

LETTER TO THE EDITOR

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# Loss of Angiopoietin-like 7 diminishes the regeneration capacity of hematopoietic stem and progenitor cells

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## Abstract

Successful expansion of hematopoietic stem cells (HSCs) would benefit the use of HSC transplants in the clinic. Angiopoietin-like 7 promotes the expansion of hematopoietic stem and progenitor cells (HSPC) *in vitro* and *ex vivo*. However, the impact of loss of *Angptl7* on HSPCs *in vivo* has not been characterized. Here, we generated *Angptl7*-deficient mice by TALEN-mediated gene targeting and found that HSC compartments in *Angptl7*-null mice were compromised. In addition, wild type (WT) HSPCs failed to repopulate in the BM of *Angptl7*-null mice after serial transplantations while the engraftment of *Angptl7*-deficient HSPCs in WT mice was not impaired. These results suggest that *Angptl7* is required for HSPCs repopulation in a non-cell autonomous manner.

**Keywords:** *Angptl7*, Knockout-mice, Hematopoietic stem cell, Repopulation, Homing

## Findings

The autologous-allogeneic hematopoietic cell transplantation has been developed for decades [1,2], and numerous attempts have been made to expand the HSCs population *in vitro* and *in vivo* [3-5]. Angiopoietin-like proteins belong to a 7-member family of secreted glycoproteins that share sequence homology with angiopoietins, which are important modulators of angiogenesis [6]. Several *Angptl* family proteins promote the expansion of murine and human HSPCs *in vitro* and *ex vivo* [7-9]. *Angptl7* in the ECM of the trabecular meshwork plays an important role in the deposition and organization of the matrix of the outflow tissue [10]. Recently, we found that *ANGPTL7* stimulated the proliferation of human HSPCs *ex vivo* (Yiren Xiao, unpublished data). In addition, we uncovered that *Angptl7*,

which was secreted by murine bone marrow SSEA4+ mesenchymal cells (Additional file 1: Figure S1), stimulated expansion of murine HSCs *ex vivo* (Additional file 2: Figure S2). However, whether *Angptl7* is redundant and dispensable or not for repopulation of HSPCs *in vivo* remains unknown. In this study, we generated *Angptl7* knockout mice and revealed that *Angptl7* is essential for HSPC repopulation in a non-cell autonomous way.

To investigate whether the loss of *Angptl7* affected HSPCs *in vivo*, we generated *Angptl7*-null mice by TALEN-mediated gene targeting (Additional file 1: Figure S1a-1c). The procedures of gene targeting were described in (Additional file 3: Supplementary methods) and (Additional file 4: Table S1). Loss of *Angptl7* in *Angptl7*-null mice was confirmed in the bone marrow (BM) by RT-PCR and Western blotting (Additional file 5: Figure S3d-3e). The *Angptl7*-null mice did not display an overt phenotype. There were no significant differences in either birth weights or adult weights among *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup>, and *Angptl7*<sup>+/+</sup> mice (Additional file 6: Table S2). In addition, we found no significant differences in staining profiles T cells, B cells, myeloid cells, and erythroid cells among the *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> mice (Figure 1a).

