC1 (ras-related C3 botulinum toxin substrate 1)/CDC42 (cell division control protein 42 homolog)-ERK1/2 (extracellular signal–regulated kinase 1/2)-pFAK (phosphorylated focal adhesion kinase) pathway.
- Lack of OBR strongly diminished cell migration, resulting in decreased neointimal formation in wire injured arteries.
- Transplantation of Lepr+/+ progenitor cells into leptin receptor deficient mice resulted in enhanced lesion formation.

JOURNAL OF THE AMERICAN HEART ASSOCIATION

Vessel-Injury Mouse Models Leptin Induces Sca-1⁺ Progenitor Cell Migration Enhancing Neointimal Lesions in

Bhaloo, Zhongyi Zhang, Yanhua Hu, Li Zhang and Qingbo Xu Yao Xie, Claire M.F. Potter, Alexandra Le Bras, Witold N. Nowak, Wenduo Gu, Shirin Issa

2017; *Arterioscler Thromb Vasc Biol.* 2017;37:2114-2127; originally published online September 21,

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SUPPLEMENTAL MATERIAL

Materials & Methods

Experimental Mice

All animal procedures were approved by the UK Home Office (PPL70/8944). C57BL/6J mice were purchased from Harlan, Blackthorn, Bicester, UK. RFP mice (Stock No: 005884) and leptin receptor deficient mice (Stock No: 00697) were purchased from the Jackson Laboratory, USA. Male wild-type (Lepr^{+/+}) and db/db (Lepr^{-/-}) mice aged 10-12 weeks were selected for surgery. Heterozygous (Lepr^{+/-}) mice were not used. All mice were kept on a chow diet in a 12-hour light and 12-hour dark environment at 25°C in the biological service unit (BSU) at the James Black Centre of King's College London.

Mouse Adventitial Progenitor Cell Culture

The procedure used for adventitial progenitor cell culture was similar to that described previously.¹ In brief, the aortic arch and root, as well as part of the heart from Lepr^{+/+} or Lepr^{+/-} mice, were harvested under sterile conditions. Adventitial tissues were carefully collected under a dissection microscope by removing the aortic media, intima and heart tissue. The adventitial tissues were then cut into pieces and seeded onto a gelatin-coated T25 flask. The flask was incubated at 37℃, 5% CO2 upside down for 2 to 3 hours. After the attachment of adventitial tissues onto the flask, complete stem cell medium (Dulbecco's Modified Eagle's Medium (DMEM ATCC, 30-2002)) with leukemia inhibitory factor (10 ng/mL, Merck Millipore, LIF1050), 2-mercaptoethanol (0.1 mM, GIBCO), 100U/mL penicillin (GIBCO), 100U/mL streptomycin (GIBCO) and 2 mM L-glutamine (GIBCO) was added for 5 days. When the cells reached 90% confluency, they were washed by PBS and trypsinized. The cells were passaged at a ratio of 1:4 every two or three days. Medium was changed every 2 days.

Mouse Vascular Smooth Muscle Cell Culture

The procedure used for vascular smooth muscle cells' culture was similar to that described previously.¹ In brief, the aortic arch and root, as well as part of the heart from Lepr^{+/+} or Lepr^{+/-} mice, were harvested under sterile conditions. Medial tissues were carefully collected under a dissection microscope by removing the aortic adventitia, intima and heart tissue. The media was then cut into pieces and seeded onto a 0.04% gelatin-coated T25 flask. The flask was incubated at 37℃, 5% CO2 upside down for 2 to 3 hours. After the attachment of medial tissues onto the flask, complete medium (Invitrogen 11960085, Dulbecco's Modified Eagle's Medium, high glucose) with 10% fetal bovine serum (GIBICO, 10270), 100U/mL penicillin (GIBCO, 15140122), 100U/mL streptomycin (GIBCO, 15140122) and 2 mM L-glutamine (GIBCO, 25030081) was added for 5 days. When the cells reached 90% confluency, they were washed by PBS and trypsinized. The cells were passaged at a ratio of 1:3 every two or three days. Medium was changed every 2 days.

Sca-1+ Progenitor Cell Isolation and Culture

When reached 90% confluency, the primary Sca-1⁺ adventitial progenitor cells were isolated using the Sca-1⁺ microbeads kit (Miltenyi Biotec, Bergisch Gladbach, 130-092-529) according to the manufacturer's instruction. Sca-1⁺ progenitor cells were selected using a magnetic cell separator. The Sca-1⁺ progenitor cells were cultured on 2% gelatin-coated flasks in complete stem cell medium (Dulbecco's Modified Eagle's Medium (DMEM ATCC, 30-2002)) with 10% Embryomax, leukemia inhibitory factor (10 ng/mL) and 2-mercaptoethanol (0.1 mM), penicillin (100 U/mL) and streptomycin (100 mg/mL) for both *in vitro* and *in vivo* studies.

