

**SUPPLEMENTAL MATERIAL FOR**

**SMAD4 maintains the Fluid Shear Stress set point to protect against Arterial-Venous Malformations**

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Mariona Graupera<sup>4</sup>, Gergana Dobрева<sup>2,3</sup>, Martin A. Schwartz<sup>5</sup>, Roxana Ola<sup>1\*</sup>

## Supplemental Figure Legends

### Supplemental Figure 1. *SMAD4*KD augments flow-induced gene expression.

(A) qPCR for the most 10 significant upregulated genes upon 24 hours 12 DYNES/cm<sup>2</sup> FSS in *CTRL* versus *SMAD4* siRNA HUVECs (n = 3/group): *ACKR4* (Atypical Chemokine Receptor 4); *APLNR* (Apelin receptor); *FBLN2* (EGF Containing Fibulin Extracellular Matrix Protein 2); *ITGB4* (Integrin Subunit Beta 4); *ELN* (Elastin); *MRAS* (Muscle RAS Oncogene Homolog); *PLCG2* (Phospholipase C Gamma 2); *KLF4* (Kruppel-Like Factor 4); *SLCO2A1* (Solute Carrier Organic Anion Transporter Family Member 2A1); *TNXB* (Tenascin XB). Data are represented as mean ± SEM: ns- non-significant, \**P*<0.05, \*\**P*<0.01, \*\*\**P*<0.001. One-way Anova.

### Supplemental Figure 2. Loss of *SMAD4* leads to dysregulated cell cycle in vitro and in vivo.

(A) Representative immunofluorescence staining of *CTRL* versus *SMAD4* siRNAs HUVECs grown in static versus 1 or 12 DYNES/cm<sup>2</sup> for EdU (green) and DAPI (blue). (B) Representative co-labeling of postnatal day 6 (P6) Tx induced *Smad4 fl/fl* and *Smad4<sup>iAEC</sup>* retinas for EdU+ nuclei (green-upper panel), and EdU (green), ERG (white), and Isolectin B4 (IB4-red) (lower panel). (C) Quantification of the total number of ECs (ERG+) in representative images shown in B. (D) S-phase ratio (EdU+/ERG+) per total ECs (ERG+) in capillaries of *Smad4 fl/fl* and *Smad4<sup>iAEC</sup>* retinas engaged or not in AVMs. (E) Representative co-labeling of postnatal day 6 (P6) Tx induced *Smad4 fl/fl* and *Smad4<sup>iAEC</sup>* retinas for phospho Histone 3 (PH3-blue) (upper panel) and PH3 (blue), ERG (white), and Isolectin B4 (IB4-red) (lower panel). Yellow arrowheads in B and E mark ECs+ for EdU and PH3, respectively in capillaries. Red/blue arrowheads in B mark the EdU+ ECs in arteries/veins. (F) M-phase ratio (PH3+/ERG+) per total ECs (ERG+) in capillaries of *Smad4 fl/fl* and *Smad4<sup>iAEC</sup>* retinas engaged or not in AVMs. n=9 (3 images (200-600 cells/image)/retina/group). Scale Bars: 100µm in A,B,E. a: artery, v: vein. Data are represented as mean ± SEM. \**P*<0.05, \*\**P*<0.01, \*\*\**P*<0.001. One-way Anova (C,D,F).

**Supplemental Figure 3. KLF4 regulates cell size but not VE Cadherin at cell junctions.**

(A,B) Quantification of cell area in *CTRL*, *SMAD4*, *KLF4* and *SMAD4;KLF4* siRNAs HUVECs subject to 12 DYNES/cm<sup>2</sup> (A) and 1 DYNE/cm<sup>2</sup> (B) for 24 hours (n=8 images (70-140 cells/image) per 4 independent experiments/group). (C) Representative VE-Cadherin staining (negative images) of *CTRL*, *SMAD4*, *KLF4* and *SMAD4;KLF4* siRNAs HUVECs subject to 12 DYNES/cm<sup>2</sup> for 24 hours. (D) Quantification of VE-Cadherin labeling intensity from experiments in C (n=6 images per 3 independent experiments/group). Scale Bars: 100µm. Data are represented as mean ± SEM. \**P*<0.05,\*\**P*<0.01,\*\*\**P*<0.001, ns- non-significant. One-way Anova (A,B,D).

**Supplemental Figure 4. Validation of KLF4 antibody in vivo.**

(A) Representative images of the vascular plexus in retinas from Tx induced P6 *Klf4 fl/fl* and *Klf4<sup>iΔEC</sup>* labelled for KLF4 (green) and IB4 (white). Scale Bars: 50 µm.