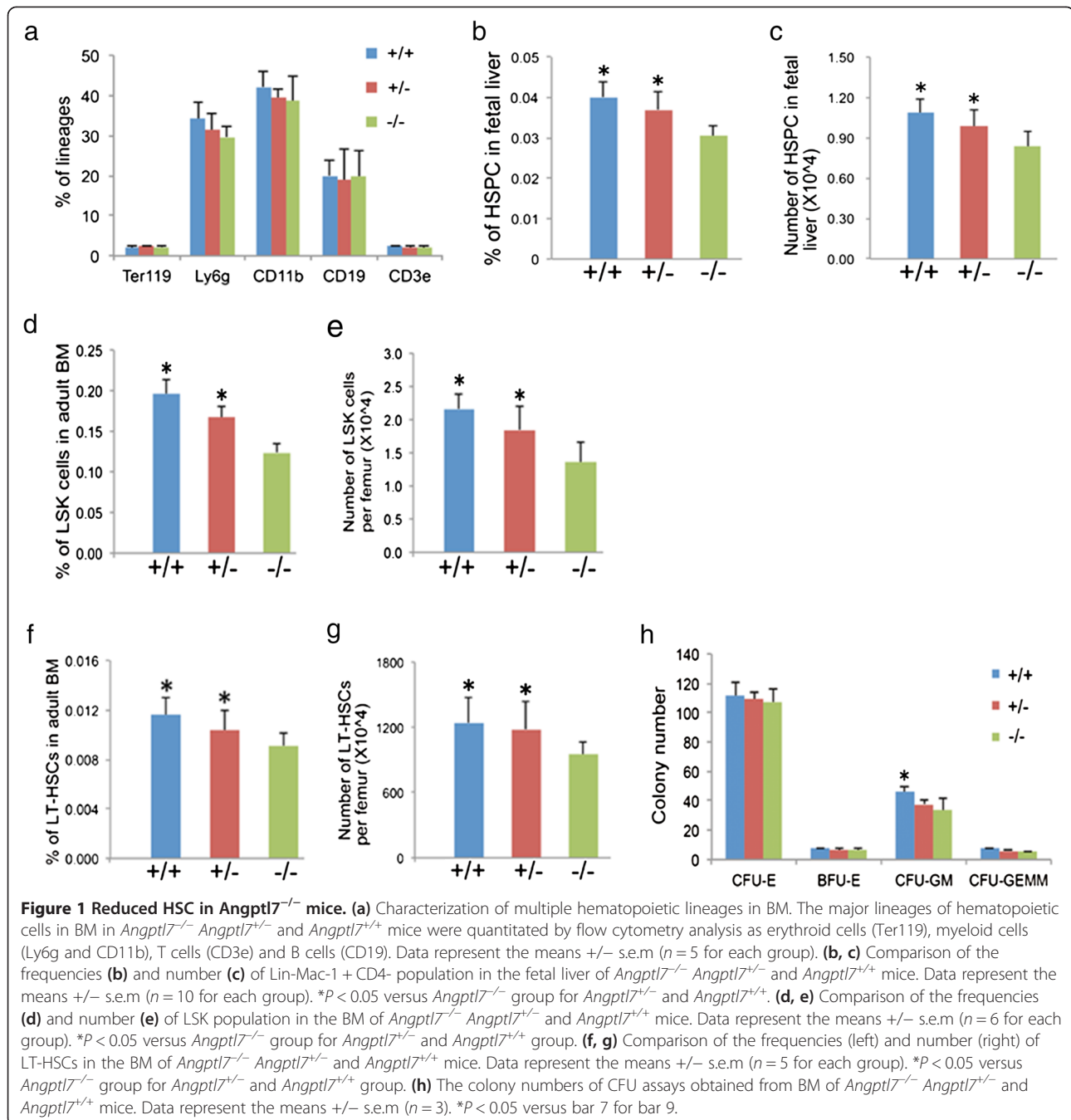
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As HSCs undergo dramatic expansion in fetal liver during embryo development [11-13], we examined the fetal liver HSC compartment and found that the percentages and numbers of Lin-Mac-1 + CD4<sup>-</sup> population, which was highly enriched of HSCs in fetal liver [14,15], were significantly lower in the fetal livers of *Angptl7*<sup>-/-</sup> mice than that in *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> mice (Figure 1b-c). In addition, we found the percentages and numbers of Lin-Sca-1 + c-Kit<sup>+</sup> (LSK) population in the BM of adult *Angptl7*<sup>-/-</sup> mice were significantly

lower than that in *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> mice (Figure 1d-e). Further analysis showed that *Angptl7*<sup>-/-</sup> mice had fewer long-term HSCs (Lin<sup>-</sup>Sca-1<sup>+</sup>Kit<sup>+</sup>Flk2<sup>-</sup>CD34<sup>-</sup>) than *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> mice (Figure 1f-g). Cloning forming assay results showed that BM cells from *Angptl7*<sup>-/-</sup> mice had fewer granulocyte/monocyte progenitors (CFU-GM), but similar numbers of erythroid precursors (CFU-E/BFU-E) and hematopoietic progenitors (CFU-GEMM) than WT BM (Figure 1h). Therefore, *Angptl7*-deficient BM had normal levels of terminally