Phenotyping of Cultured Sca-1+ Progenitor Cells And Smooth Muscle Cells

Cells were isolated from the adventitial of Lepr^{+/+} and Lepr^{-/-} (db/db) mice, and kept growing for few passages. The heterogeneous $Sca-1^+$ adventitial cells were purified by applying anti-Sca-1 microbeads. Flow cytometry was performed within 3 passages after sorting. Cultured lepr^{+/+} and Lepr^{-/-} Sca-1⁺ progenitors were harvested by using scraptase (GenDEPOT, CA110-010). Cells were then centrifuged and resuspended in cold PBS, followed by a 30-minute staining on ice with following antibodies: Anti-Sca-1-FITC (clone D7, abcam, ab25031), Anti-CD45-APC-Cy7 (clone 30-F11, biolegend, 103115), Anti-CD29-PE (clone HMβ1-1, BD Pharmingen, 562801), Anti-CD11b-BV785 (clone M1/70, biolegend, 101243), Anti-CD31-BV510 (clone MEC13.3, BD Horizon, 563089), Anti-CD140a-APC (clone APA5, eBioscience, 17-1401-81), CD105-PE (clone MJ7/18, BioLegend, 120407), CD117(ckit)-PE (clone 2B8, eBioscience, 12-1171-82), CD146-FITC (clone P1H12, eBioscience, 11-1469-41), Flk1-PE (clone AVAS 12α1, BD Pharmingen, 555308), anti-NG2 antibody (rabbit polyclonal, Abcam, ab129051), anti-Lepr (rabbit polyclonal, Sigma, HPA030899). Cells stained with secondary control only were used as negative control. Cells were analyzed with BD accuri C6 or BD LSR fortessa II (Becton Dickinson) flow cytometers.

Transwell Assay

Cell migration assessments were performed using transwell inserts with 8.0 μm micron pore membrane filters in a 24-well plate (#3422, Corning Life Science, USA) as established previously.² Sca-1⁺ progenitor cells were trypsinized and transferred into the upper chamber at 10^5 cells/mL in serum-free media with or without inhibitors (hoelzel-biotech, CYT354 / Santa Cruz, WP1066 / Merck Millipore, PD98059). Meanwhile, the lower chamber was loaded with 800 mL of serum-free media with treatment of leptin (Peprotech, 450-31). After 16-hour incubation, the upper side of the filters was carefully washed using a cotton bud to remove any non-migratory cells. Migratory cells on the underside of the filters were fixed with 4% PFA for 15 minutes, followed by a 15-minutes staining with 1% crystal violet at room temperature. Data is representative of mean cell numbers of migratory progenitor cells in 5 random fields at 20x magnification. For the experiments involving inhibitors or antagonists, Sca-1⁺ progenitor cells were cultured with the respective chemicals in the upper chamber only.

Scratch-wound Assay

 $1x10⁴$ cells per well of Sca-1⁺ progenitor cells were seeded in a 12-well plate in complete culture media. Once the cells reached 90% confluency, a scratch wound was made from top to bottom using a 1 mL pipette tip. The pipette was kept at a consistent angle and pressure during the scratch to ensure uniformity of width of the scratch. The wells were carefully washed twice using PBS to remove any cell debris caused by the "scratch" procedure. Treatment and/or inhibitors were added into wells with serum-free media. After 16-hour incubation, cells were fixed with 4% PFA and stained with 1% crystal violet for 15 minutes at room temperature. The migration of Sca-1⁺ progenitor cells into the "wound" area was evaluated using a phase contrast microscope. Data indicated are the mean area occupied by migratory progenitor cells in 5 random fields of view at 10x magnification.