**Supplemental Figure 5. Increased PI3K signaling mediates EC aberrant responses in *Smad4*<sup>ECko</sup>.**

(A) Representative WB for pAkt in HUVECs subject to 12 DYNES/cm<sup>2</sup> treated with PBS or Pictilisib (PI3Ki-75nM) for 4 hours (n=3/group). (B) Representative VE-Cadherin staining (negative images) of *CTRL* and *SMAD4* siRNAs HUVECs subject to 24 hours 12 DYNES/cm<sup>2</sup> and treated with PBS, PI3K inhibitor or with *AKT* siRNA. Flow direction: right to left. (C) Quantification of the length/width ratio (n=6 average of images (70-140 cells/image) per 3 independent experiments/group) and of EC alignment parallel to flow direction (%) (n=12 average of images (100-240 cells/image) per 3 independent experiments/group). (D) Representative images of retinas from P6 *Smad4 fl/fl* and *Smad4<sup>iΔEC</sup>* from pups treated with PBS or PI3K inhibitor labeled for ERG (white), GOLPH4 (red) and IB4 (green)-upper panel and ERG (white), GOLPH4 (red) and IB4 (green line)-lower panel. Yellow arrowheads mark the EC orientation within the AVMs. (E) Quantification of EC polarization: against or with flow and neutral (non-oriented) in capillaries and AVMs from P6 retinas of *Smad4 fl/fl* and *Smad4<sup>iΔEC</sup>* pups treated with PBS or PI3Ki (n=3 retinas/group). (F) S-phase ratio (EdU+/ERG+) per

total ECs (ERG+) in the vascular plexus of *Smad4*<sup>iΔEC</sup> retinas in PBS versus PI3Ki (Pictilisib) treated pups (n=8 (2 images (200-600 cells/image)/retina/group)). (G) S-phase ratio (EdU+) per total DAPI+ cells in response to 24 hours 12 DYNES/cm<sup>2</sup> of CTRL and *SMAD4* siRNAs HUVECs treated with PBS versus PI3Ki (n=6 (2 images (200-300 cells/image)/experiment/group)). (H,I) *KLF4* mRNA expression by qPCR in HUVECs subject to 12 DYNES/cm<sup>2</sup> and treated with PBS versus Pictilisib (H) (n=5/group) and in CTRL, *CD31*, *KDR*, and *CDH5* siRNAs HUVECs (I) (n=4/group). Scale Bars: 100μm in D. a: artery, v: vein. Data are represented as mean ± SEM. \**P*<0.05, \*\**P*<0.01, \*\*\**P*<0.001, ns- non-significant. One-way Anova (C,E,G), Mann-Whitney test (F,H,I).

### Supplemental Figure 6. *Klf4* inactivation restores arterial identity in *Smad4*<sup>iΔEC</sup> retinas

(A) qPCR for *CCNA2*, *CDKN1A* and *CDKN2D* in CTRL versus *SMAD4* siRNAs HUVECs grown in static versus subject to 12 DYNES/cm<sup>2</sup> (n=3/group). (B,D) Representative confocal images of labeled retinas for CX37 (white) (B), CX40 (green) (D) and IB4 (red) from Tx induced P6 *fl/fl*, *Smad4*<sup>iΔEC</sup>, *Klf4*<sup>iΔEC</sup> and *Smad4;Klf4*<sup>iΔEC</sup>. Yellow arrowheads indicate AVMs. Blue arrowheads indicate expression of arterial markers in arteries and arterioli. (C,E) Quantification of CX37 (C) and CX40 (E) signals in the vascular plexus from the indicated genotypes (n = 4 retinas/group). Scale Bars in B,D: 100μm. a: artery, v: vein. Data are represented as mean ± SEM. n.s- non-significant, \**P*<0.05, \*\**P*<0.01, \*\*\**P*<0.001. One-way Anova (A,C,E).

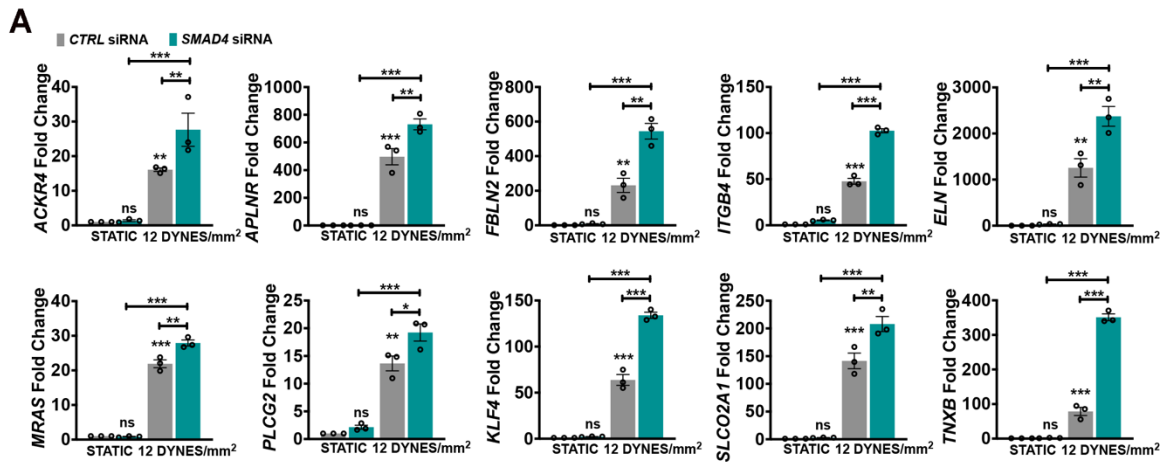
### Supplemental Figure 7. Palbociclib reduces vascular density in *Smad4*<sup>iΔEC</sup> retinas.

