

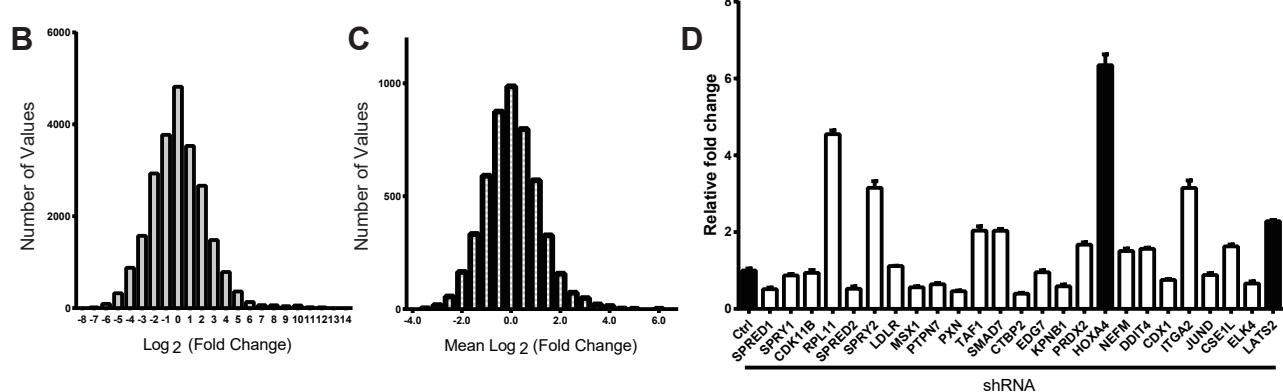
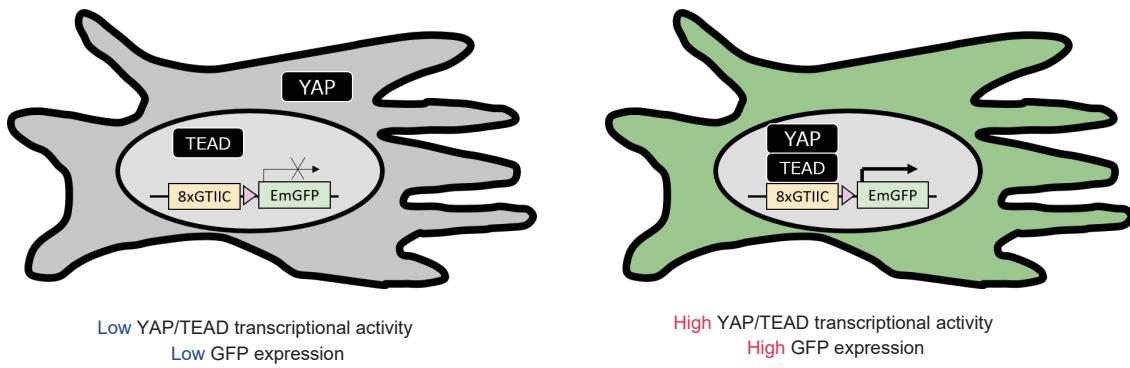
# **Appendix for:**