RNA Extraction

Cells after overnight starvation in serum-free medium were treated with 100 ng/mL leptin for 5 min, 15 min, 30 min, 1 h and 4 h. Total RNA extraction from Sca-1⁺ progenitor cells was performed by applying RNeasy Mini kit (QIAGEN Inc., 74106) According to manufacturer's instruction, cells were washed twice by PBS, disrupted by proportional amount of RLT lysis buffer and then scraped off from the 6-well plate. The lysate was transferred into a mini QIAshredder spin column and centrifuged at full speed for 2 minutes. High-molecular weight DNA and other substances in lysates were hence removed. Same volume of 70% ethanol was added to the lysate and the mixture was transferred to RNeasy mini column for a 30-second centrifuge at full speed. The flow was discarded and 700μL of RW1 was added to RNeasy mini column. After 30-second centrifuge, the flow was discarded and 500μL of RPE was added to the column twice for washing away the ethanol. The flow was the discarded and the column was centrifuged within a new collection tube for 2 minutes at full speed to ensure no solution outside the column. At last, the RNeasy mini column was transferred into a new 1.5 mL RNA-free tube and 40μL DEPC water (Invitrogen) was added to the membrane of the column, followed by 1-minute centrifuge at full speed.. The RNA concentration was measured by a nanodrop spectrophotometer ND-1000 (Thermo Scientific, UK) at the absorbance at 280nm.

Reverse Transcription (RT)

Reverse transcription was achieved by QuantiTect Reverse Transcription Kit (Qiagen, 205311), according to the manufacturer's instructions. Briefly, 1 μg of RNA template, 2 μL of gDNA wipeout buffer and enough volume of RNase-free water were mixed in the tube with a total volume 14 μL. The tube was placed in a RT-PCR machine (TECHNE TC-412, Bibby Scientific, UK) at 42℃ for 2 minutes, after which 1 µL of RT enzyme, 1 µL of primer mix and 4 μL of RT buffer, making up to 6μL of reaction volume, was added to the tube mentioned above. Then the mixture was incubated at 42℃ for 15 minutes and subsequently 95℃ for 3 minutes. The cDNA obtained was diluted into 100 μL by using DEPC-treated water, acquiring a final concentration of 10 ng/μL.

Quantitative Real-Time Polymerase Chain Reaction (qPCR)

Quantitative real-time PCR was performed by applying SYBR green system (Qiagen, 204057). The target gene was amplified in a duplex in 20 μL PCR mixtures (10 μL Sybr Green, 2 μL cDNA template, 1.6 μL optimized primers and 6.4 μL DEPC water) which was loaded into a 96-well plate (Eppendorf White, Eppendorf, UK). The plate was centrifuged at 1000 rpm for 5 minutes before running the program in qPCR machine. Ct values were established using EPPENDORF Mastercycler ep realplex. GAPDH served as an endogenous control. Sequences of primer sets used in this experiment are listed in Table S1.

Primers were designed by using DNA Integrated Technologies (IDT). (http://eu.idtdna.com/scitools/Applications/RealTimePCR/)

Conventional Polymerase Chain Reaction (PCR)

Conventional polymerase chain reaction was performed in this project for the identification of Lepr^{-/-} mouse. 50 ng cDNA was amplified to the corresponding DNA by applying Tag DNA polumerase (Invitrogen, 1034253). The products were transferred into 2% agarose gel and observed by the Biospectrum AC Imaging system and Vision worksLS software. All primers for the identification of mice were provided by Jackson laboratory, USA. The sequences of the primers were listed in Table S2.

Western Blot Analysis

Femoral artery or Sca-1+ progenitor cells with or without treatment were lysed using RIPA buffer (Life Tech, 89901) with phosphatase inhibitor tablets (Roche, 04906845001,) and protease inhibitors (Roche, 11873580001). The lysate was sonicated using a Branson Sonifier 150 at level 1 for 8 seconds twice prior to 45-minutes incubation on ice. The lysate was then centrifuged at 15000 g for 10 minutes at 4 °C. The supernatant was collected and transferred to a new 1.5 mL tube. The concentration of proteins was measured by performing Biorad Protein Assay (BIO-RAD, 5000006). 20 μg of lysate mixed with SDS loading buffer was loaded into a NuPage 4-12% Bis Tris-gel immersed in NuPage MOPS SDS running buffer, followed by standard Western blot procedures. Primary antibodies against pJAK2 (Cell signaling, 3771S, 1:1000), JAK2 (Cell signaling, 3230, 1:1000), pSTAT3 (Cell Signaling, 9131S, 1:1000), STAT3 (Cell signaling, 4904, 1:1000), pERK 1/2 (Santa Cruz, sc-16982-R, 1:500), ERK 1/2 (Cell Signaling, 4695P, 1:1000), pOBr (Invitrogen, PA5-64638, 1:1000), OBR (Invitrogen, PA1-053,1:1000), pFAK (Abcam, ab39967, 1:600), FAK (Santa Cruz, sc557, 1:600) and GAPDH (Santa Cruz, sc25778, 1:1000) were used to detect the respective proteins. The ECL detection system (Invitrogen, RPN2106) was used to detect the membrane-bound primary antibodies.