(A) Representative WB images for the indicated proteins of whole lung lysates from pups treated with DMSO and Palbociclib (n = 3/group). (B) Confocal images of vascular front of P6 *Smad4 fl/fl* and *Smad4*<sup>iΔEC</sup> retinas treated with DMSO or Palbociclib labeled for IB4 (red) - upper panel, EdU (green) and ERG (white) -middle panel and IB4/ERG/EdU (lower panel) of the vascular front in indicated genotypes. (C) Quantification of the vascular density, the number of EdU+/ERG+ ECs per total number of ERG+ ECs (%) and of the total number of ERG+ ECs at the vascular front of *Smad4*<sup>iΔEC</sup> retinas



DMSO or Palbociclib treated. Scale Bars in **B**: 50 $\mu$ m. Data are represented as mean  $\pm$  SEM. n.s- non-significant, \* $P$ <0.05, \*\* $P$ <0.01, \*\*\* $P$ <0.001. One-way Anova (**C**).

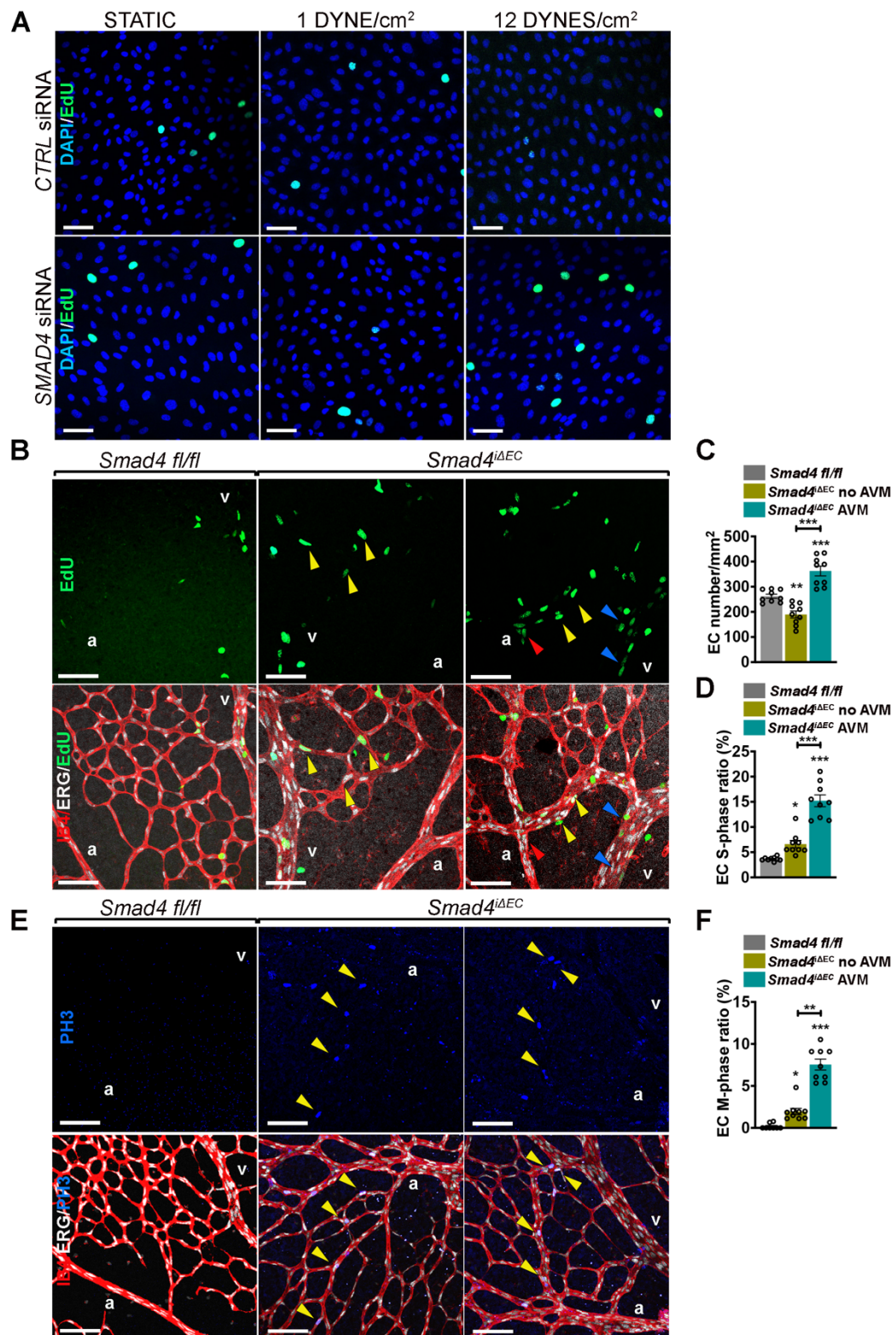
## Supplemental Figure 1



### Supplemental Figure 1. *SMAD4*KD augments flow-induced gene expression.

(A) qPCR for the most 10 significant upregulated genes upon 24 hours 12 DYNES/cm<sup>2</sup> FSS in *CTRL* versus *SMAD4* siRNA HUVECs (n = 3/group): *ACKR4* (Atypical Chemokine Receptor 4); *APLNR* (Apelin receptor); *FBLN2* (EGF Containing Fibulin Extracellular Matrix Protein 2); *ITGB4* (Integrin Subunit Beta 4); *ELN* (Elastin); *MRAS* (Muscle RAS Oncogene Homolog); *PLCG2* (Phospholipase C Gamma 2); *KLF4* (Kruppel-Like Factor 4); *SLCO2A1* (Solute Carrier Organic Anion Transporter Family Member 2A1); *TNXB* (Tenascin XB). Data are represented as mean ± SEM: ns- non-significant, \**P*<0.05, \*\**P*<0.01, \*\*\**P*<0.001. One-way Anova.