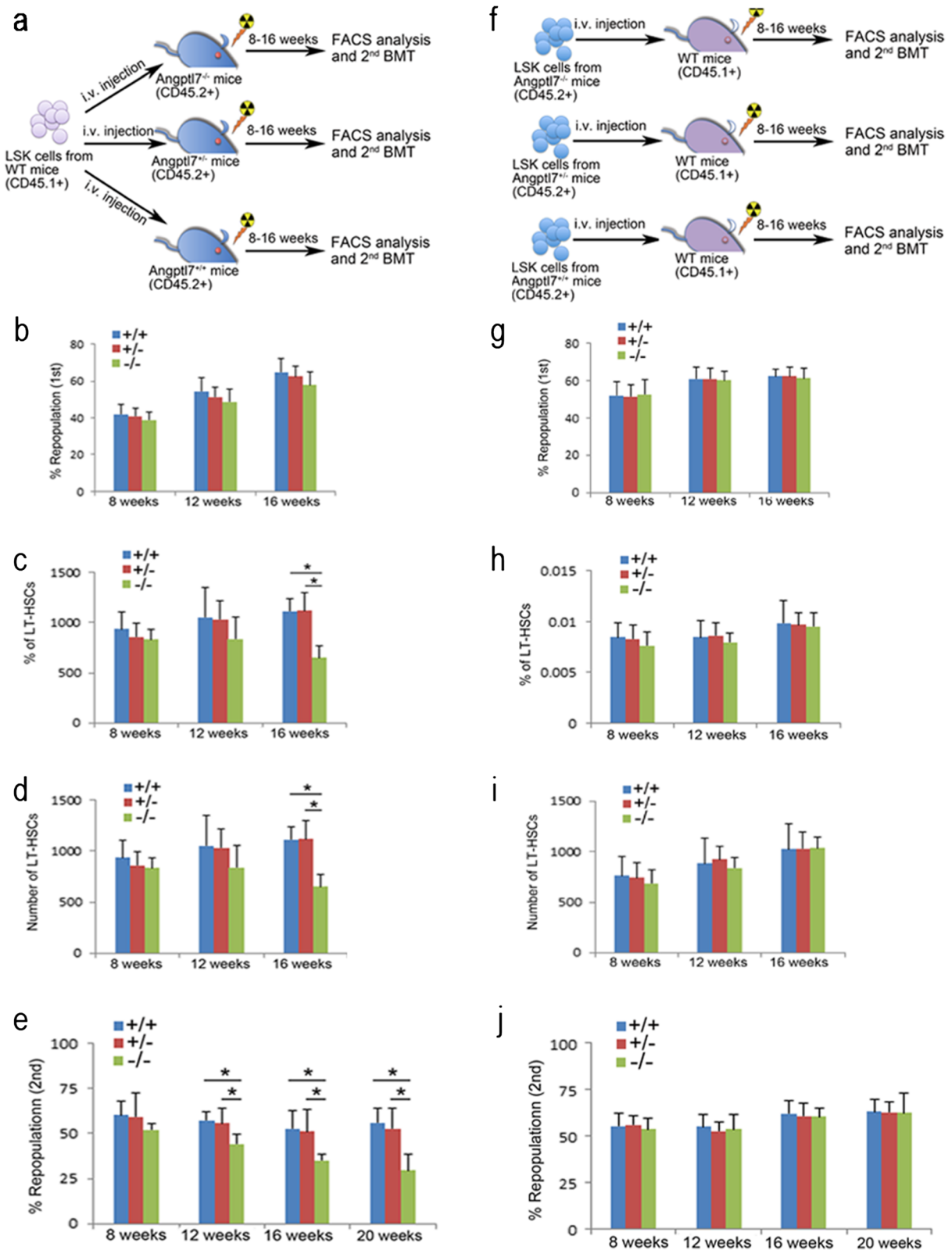


Figure 2 (See legend on next page.)

(See figure on previous page.)