## **Homeobox A4 Suppresses Vascular Remodeling as a Novel Regulator of YAP/TEAD Transcriptional Activity**

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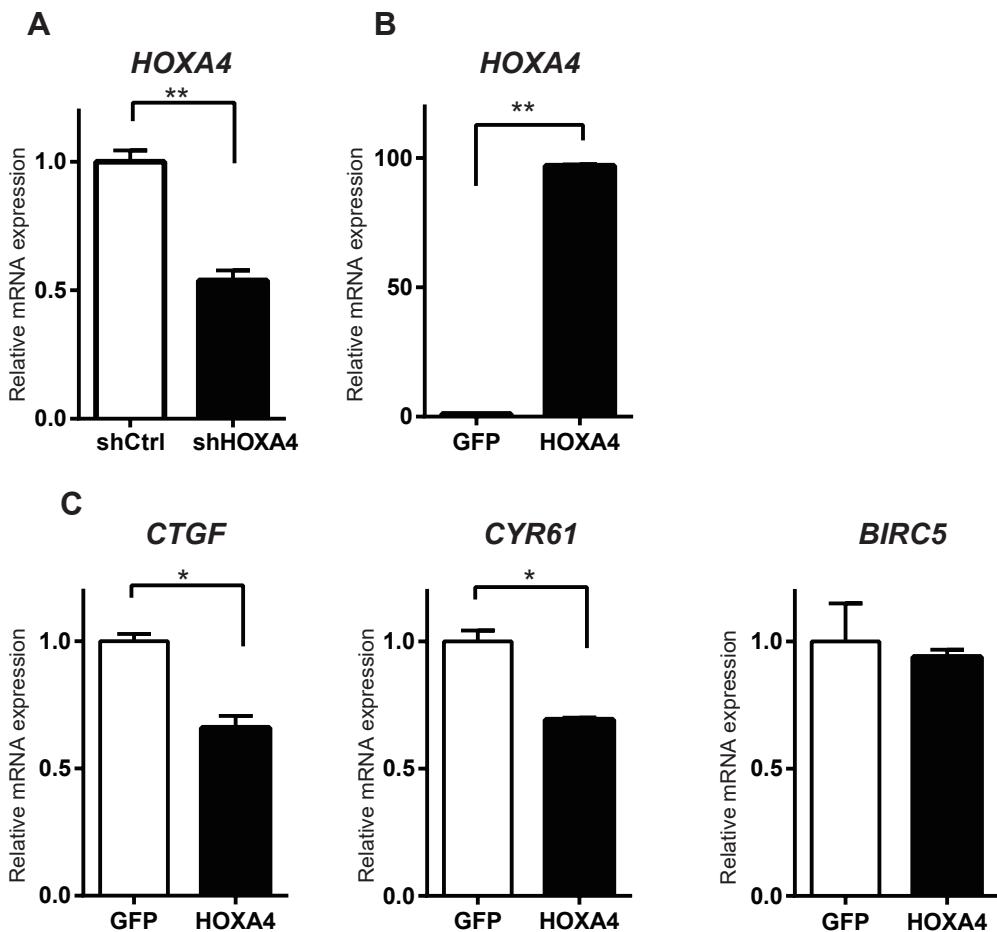
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**A**

### Appendix Figure S1 - Identification of novel modulators of YAP/TEAD transcriptional activity using pooled shRNA screen.

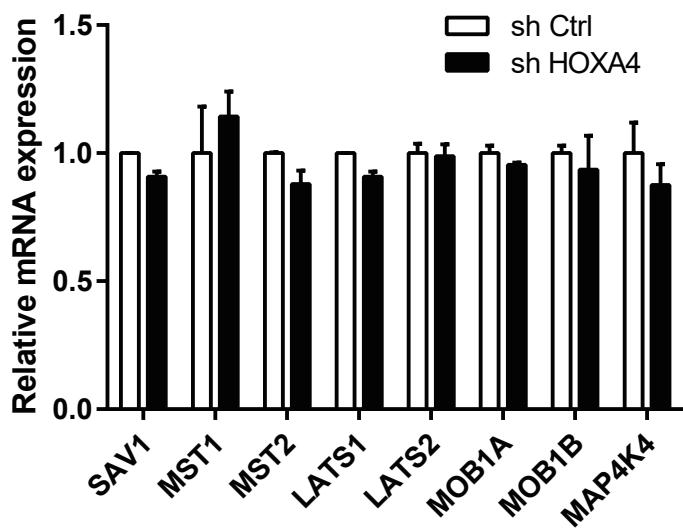
- A**, Schematic drawing of 8xGTIIC-EmGFP 293T cell line carrying eight TEAD-binding sequences upstream of a minimal promoter and EmGFP.
- B** and **C**, Distribution of read abundance per shRNA (B) or average read abundance of shRNA targeting each gene (C) followed an approximately Gaussian curve.
- D**, Validation of YAP/TEAD transactivation by knockdown of candidate genes using 8xGTIIC luciferase assay (n=3). Data are presented as mean  $\pm$  SEM.



**Appendix Figure S2 - HOXA4 attenuated the expression of YAP/TEAD target genes.**

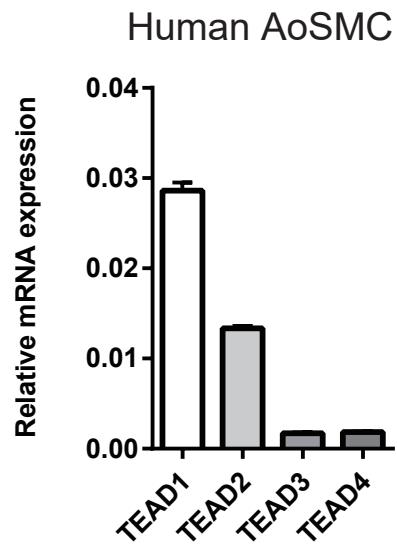
**A**, Knockdown efficacy of shHOXA4 vector in HEK 293T cells assessed by quantitative real-time PCR (n=3).

**B** and **C**, Quantitative real-time PCR analysis of HOXA4 (B) and YAP/TEAD target gene expression (C) in HEK 293T cells transfected with HOXA4 expression vector (n=3). All data are presented as mean  $\pm$  SEM.