Supplemental Figure 2

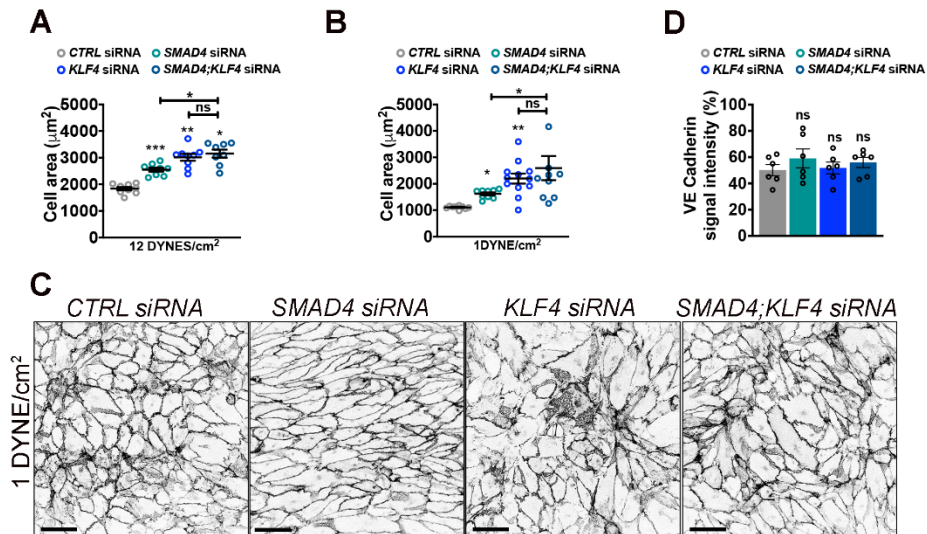


Supplemental Figure 2. Loss of *SMAD4* leads to dysregulated cell cycle in vitro and in vivo.

(A) Representative immunofluorescence staining of CTRL versus *SMAD4* siRNAs HUVECs grown in static versus 1 or 12 DYNES/cm<sup>2</sup> for EdU (green) and DAPI (blue). (B) Representative co-labeling of postnatal day 6 (P6) Tx induced *Smad4* fl/fl and *Smad4*<sup>ΔEC</sup> retinas for EdU+ nuclei (green-upper panel), and EdU (green), ERG (white), and Isolectin B4 (IB4-red) (lower panel). (C) Quantification

of the total number of ECs (ERG+) in representative images shown in **B**. **(D)** S-phase ratio (EdU+/ERG+) per total ECs (ERG+) in capillaries of *Smad4 fl/fl* and *Smad4<sup>iΔEC</sup>* retinas engaged or not in AVMs. **(E)** Representative co-labeling of postnatal day 6 (P6) Tx induced *Smad4 fl/fl* and *Smad4<sup>iΔEC</sup>* retinas for phospho Histone 3 (PH3-blue) (upper panel) and PH3 (blue), ERG (white), and Isolectin B4 (IB4-red) (lower panel). Yellow arrowheads in **B** and **E** mark ECs+ for EdU and PH3, respectively in capillaries. Red/blue arrowheads in **B** mark the EdU+ ECs in arteries/veins. **(F)** M-phase ratio (PH3+/ERG+) per total ECs (ERG+) in capillaries of *Smad4 fl/fl* and *Smad4<sup>iΔEC</sup>* retinas engaged or not in AVMs. n=9 (3 images (200-600 cells/image)/retina/group). Scale Bars: 100μm in **A,B,E**. a: artery, v: vein. Data are represented as mean ± SEM. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ . One-way Anova (**C,D,F**).

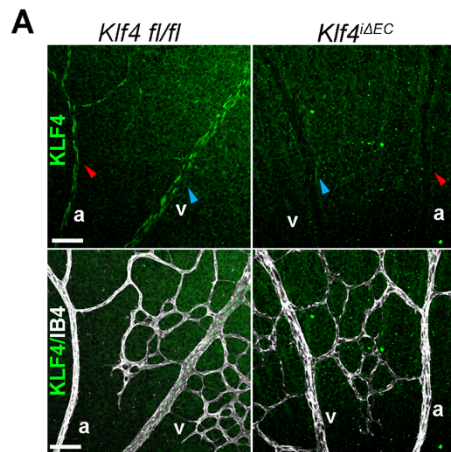
### Supplemental Figure 3



#### Supplemental Figure 3. KLF4 regulates cell size but not VE Cadherin at cell junctions.

(A,B) Quantification of cell area in *CTRL*, *SMAD4*, *KLF4* and *SMAD4;KLF4* siRNAs HUVECs subject to 12 DYNES/cm<sup>2</sup> (A) and 1 DYNE/cm<sup>2</sup> (B) for 24 hours (n=8 images (70-140 cells/image) per 4 independent experiments/group). (C) Representative VE-Cadherin staining (negative images) of *CTRL*, *SMAD4*, *KLF4* and *SMAD4;KLF4* siRNAs HUVECs subject to 12 DYNES/cm<sup>2</sup> for 24 hours. (D) Quantification of VE-Cadherin labeling intensity from experiments in C (n=6 images per 3 independent experiments/group). Scale Bars: 100 $\mu\text{m}$ . Data are represented as mean  $\pm$  SEM. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ , ns- non-significant. One-way Anova (A,B,D).