**Figure 2 Comparison of reconstitution capacity of HSC in *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> mice.** (a) Experimental design for assessing the repopulation efficiency in *Angptl7*-null BM environment. (b) Comparison of the repopulation efficiency in BM of *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> recipients. Data represent the means  $\pm$  s.e.m. ( $n = 8$  for each group). (c, d) Comparison of the frequencies (c) and number (d) of LT-HSCs in BM of *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> recipients. Data represent the means  $\pm$  s.e.m. ( $n = 8$  for each group). \* $P \leq 0.05$  for bar 9 versus bar 7 and bar 8. (e) Comparison of the second transplantation repopulation efficiency in BM of *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> recipients as referred in (a). Data represent the means  $\pm$  s.e.m ( $n = 7$  for each group). \* $P \leq 0.05$  for bar 6 versus bar 4 and bar 5, for bar 9 versus bar 7 and bar 8, for bar 12 versus bar 10 and bar 11. (f) Experimental design effect of endogenous deletion of *Angptl7* in HSCs. *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> or *Angptl7*<sup>+/+</sup> donor BM CD45.2 cells ( $5 \times 10^5$  cells) were transplanted into lethally irradiated WT CD45.1 recipient mice. (g) Comparison of the repopulation efficiency of donor BM *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> cells. Data represent the means  $\pm$  s.e.m. ( $n = 8$  for each group). (h, i) Comparison of the frequencies (h) and number (i) of donor *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> LT-HSCs in BM of WT recipients. Data represent the means  $\pm$  s.e.m. ( $n = 8$  for each group). (j) Comparison of the repopulation efficiency of *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> donor BM cells in second recipients mice as described in (f). Data represent the means  $\pm$  s.e.m ( $n = 8$  for each group).

differentiated hematopoietic cells, but decreased myeloid progenitors. To investigate whether other angiopoietin-like proteins compensate the loss of *Angptl7* in BM, we compared the expression levels of other angiopoietin-like proteins in *Angptl7*-deficient BM stromal cells to that in WT BM stromal cells and found that their expression levels did not significantly change due to loss of *Angptl7* (Additional file 7: Figure S4).

Since *Angptl7* is secreted by stromal cells and binds to HSPCs [9], we speculated that *Angptl7* may play a non-cell autonomous role in reconstitution of the hematopoietic system. To evaluate the hypothesis, we compared the extent of repopulation of HSCs in *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and *Angptl7*<sup>+/+</sup> recipient mice. After WT donor BM cells were injected into lethally irradiated *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> or *Angptl7*<sup>+/+</sup> recipients without competitors (Figure 2a), we found that the reconstitution efficiencies of HSCs were similar among *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and WT recipient mice (Figure 2b), but the frequencies and numbers of LT-HSCs in *Angptl7*<sup>-/-</sup> recipients were significantly lower than that in *Angptl7*<sup>+/-</sup> and WT recipients, suggesting that *Angptl7* supports the maintenance of the HSC pool in BM (Figure 2c-d). Continuously the BM cells from the primarily repopulated *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and WT recipient mice were collected for secondary transplantation. We found that the repopulating activity of cells originating from the primary *Angptl7*<sup>-/-</sup> recipients was significantly decreased compared with those from the primary *Angptl7*<sup>+/-</sup> and WT recipients at different time points (Figure 2e). To test whether *Angptl7* had a cell-intrinsic effect on HSCs, we transplanted *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> or WT BM cells into lethally irradiated WT recipients and measured the frequencies of donor cells in recipient mice (Figure 2f). We found the repopulation efficiencies, percentages, and numbers of donor *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and WT LT-HSCs were similar in WT recipients (Figure 2g-i). We further transplanted the BM cells from the primary recipient mice to lethally irradiated WT mice. The repopulation efficiencies of secondary transplantation were similar among *Angptl7*<sup>-/-</sup>, *Angptl7*<sup>+/-</sup> and

WT donors at indicated time points (Figure 2j). Overall, these results suggest that *Angptl7* promoted HSC repopulation in a non-cell autonomous manner *in vivo*.

In this study, we demonstrated that deficiency of *Angptl7* in the BM niche as a lack of exogenous *Angptl7* stimulation impaired the reconstitution of HSCs in lethally irradiated mice, whereas endogenous deletion of *Angptl7* in HSCs did not affect repopulation of HSCs in lethally irradiated mice. Thus, *Angptl7* was indispensable for BM microenvironment to support HSC repopulation.