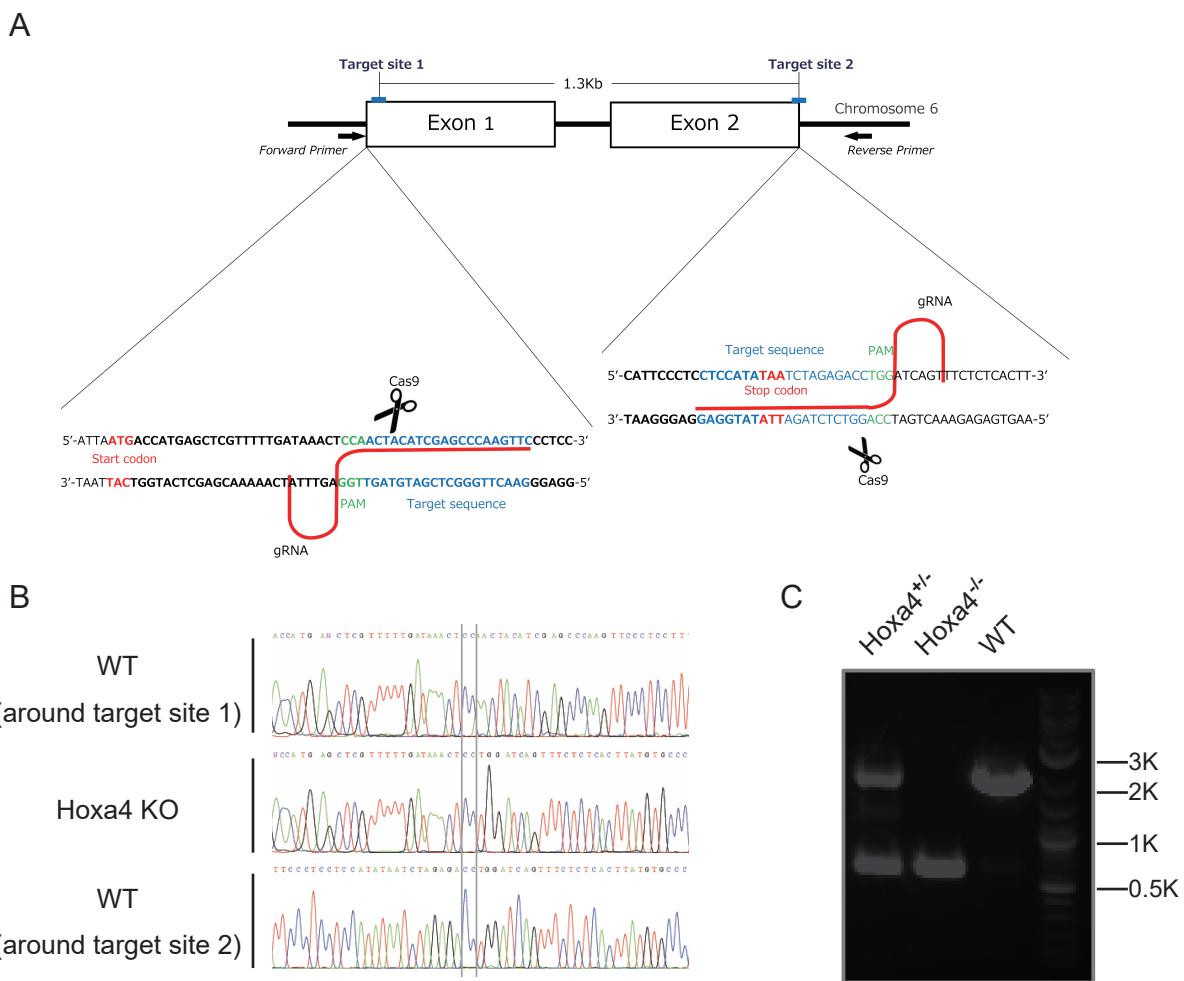
**Appendix Figure S3 - Gene expression levels of canonical Hippo pathway kinases were unaffected by HOXA4.**

Quantitative real-time PCR analysis of core Hippo pathway kinases in HEK 293T cells transfected with shHOXA4 (n=3). Data are presented as mean  $\pm$  SEM.



**Appendix Figure S4 - Expression of TEAD family genes.**

Quantitative real-time PCR analysis of TEAD family genes in human vascular smooth muscle cells (n=3). Data are presented as mean  $\pm$  SEM.