## Supplemental Figure 4

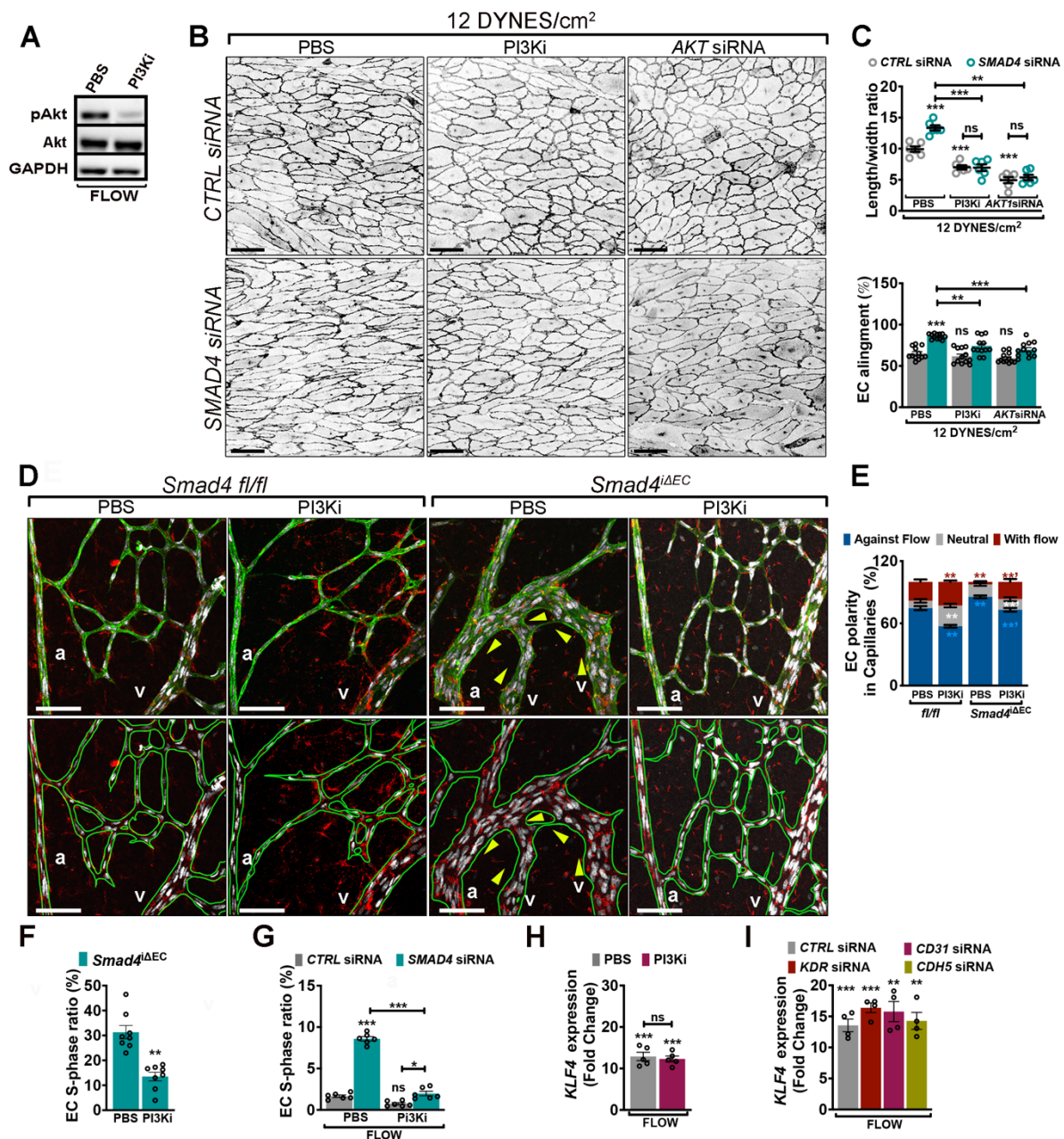


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(A) Representative images of the vascular plexus in retinas from Tx induced P6 *Klf4 fl/fl* and *Klf4<sup>iΔEC</sup>* labelled for KLF4 (green) and IB4 (white). Scale Bars: 50  $\mu$ m.