MAPK Pathway Phosphorylation Array

The detection of phosphorylated MAPK-related protein was performed by using MAPK pathway phosphorylation array (AAHMAPK1-4, Raybiotech). The proteins of cells with or without treatment were collected by applying lysis buffer provided in the kit. The concentration of cell lysate was measured by using detection buffer provided in the kit. Array membrane was put into a well of the incubation tray and incubated by blocking buffer for 30 minutes at room temperature. Discard the blocking buffer and pipette 1mL of diluted sample into each well and incubate for 2 hours at room temperature. After two times washes, added 1 mL of the well-mixed detection antibody cocktail into each well for 2 hours at room temperature. After two times washes, 2mL of 1x HRP-Anti-Rabbit IgG was added into each well for 2 hours at room temperature. After two times washes, the membrane was well cleaned and incubated with 500 μL of the well-mixed detection buffer for 2 minutes at room temperature. The membrane was analyzed by applying an X-ray film or a chemiluminescence imaging system. The proteins detected in this array were listed below.

Histological Analysis

Femoral artery was fixed with 4% formalin overnight at 4°C prior to a machine-based dehydration. The dehydrated samples were embedded in paraffin and subsequently cut into 5 µm sections. H&E staining was performed using a standard protocol with Hematoxylin and Eosin for morphology analysis.

Cell Proliferation ELISA, BrdU

BrdU assay was performed using cell proliferation ELISA kit (Roche, 11669915001) according to a standard protocol provided by manufactory. Briefly, cells were cultured with leptin for different time points and then labelled with BrdU for 2 hours in a humidified atmosphere at 37°C. Incubation with Fixdenat solution and anti-BrdU POD were subsequently performed. After the three-time washing step, substrate reaction was applied and the absorbance of samples was measured at 450 nm.

Mouse Leptin ELISA

The concentration of serum leptin was measured by applying Mouse/Rat Leptin Quantikine ELISA Kit (R&D systems, # MOB00) according to the standard protocol provided by manufactory. Briefly, blood from wild-type and db/db mice was collected and allowed to clot for 2 hours at room temperature before centrifuging for 20 minutes at 2000g. Serum was collected and diluted 20 times for assays. ELISA microplate strips was firstly added 50 μL of Assay Diluent RD1W for each well. 50 μL of standard, control, or samples were then added to each well and incubated for 2 hours at room temperature. After five times wash, 100 μL of Mouse/Rat Leptin Conjugate was added to each well for 2-hour incubation at room temperature. Repeat the washing procedure and add 100 μL of Substrate Solution to each well for 30 minutes at room temperature away from the light. 100 μL of Stop Solution was finally added to each well and the absorbance was determined by using a microplate reader set to 450 nm within 30 minutes. The absorbance at 540 nm or 570 nm was used as wavelength correction.

G-LISA RhoA/Rac1/Cdc42 Activation Assay

The activation of GTPase family was determined by performing G-LISA RhoA/Rac1/Cdc42 Activation Assay (Cytoskeleton, Inc. #BK135) according to the standard protocol provided by manufactory. Briefly, cells were seeded and grown to 70% confluency, followed by the stimulation of leptin from 2 to 30 minutes. Cells were collected and lysed on ice and the concentration of cell lysate was measured by using the detection solution provided in the kit. After adding buffer blank and buffer positive control, 50µl of equalized lysate were added to duplicate wells in an incubation plate. The plate was immediately placed on an orbital shaker at 400rpm for 15 (Rac1 and Cdc42) or 30 (RhoA) minutes in the cold room. After two times wash, 200 µl of room temperature Antigen Presenting Buffer was added to each well at room temperature for 2 minutes. After three times wash, 50 µl of diluted anti-RhoA/Rac1/Cdc42 primary antibody was added to each well on an orbital shaker at 400rpm for 30 (Rac1 and Cdc42) or 45 (RhoA) minutes at room temperature, followed by the incubation of 50 µl of diluted secondary antibody to each well for 30 (Rac1 and Cdc42) or 45 (RhoA) minutes at room temperature. After three times wash, 50 µl of the mixed HRP detection reagent was added into each well at 37°C for 10-15 min, the reaction of which was stopped by administration of 50 µl of HRP Stop Buffer. The absorbance was measured at 490 nm by using a microplate spectrophotometer. Buffer blank was designated as assay blank.

RFP Labelling Cells

Lentiviral particles were generated by transfecting HEK293T cells with LV H2b RFP plasmid (Addgene, 26001) and used to label the nucleus of adventitial cells. Sca-1⁺ progenitor cells were incubated with RFP lentivirus for 10 minutes, and the waste was carefully disposed. Expression of RFP was checked using a fluorescent microscope two days after the treatment with lentivirus.