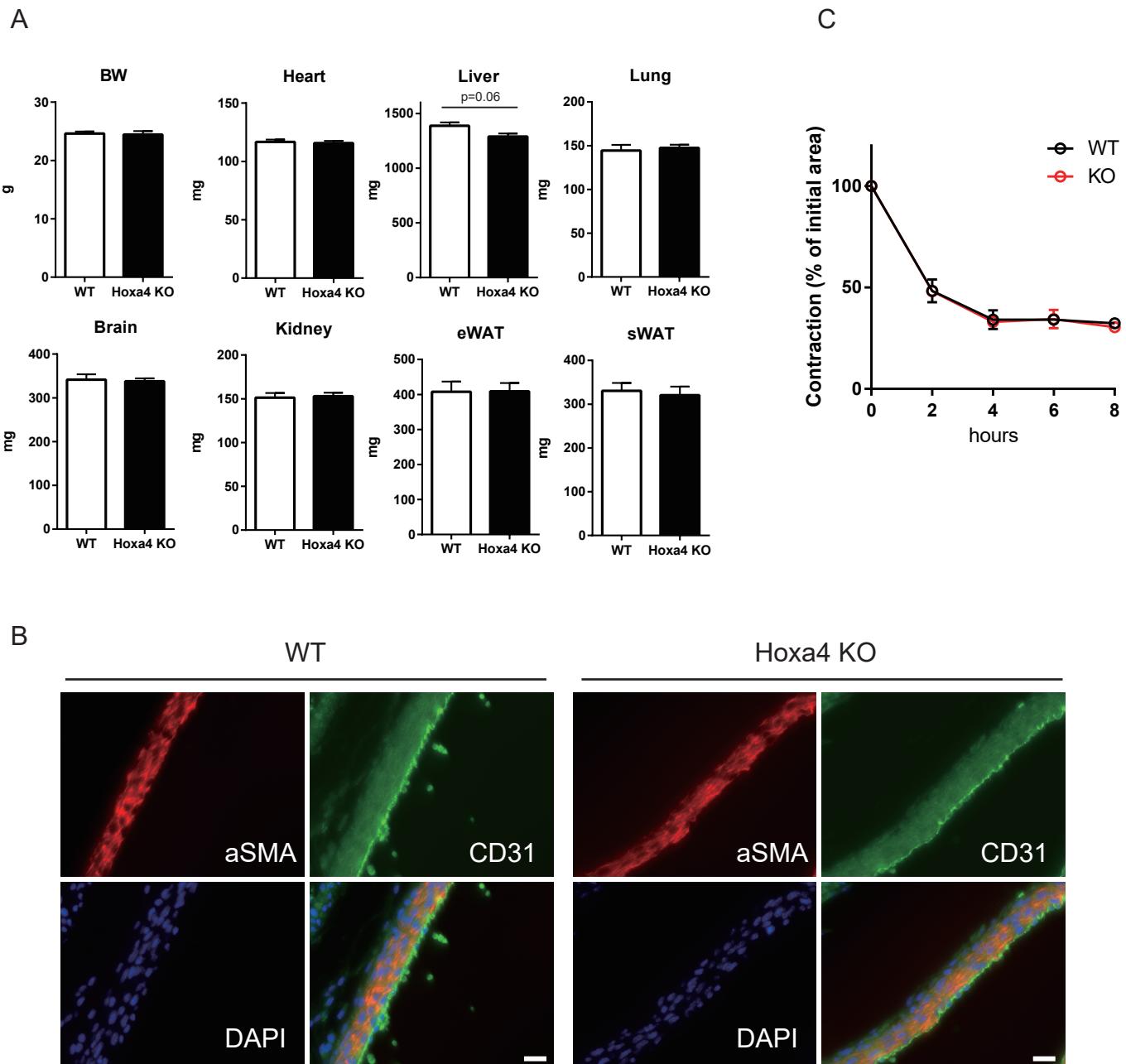


## Appendix Figure S5 - Generation of Hoxa4 deficient mice.

### A, Strategy to generate Hoxa4 KO mice.

**B**, DNA sequence around the target site of Cas9 cleavage in the knockout allele.

### C, PCR analysis of mouse tail genome.



### Appendix Figure S6 – Characteristics of Hoxa4-deficient mice.

**A**, Organ weight of 9-week-old male mice (WT: n=6, Hoxa4: n=8). Data are presented mean  $\pm$  SEM and analyzed by unpaired two-tailed Mann–Whitney test. BW; body weight, eWAT; epididymal white adipose tissue, sWAT; subcutaneous white adipose tissue.

**B**, Representative immunofluorescent staining images for  $\alpha$ -SMA and CD31 on dorsal aorta of embryonic day 18.5 (E18.5). Bars indicate 20  $\mu$ m.

**C**, Collagen gel contraction assay of primary vascular smooth muscle cells harvested from 8-week-old male mice (n=3). Data are presented as mean  $\pm$  SEM.