## Supplemental Figure 5



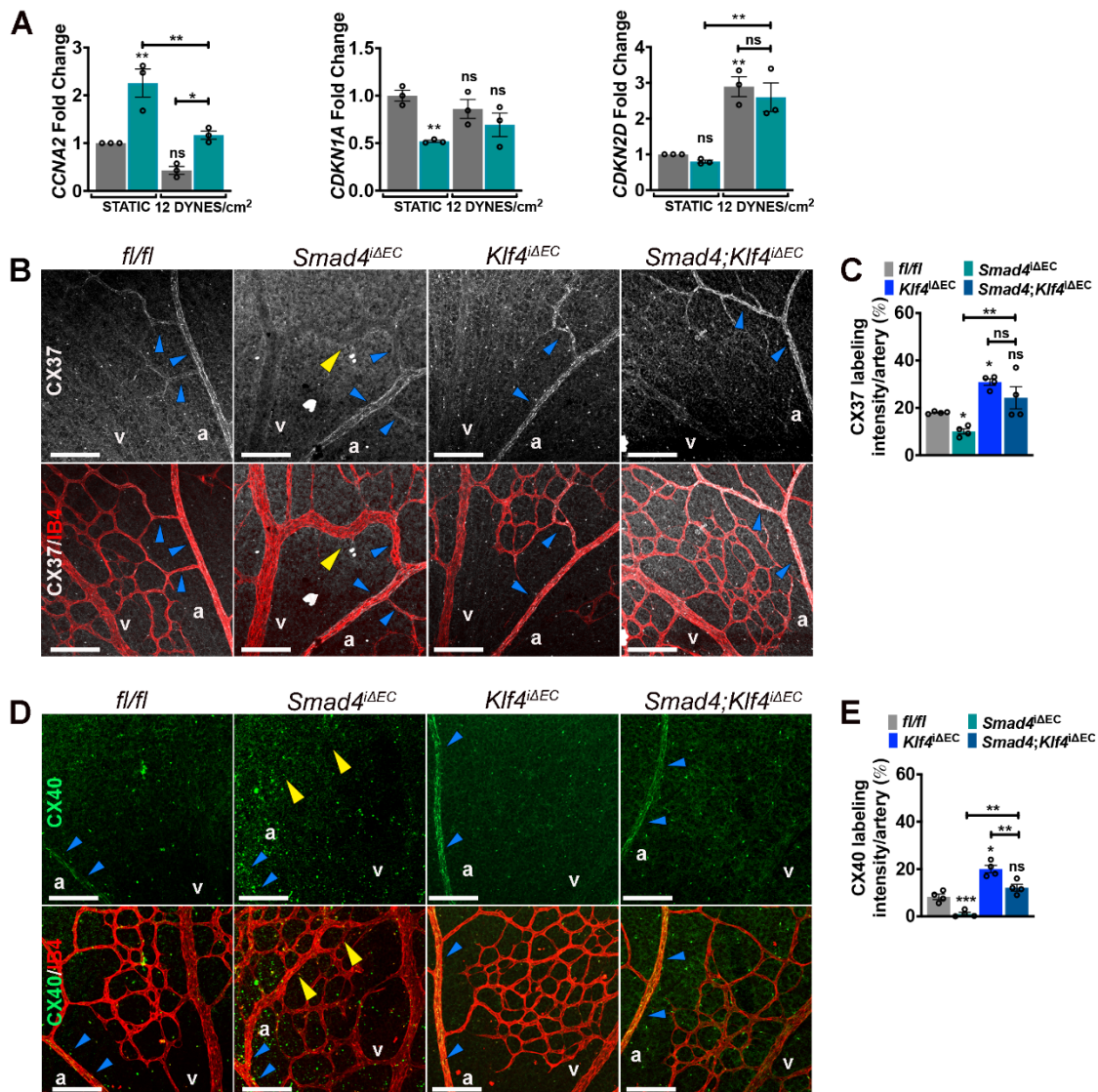
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(A) Representative WB for pAkt in HUVECs subject to 12 DYNES/cm<sup>2</sup> treated with PBS or Pictilisib (PI3Ki-75nM) for 4 hours (n=3/group). (B) Representative VE-Cadherin staining (negative images) of CTRL and SMAD4 siRNAs HUVECs subject to 24 hours 12 DYNES/cm<sup>2</sup> and treated with PBS, PI3K inhibitor or with AKT siRNA. Flow direction: right to left. (C) Quantification of the length/width ratio (n=6 average of images (70-140 cells/image) per 3 independent experiments/group) and of EC alignment parallel to flow direction (%) (n=12 average of images (100-240 cells/image) per 3 independent experiments/group). (D) Representative images of retinas from P6 *Smad4*<sup>fl/fl</sup> and *Smad4*<sup>iAEC</sup> from pups treated with PBS or PI3K inhibitor labeled for ERG (white), GOLPH4 (red) and IB4 (green)-upper panel and ERG (white), GOLPH4 (red) and IB4 (green line)-lower panel. Yellow arrowheads mark the EC orientation within the AVMs. (E) Quantification of EC polarization: against or with flow and neutral (non-oriented) in capillaries and AVMs from P6 retinas of *Smad4*<sup>fl/fl</sup> and *Smad4*<sup>iAEC</sup> pups treated with PBS or PI3Ki (n=3 retinas/group). (F) S-phase ratio (EdU+/ERG+) per total ECs (ERG+) in the vascular plexus of *Smad4*<sup>iAEC</sup> retinas in PBS versus PI3Ki (Pictilisib) treated

pups (n=8 (2 images (200-600 cells/image)/retina/group). **(G)** S-phase ratio (EdU+) per total DAPI+ cells in response to 24 hours 12 DYNES/cm<sup>2</sup> of *CTRL* and *SMAD4* siRNAs HUVECs treated with PBS versus PI3Ki (n=6 (2 images (200-300 cells/image)/experiment/group). **(H,I)** *KLF4* mRNA expression by qPCR in HUVECs subject to 12 DYNES/cm<sup>2</sup> and treated with PBS versus Pictilisib **(H)** (n=5/group) and in *CTRL*, *CD31*, *KDR*, and *CDH5* siRNAs HUVECs **(I)** (n=4/group). Scale Bars: 100µm in **D**. **a**: artery, **v**: vein. Data are represented as mean ± SEM. \**P*<0.05, \*\**P*<0.01, \*\*\**P*<0.001, ns- non-significant. One-way Anova (**C,E,G**), Mann-Whitney test (**F,H,I**).