Mouse Femoral Artery Denudation Injury and Cell Delivery

The procedure used for the mouse model is similar to that established previously.³ Wild-type or leptin receptor deficient mice (db/db) were anesthetized with ketamine and medetomidine hydrochloride. A groin incision was made under a surgical microscope. An arteriotomy was made in the epigastric branch of femoral arteries. A 0.014' guide wire (Hi-Torque, Cross-it 200XT) was inserted into the femoral artery to the level above the bifurcation of the abdominal aorta. The guide wire was gently pulled back and forth for three times, leading to the endovascular injury of the arteries. After removal of the guide wire, the femoral artery was ligated. Sham-injury arteries without passage of the guide wire were used as a control. After the injury, 1×10⁶ (RFP) progenitor cells with or without treatment in 30 µL Matrigel[®] Basement Membrane Matrix per artery were delivered to the adventitial side of injured artery. The arteries were harvested on days 1, 7, 14 or 28 post-surgery for *En Face*, frozen section or paraffin sections.

Immunofluorescence Staining and *En Face* **preparation**

Frozen sections were fixed with 4ºC acetone and permeabilized with 0.1% Trixton X-100 in PBS for 30 minutes at room temperature. For *En Face* staining, tissues were fixed with 4% paraformaldehyde for 15 minutes and then permeabilized with 0.5% Trixton X-100 in PBS for 30 minutes. For cell staining in chamber slides (BD Biosciences), the cells were fixed with 4% paraformaldehyde for 15 minutes and then permeabilized with 0.1% Trixton X-100 in PBS for 30 minutes. Tissues or cells were blocked in 5% swine serum for 45 minutes at room temperature prior to their incubation with primary antibodies for either 1 hour at 37 ºC or overnight at 4 ºC. After three washes with PBS, secondary antibodies were administered for 45 minutes at 37 ºC, followed by DAPI (1:5000 in PBS) staining for 5 minutes at room temperature. Frozen section or cells in chamber slides were mounted with fluorescence mounting medium (Dako, S3023). For *En Face* preparation, after the staining with DAPI the vessel was cut open along the long axis with the lumen facing up.⁴ The opened vessels were then transferred onto a clean slide and mounted with fluorescence mounting media (Vectashield). A confocal microscope (Leica SP5) and AxioVision Digital Imaging System (Carl Zeiss Ltd) were used for image acquisition. Primary antibodies applied in the present study were Sca-1 (Abcam, ab25031, 1:200), SMA (Sigma, A5228, 1:200), CD31 (Abcam, ab30349, 1:200), leptin (Invitrogen, Pa1-051, 1:200), F4/80 (Abcam, ab6640, 1:200), CD68

(Santa Cruz, sc-9139, 1:200), VE-cadherin (Santa Cruz, sc-6458, 1:200), pFAK (Abcam, ab39967, 1:200), FAK (Santa Cruz, sc557, 1:200), CD45 (Biolegend, 103115, 1:200), OBR (R&D systems, AF497, 1:200), RFP (Abcam, ab62341, 1:200). Slides stained with secondary antibodies only were used as negative control. Secondary antibodies included Thermofisher AlexaFluor Donkey anti-mouse 488 (A-21202) and 594 (A-21203), AlexaFluor Donkey anti-rabbit 488 (A-21206), 594 (A-21206) and 633(A-21070), 647 (A-31573), AlexaFluor Donkey anti-goat 488 (A-11055), 594 (A-11058) and 633 (A-21082), AlexaFluor Donkey anti-rat 488 (A-21208), 594 (A-21209) and 647 (A-21094).

Immunohistochemical Staining

The procedure used in present study is similar to that established previously.³ Immunohistochemistry was achieved by using vectastain ABC HRP kit (Vector laboratories, PK-6100). Briefly, praffin sections were incubated with SMA (Sigma, A5228, 1:200) primary antibody for 1 hour at room temperature after fixation and blocking. Diluted biotinylated secondary antibody was then applied for 30 minutes at room temperature. After washing, sections were incubated in peroxidase substrate solution until desired stain intensity showed up. The sections were then rinsed, counterstained, cleared and mounted.