## Additional files

**Additional file 1: Figure S1.** Generation of *Angptl7* knockout mice. (A) *Angptl7* is highly expressed in BM CD45-SSEA4+ cells. BM cells were collected by flow cytometry and *Angptl7* expression was measured by real-time RT-PCR. All the cells sorted for analysis were gated from the CD45- fraction. The results were normalized to  $\beta$ -actin mRNA levels and represent the means  $\pm$  s.e.m. \* $P < 0.05$  versus bar 13 for bar 14. (B) Fluorescent microscopy imaging analyzes *Angptl7* (green) and SSEA4 (red) cells in mouse BM cells. Nuclei were counterstained with DAPI (blue).

**Additional file 2: Figure S2.** Treatment with *Angptl7* induces the expansion of mouse HSPC. Left, Representative flow cytometric analysis of peripheral blood (PB) donor-derived CD45.1+ engraftment at 12 weeks after transplantation in mice transplanted with the progeny of  $1 \times 10^5$  bone marrow lin- cells after 7 days culture with TSF or TSF plus 500 ng ml<sup>-1</sup> *Angptl7*. Right, The mean levels of donor CD45.1+ cells in the PB of CD45.2 + recipient mice at 12 weeks after transplantation of CD45.1+ bone marrow Lin- cells ( $1 \times 10^5$ ) or their progeny after 7 d of culture with TSF or TSF plus 500 ng ml<sup>-1</sup> *Angptl7*. Data represent the means  $\pm$  s.e.m ( $n = 6$  mice per group). \* $P < 0.05$  versus bar 2 and bar 3 for bar 1, bar 2 for bar 3.

**Additional file 3: Supplemental methods.**

**Additional file 4: Table S1.** Primer list used in generation of *Angptl7*-deficient mice.

**Additional file 5: Figure S3.** (a) Top, the DNA sequences of *Angptl7* locus targeted by *Angptl7*-TALENs are shown. The DNA-binding sites are in red, and the sequence between the DNA-binding sites is spacer region blue. There is a restriction site for the endonuclease DrrI between the two binding sites (blue). Bottom, schematic view of the design of TALENs targeting the *Angptl7* locus. The *Angptl7* donor vector contains two homologous arms on both sides of exon 1 of *Angptl7*, a splice acceptor, encoding cDNA of Venus, and puromycin (Puro) that was driven by PGK promoter. *Angptl7*-TALEN recognition sites (purple box), SD: splice donor, SA: splice acceptor. (b) Electropherograms around the TALEN spacer in the *Angptl7* locus. The red boxes highlight the mice identified with *Angptl7* mutations. (c) DNA sequences of the *Angptl7* locus from live F0 mice identified in (b). '-' represent deleted nucleotides. (d) The lack of *Angptl7* in the BMs of *Angptl7* knockout mice was confirmed by RT-PCR. (e)

The lack of Angptl7 in the BMs of Angptl7 knockout mice was confirmed by western blots.

**Additional file 6: Table S2.** The organ weight of *Angptl7*-deficient mice.

**Additional file 7: Figure S4.** Relative expression levels of angiotensin-like proteins in WT (+/+) and *Angptl7*-/- bone marrow stromal cells. The results were normalized to  $\beta$ -actin mRNA levels and represent the means  $\pm$  s.e.m. \* $P < 0.05$  versus bar 11 for bar 12.

#### Competing interests

The authors declare that they have no competing interests.

#### Authors' contributions

Contribution: YX, XW, DP, and PL conceived the study and designed the experiments; YX, and XW, and XW designed the constructs used in this study; WY, YX, and WY performed the *in vivo* studies; YY and YX helped to perform FACS analysis and western blots; XW, LL, and DP generated genetic modified mouse strains in this study; YL, BX, PL, XL, LL, and DW contributed the discussion part of the manuscript; XL and BX provided vital new reagents and revised the manuscript; and PL, YY, and DP discussed and wrote the manuscript. All authors read and approved the final manuscript.

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