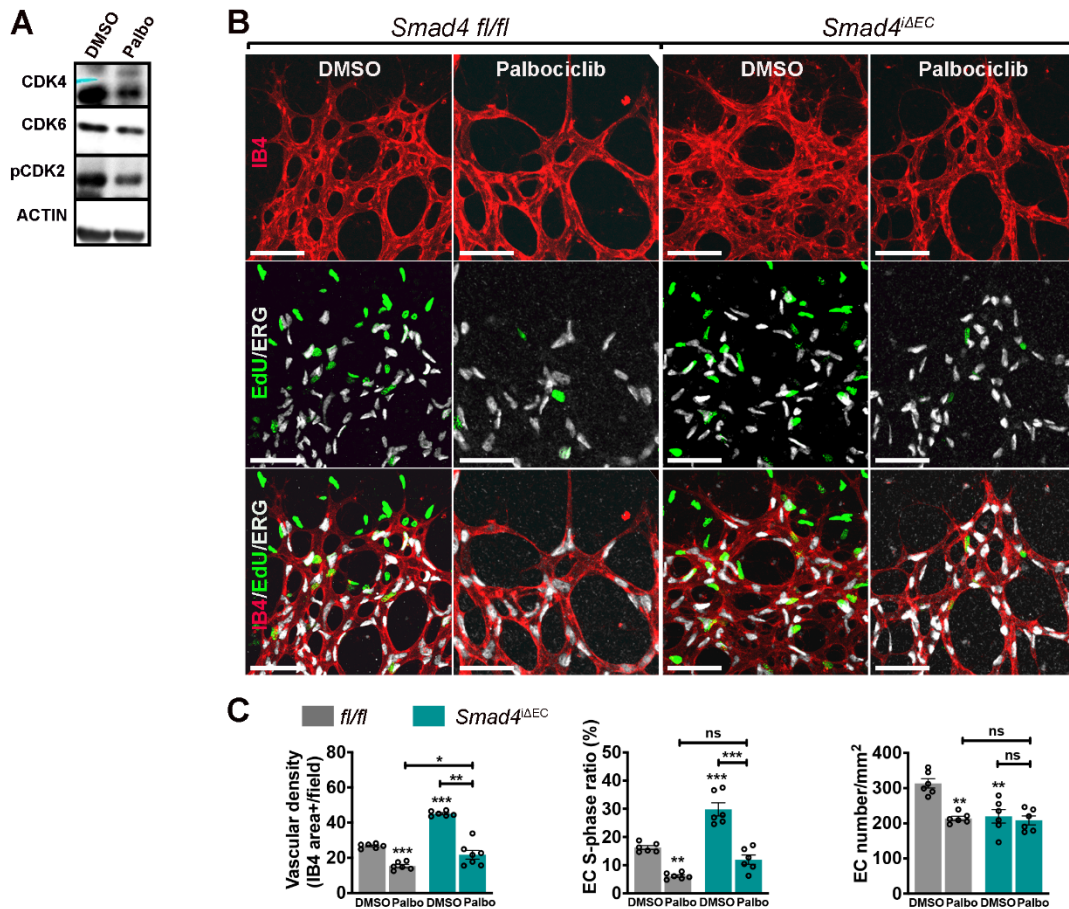
## Supplemental Figure 6



### Supplemental Figure 6. *Klf4* inactivation restores arterial identity in *Smad4*<sup>ΔEC</sup> retinas

(A) qPCR for *CCNA2*, *CDKN1A* and *CDKN2D* in CTRL versus *SMAD4* siRNAs HUVECs grown in static versus subject to 12 DYNES/cm<sup>2</sup> (n=3/group). (B,D) Representative confocal images of labeled retinas for CX37 (white) (B), CX40 (green) (D) and IB4 (red) from Tx induced P6 *fl/fl*, *Smad4*<sup>ΔEC</sup>, *Klf4*<sup>ΔEC</sup> and *Smad4;Klf4*<sup>ΔEC</sup>. Yellow arrowheads indicate AVMs. Blue arrowheads indicate expression of arterial markers in arteries and arterioli. (C,E) Quantification of CX37 (C) and CX40 (E) signals in the vascular plexus from the indicated genotypes (n = 4 retinas/group). Scale Bars in B,D: 100μm. a: artery, v: vein. Data are represented as mean ± SEM. n.s- non-significant, \*P<0.05, \*\*P<0.01, \*\*\*P<0.001. One-way Anova (A,C,E).

## Supplemental Figure 7



### Supplemental Figure 7. Palbociclib reduces vascular density in *Smad4*<sup>iΔEC</sup> retinas.

(A) Representative WB images for the indicated proteins of whole lung lysates from pups treated with DMSO and Palbociclib ( $n = 3/\text{group}$ ). (B) Confocal images of vascular front of P6 *Smad4*<sup>fl/fl</sup> and *Smad4*<sup>iΔEC</sup> retinas treated with DMSO or Palbociclib labeled for IB4 (red) - upper panel, EdU (green) and ERG (white) -middle panel and IB4/ERG/EdU (lower panel) of the vascular front in indicated genotypes. (C) Quantification of the vascular density, the number of EdU+/ERG+ ECs per total number of ERG+ ECs (%) and of the total number of ERG+ ECs at the vascular front of *Smad4*<sup>iΔEC</sup> retinas DMSO or Palbociclib treated. Scale Bars in B: 50 $\mu\text{m}$ . Data are represented as mean  $\pm$  SEM. n.s- non-significant, \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ . One-way Anova (C).