Van Gieson Staining

Slides were underwent deparaffin and rehydration, following the immersion in 32 mmol/L of potassium permanganate for 10 minutes. After rinsing in dH₂O, the slides were immersed in 70 mmol/L of oxalicacid dehydrate for 5 minutes, washing in several changes of $dH₂O$. The slides were again rinsed in 70% ethanol, followed by a 3-hour incubation in Miller's stain. After rinsing in 70% ethanol and dH_2O , the slides were incubated in Van Gieson solution for 5 minutes. After washing in dH_2O , the slides were dried and mounted with mounting medium.

Statistical Analysis

Data represented as the mean and standard error of the mean (S.E.M.) of at least three individual experiments. Data were analyzed using Graphpad Prism 6. For data involved with two groups, unpaired and two-tailed Student's t-test was applied. For data involved with more than two groups, ANOVA test was applied followed by Dunnett's multiple comparison test. The mean of each column with different treatments was compared to the mean of the column named either "Ctr", "Control" or "0". Significance was considered as p-value <0.05.

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Supplemental Data Figures

Figure I. The expression of progenitor and hematopoietic markers in leptin-induced migrated cells. Migrated Lepr^{+/+} Sca-1⁺ progenitor cells after transwell assay were analyzed by immunofluorescence for OBR, Sca-1 (**A**), CD29 (**B**), CD34 (**C**) and CD45 (**D**) (scale bars, 3 µm, n=3).

Figure II. Sca-1⁺ OBR⁺ cells reside mainly in the adventitia of aorta and femoral artery. **A**, Cross sections of aorta from wild-type mice were analyzed by immunofluorescence for Sca-1 and OBR (scale bars, 100 and 50 µm, n=4). Red arrow indicated Sca-1+ OBR+ adventitial cells. White arrow indicated Sca-1+ OBR+ medial cells. Purple arrow indicated Sca-1+ OBR+ endothelial cells. **B**, Cross sections of femoral artery from wild-type mice were analyzed by immunofluorescence for Sca-1 and OBR (scale bars, 50 and 25 µm, n=4).

Figure III. Phenotyping of Lepr^{+/+} and Lepr^{-/-} of Sca-1⁺ adventitia progenitor cells. **A**, Phenotyping of Lepr^{+/+} Sca-1⁺ adventitial progenitor cells with primary antibodies of CD45, CD11b, CD31, CD29, CD140a and Sca-1. **B**, Phenotyping of Lepr¹ Sca-1⁺ adventitial progenitor cells with primary antibodies of CD45, CD11b, CD31, CD29, CD140a and Sca-1.

Figure IV. Phenotyping of Lepr^{+/+} and Lepr^{-/-} of Sca-1⁺ adventitial progenitor cells. **A**, Phenotyping of Lepr^{+/+} Sca-1⁺ adventitial progenitor cells with primary antibodies of CD29, CD34, CD105, c-kit, CD146 and Kdr. **B**, Phenotyping of Lepr-/- Sca-1+ adventitial progenitor cells with primary antibodies of CD29, CD34, CD105, C-kit, CD146 and KDR.

Figure V. The identification of Lepr¹ mice was achieved by applying conventional PCR and immunostaining. **A**, Cross sections of uninjured femoral artery from wildtype were analyzed by immunofluorescence for αSMA, Sca-1 and Lepr (scale bars, 50 µm, n=4). **B**, Cross sections of uninjured femoral artery from db/db were analyzed by immunofluorescence for αSMA, Sca-1 and Lepr (scale bars, 50 µm, n=4). **C**, The semi-quantification of the numbers of Sca-1+ adventitial cells was evaluated in the femoral artery of db/db and WT mice (n=5). **D**, Conventional PCR was performed for the identification of every Lepr^{-/-} mice (n=29).

Figure VI. Leptin activated pMEK1/2 but not pMKK3/6 signaling pathways on Sca-1+ adventitial progenitor cells. **A,** Western blotting was performed on Sca-1+ progenitor cells in response to 100 ng/ml of leptin for the detection of pMKK3/6 (n=3), pMEK1/2 (n=4). **B** and **C**, Quantification of the activation of pMKK3/6 (B) and pMEK1/2 (C).