## Supplemental Methods

**Supplemental Table1: List of mouse and human primers used in the study.**

Gene name (Protein name)	Forward Sequence	Reverse Sequence
<b>Primers for HUVECs</b>		
<i>ACKR4</i> (human-Atypical Chemokine Receptor 4)	GTTTTTCGTCATTGGACTTGCAG	GCTACAGCCAAATTCAGGATGT
<i>APLNR</i> (human-Apelin Receptor)	CTCTGGACCGTGTTTCGGAG	GGTACGTGTAGGTAGCCCACA
<i>CCNA2</i> (human-Cyclin A2)	ACCCAGAAAACCATTGGTCC	CATTTAACCTCCATTTCCCTAAGGT
<i>CCNB1</i> (human-Cyclin B1)	AATAAGGCGAAGATCAACATG GC	TTTGTTACCAATGTCCCCAAGAG
<i>CCNB2</i> (human-Cyclin B2)	CCGACGGTGTCCAGTGATTT	TGTTGTTTTGGTGGGTGAACT
<i>CDK1</i> (human-CDK1)	AAACTACAGGTCAAGTGGTAGC C	TCCTGCATAAGCACATCCTGA
<i>CDKN1A</i> (human-p21)	TGTCCGTCAGAACCCATGC	AAAGTCGAAGTTCATCGCTC
<i>CDKN2A</i> (human-p16)	CAACGCACCGAATAGTTACG	AGCACCACCAGCGTGTC
<i>CDKN2B</i> (human-p15)	CACCGTTGGCCGTAAACTTAAC	TAATGAAGCTGAGCCCAGTCT
<i>GJA4</i> (human- CX37)	ACACCCACCCTGGTCTACC	CACTGGCGACATAGGTGCC
<i>GJA5</i> (human-CX40)	CCGTGGTAGGCAAGGTCTG	ATCACACCGGAAATCAGCCTG
<i>GJA1</i> (human-CX43)	GGTGACTGGAGCGCCTTAG	GCGCACATGAGAGATTGGGA
<i>ELN</i> (human-Elastin)	GCAGGAGTTAAGCCCAAGG	TGTAGGGCAGTCCATAGCCA
<i>EPHRINB2</i> (human-Ephrin B2)	TATGCAGAACTGCGATTTCCAA	TGGGTATAGTACCAGTCCTTGTC
<i>FBLN2</i> (human-Fibulin 2)	ACTGTGGGTTCTTACCACTGT	CCACCTGGGAAAATTCTGACTT
<i>FLT4</i> (human-VEGFR3)	TGCACGAGGTACATGCCAAC	GCTGCTCAAAGTCTCTCACGAA
<i>GAPDH</i> (human-GAPDH)	CTGGGCTACACTGAGCACC	AAGTGGTCGTTGAGGGCAATG
<i>HPRT</i> (human-Hypoxanthine Phosphoribosyltransferase 1)	GACCAGTCAACAGGGGACAT	CCTGACCAAGGAAAGCAAAG
<i>ITGB4</i> (human-Integrin beta-4)	CTCCACCGAGTCAGCCTTC	CGGGTAGTCCTGTGTCCTGTA
<i>KLF4</i> (human-KLF4)	CCCACATGAAGCGACTTCCC	CAGGTCCAGGAGATCGTTGAA

<i>MRAS</i> (human-Ras-related protein M-Ras)	TTCCTCATCGTCTACTCCGTC	AGGATCATCGGGAATGACTCC
<i>PECAM-1</i> (human-PECAM1)	AAGTGGAGTCCAGCCGCATATC	ATGGAGCAGGACAGGTTTCAGTC
<i>PLCG2</i> (human-PLCG2)	CATCCTATATGGCACTCAGTTC G	TCCTGGTGTAAGATTTTCAAGCC
<i>SLCO2A1</i> (human-SLCO2A1)	TCGGTCTTCGGCAACATTAAG	GCTCTTGAAGTAGGCGCTGTA
<i>SOX17</i> (human-SOX17)	GTGGAACCGCACGGAATTTG	GGAGATTCACACCGGAGTTCA
<i>TEK</i> (human-TIE2)	TTAGCCAGCTTAGTTCTCTGTGG	AGCATCAGATACAAGAGGTAGGG
<i>TNXB</i> (human-TN-X)	GCCCTGCTCACTTGGACTG	GGAGCCGTGCATTGTAGGAG
<i>CDH5</i> (human-VE-Cadherin)	QT00013244, Qiagen	
<i>KDR</i> (human-VEGFR2)	QT00069818, Qiagen	
<i>SMAD4</i> (human-SMAD4)	QT00013174, Qiagen	
<b>Primers for mouse lung ECs</b>		
<i>Gapdh</i> (mouse-GAPDH)	AGGTCGGTGTGAACGGATTTG	TGTAGACCATGTAGTTGAGGTCA
<i>Klf4</i> (mouse-KLF4)	QT00095431, Qiagen	
<i>Smad4</i> (mouse-SMAD4)	QT00130585, Qiagen	