Figure VII. Leptin activated the GTPase Cdc42 and Rac1 on Sca-1⁺ adventitial progenitor cells. **A** and **B**, Quantification of activated GTPase Cdc42 (**A**) and Rac1 (**B**) was evaluated in response to 100 ng/mL of leptin by performing G-LISA G-protein activation assays (n=3). **C**, The expression of MAPK-related protein was detected by applying MAPK protein microarrays over time in response to 100 ng/mL of leptin (n=3). Untreated cells were served as control. All graphs are shown as mean

Figure VIII. Inhibition of OBR and ERK pathway significantly reduced the migration of Sca-1+ adventitial progenitor cells in response to 100 ng/mL of leptin. **A** and **B**, Migration of Sca-1+ progenitor cells in response to 100 ng/mL of leptin with or without ERK inhibitor was evaluated by applying 1% crystal violet staining after 16-hour incubation (scale bars, 50 µm, n=5). **C** and **D**, Migration of Sca-1+ progenitor cells from db/db mice in response to an increasing gradient concentration of leptin was evaluated by 1% crystal violet staining after 16-hour incubation (scale bars, 50 µm, n=5). **E** and **G**, Chemotaxis of Sca-1+ progenitor cells in response to 100 ng/mL of leptin and ERK inhibitor was documented by using an 8.0 µm transwell system (scale bars, 50 µm, n=5). **F** and **H**, Chemotaxis of Sca-1+ progenitor cells from db/db mice in response to an increasing gradient of leptin in an 8.0 µm transwell system was identified by applying 1% crystal violet staining after 16-hour incubation (scale bars, 50 µm, n=5). Serum-free cultured medium and dimethyl sulfoxide (DMSO) without leptin treatment was used as controls for the migratory assays above. Migration index for transwell and wound healing assays was defined as the mean ratio of treatment to control of cell numbers counted per 5 random fields at 20X magnification. All graphs are shown as mean \pm SEM. *p<0.05, **p<0.01, ***p<0.001.

Figure IX. OBR and STAT3 were the upstream of ERK1/2. **A**, Western blotting was performed on Sca-1⁺ progenitor cells in response to 100 ng/ml of leptin with ERK inhibitor for the detection of pSTAT3 (n=8) and pERK1/2 (n=10). **B** and **C**, Quantification of the activation of pSTAT3 and pERK1/2. **D**, Western blotting was performed on Sca-1⁺ progenitor cells in response to 100 ng/ml of leptin with STAT3 inhibitor for the detection of pSTAT3 (n=4) and pERK1/2 (n=4). **E** and **F**, Quantification of the activation of pSTAT3 and pERK1/2. **G**, Western blotting was performed on Lepr^{./-}Sca-1⁺ progenitor cells in response to 100 ng/ml of leptin for the detection of pSTAT3 (n=4) and pERK1/2 (n=4). **H** and **I**, Quantification of the activation of pSTAT3 and pERK1/2.

A

Figure X. Identification of RFP and non-RFP Sca-1+ adventitia progenitor cells after lentivirus transfection. **A**, Confirmation of the existence of RFP in Sca-1+ adventitia progenitor cells with primary antibodies of RFP by applying FACS assay.

Figure XI. Guide-wire injury induced significant neointimal formation 2 weeks and 4 weeks after surgery. **A**, H&E staining of the femoral artery after guide-wire injury was performed (scale bar, 50 and 10 µm, n=10). **B** and **C**, The composition of the neointima was evaluated by immunohistochemistry (B) and immunofluorescence (C) for smooth muscle actin on frozen sections of femoral artery at 2 or 4 weeks after surgery (scale bar, 50 and 10 µm, n=10). **D**, Quantification of lumen area of injured arteries (n=10). **E**, Quantification of media area of injured arteries (n=10). **F**, Quantification of neointimal area of injured arteries (n=10). **G**, Quantification of ratio of neointima to media of injured arteries (n=10). **H**, Quantification of ratio of media to adventitia of controls and injured arteries (n=10). Graphs are shown as mean \pm SEM. *P<0.05, **P<0.01, ***P<0.001. Dashed box represented the magnified field. Arrows indicate SMCs or ECs. EC indicates endothelium; SMC, smooth muscle cells; Neo, neointima; M, media; SMA, smooth muscle actin; 2W, 2 weeks; 4W, 4 weeks.

Figure XII. The expression of macrophage marker F4/80 was upregulated after surgery around the injured artery. Expression of macrophage (Alexa 488; green) in injured artery was analyzed on day 1(**A,** n=10), 3(**B,** n=7), 5(**C**, n=5) and 7(**D**, n=10) post-operation by performing immunofluorescence. (scale bars, 50 and 10 µm). Dashed box represented the magnified field.

Figure XIII. Transplantation of exogenous Sca-1+ progenitor cells did not affect the extent of fibrosis during the neointimal formation. **A**, Cross sections of injured femoral artery at 1 week post-surgery were analyzed by Elastica-van Gieson staining (scale bars, 50 and 10um, n=3). **B**, Lepr^{+/+} Sca-1⁺ progenitor cells were seeded on the adventitial side of injured artery in wild-type mice. Cross sections of femoral artery at 1 week post-surgery were analyzed by Elastica-van Gieson staining (scale bars, 50 and 10 µm, n=3). **C**, Cross sections of injured femoral artery at 2 week postsurgery were analyzed by Elastica-van Gieson staining (scale bars, 50 and 10µm, n=3). **D**, Lepr+/+ Sca-1+ progenitor cells were seeded on the adventitial side of injured artery in wild-type mice. Cross sections of femoral artery at 2 week post-surgery were analyzed by Elastica-van Gieson staining (scale bars, 50 and 10µm, n=3). Images shown are representative of at least 3 independent experiments.

Figure XIV. Both adventitial Sca-1⁺ cells and hematopoietic cells contributed to neointimal formation. **A**, Lepr^{+/+} Sca-1⁺ progenitor cells were seeded on the adventitial side of injured artery in db/db mice. Cross sections of femoral artery from db/db mice at 2 weeks after the surgery were analyzed by immunofluorescence for CD45 and OBR (scale bars, 50 and 10µm, n=4). Dashed box represented the magnified field. Images shown are representative of at least 3 independent experiments.

Figure XV. The migration of Sca-1⁺ adventitial progenitor cells could be inhibited or induced by different adipokines. **A**, Chemotaxis of Sca-1+ progenitor cells in response to 10 ng/mL of adiponectin (n=3), 10 ng/mL of resistin (n=3) and 100 ng/mL of leptin (n=6) was evaluated by applying 1% crystal violet staining after 16-hour incubation (scale bars, 50µm). **B**, Migration of Sca-1+ progenitor cells in response to 10 ng/mL of adiponectin (n=3), 10 ng/mL of resistin (n=3) and 100 ng/mL of leptin (n=6) was documented by using an 8.0 µm transwell system. Serumfree cultured medium and dimethyl sulfoxide (DMSO) without treatment was used as controls for the migration assays above.

Figure XVI. Leptin could enhance the expression of CXCR5 except for OBR. **A**, Expression of CXCR5 for Sca-1+ progenitor cells was evaluated by qPCR 24 hours after the treatment of leptin (n=5). All graphs are shown as mean \pm SEM. *p<0.05, **p<0.01, ***p<0.001.

Figure XVII. Leptin might induce the differentiation of Sca-1⁺ progenitor cells towards SMC. **A**, Expression of Calponin (n=5) for Sca-1+ progenitor cells was evaluated by qPCR in response to 100 ng/mL of leptin. All graphs are shown as mean \pm SEM. *p<0.05, **p<0.01, ***p<0.001.

vessel injury was analyzed in db/db and wild-type. **A**, Cross sections of injured femoral artery from wild-type mice were analyzed by immunofluorescence for Sca-1 (scale bars, 50 µm, n=4). **B**, Cross sections of injured femoral artery from db/db mice were analyzed by immunofluorescence for Sca-1 (scale bars, 50 µm, n=4). **C**, The semi-quantification of number of Sca-1+ adventitial cells was evaluated in wild-type mice before and at two weeks post-surgery (n=4). **D**, The semi-quantification of number of Sca-1⁺ adventitial cells was evaluated in db/db mice before and at two weeks post-surgery $(n=4)$.

Figure XIX. APCs and SMCs from db/db mice showed a difficulty to expand *in vitro.* **A**, Morphology of Sca-1+ progenitor cells from db/db mice. **B**, Morphology of SMCs from db/db mice. **C** and **D**, Proliferation between Lepr-/- or lepr+/+ Sca-1+ progenitors (**C,** n=4) and SMC (**D,** n=4) was examined by BrdU assay after 16-hour incubation with 100 ng/mL of leptin. Serum-free cultured medium without leptin treatment was used as a control for the migration assays above. Figures shown above are representative of at least 3 separate experiments (scale bars, 50 µm). APC indicates Sca-1+ adventitial progenitor cells; SMC indicates smooth muscle cells. All graphs are shown as mean \pm SEM. *p<0.05, **p<0.01, ***p<0.001.

Migration towards Leptin

Figure XX. Schematic illustration of the roles of exogenous leptin in enhancing Sca-1+ adventitial progenitor cells chemotaxis. Elevated leptin in circulation or femoral artery after guide-wire injury binds to its receptors OBR on the Sca-1⁺ adventitial progenitor cells. The GTPases Rac1 and Cdc42 are activated leading to the phosphorylation of pERK. Leptin also induces the activation of STAT3 pathway and the expression of cytoskeleton related proteins paxillin, vinculin and phosphorylated FAK, which may eventually contribute to cell migration.

Supplemental Data Tables

Table I. Primers used

Table II. Primers used

