

Differential endothelial cell cycle status in postnatal retinal vessels revealed using a novel PIP-FUCCI reporter and zonation analysis

Ziqing Liu^{1#}, Natalie T Tanke², Alexandra Neal¹, Tianji Yu¹, Tershona Branch¹, Arya Sharma¹, Jean G Cook³, Victoria L Bautch^{1,2,4,5*}

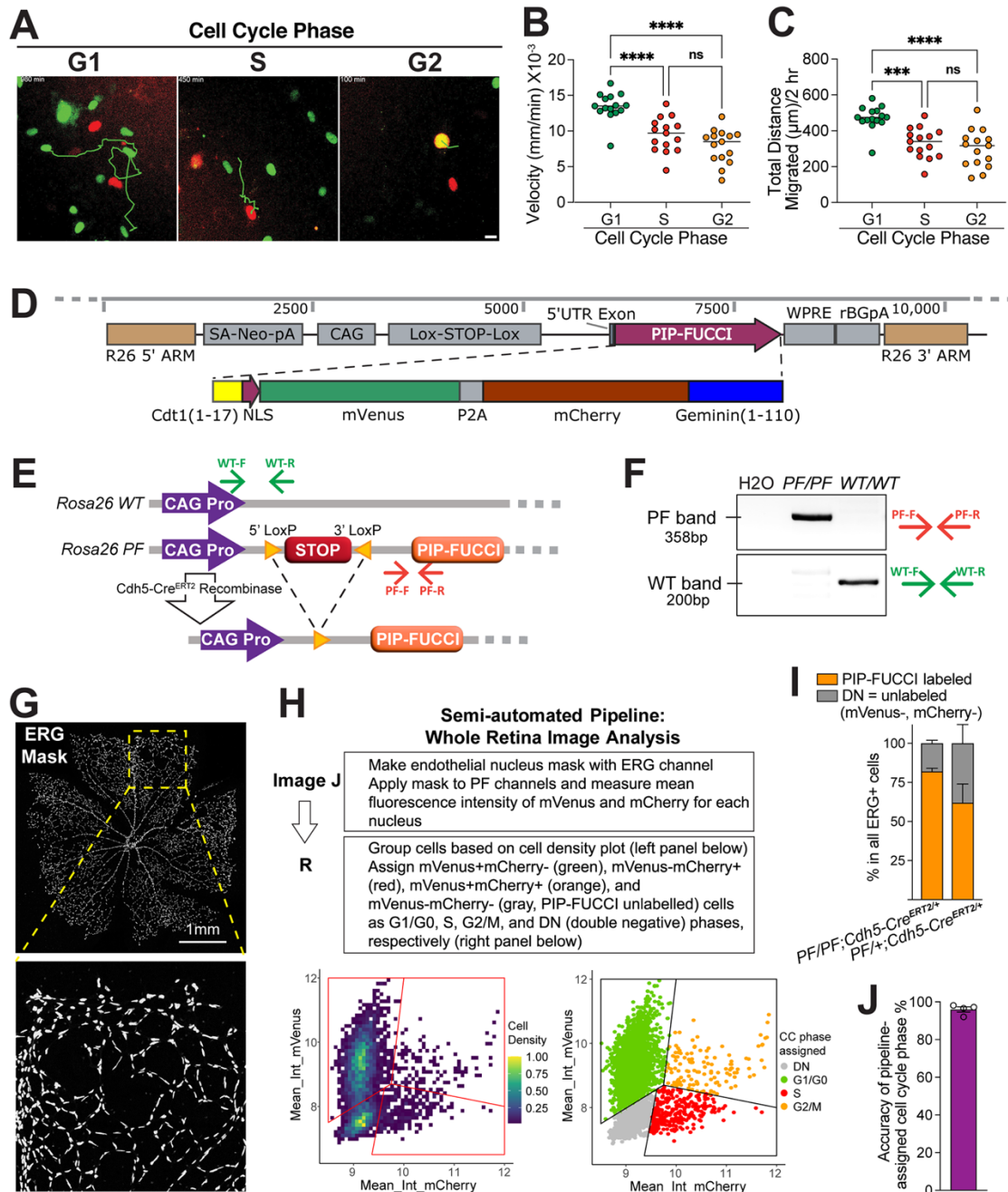
¹Department of Biology, ²Curriculum in Cell Biology and Physiology, ³Department of Biochemistry and Biophysics, ⁴McAllister Heart Institute, ⁵Lineberger Comprehensive Cancer Center, The University of North Carolina, Chapel Hill, NC USA

* corresponding author

present address: Dept of Physiology, Medical College of Wisconsin, Milwaukee WI

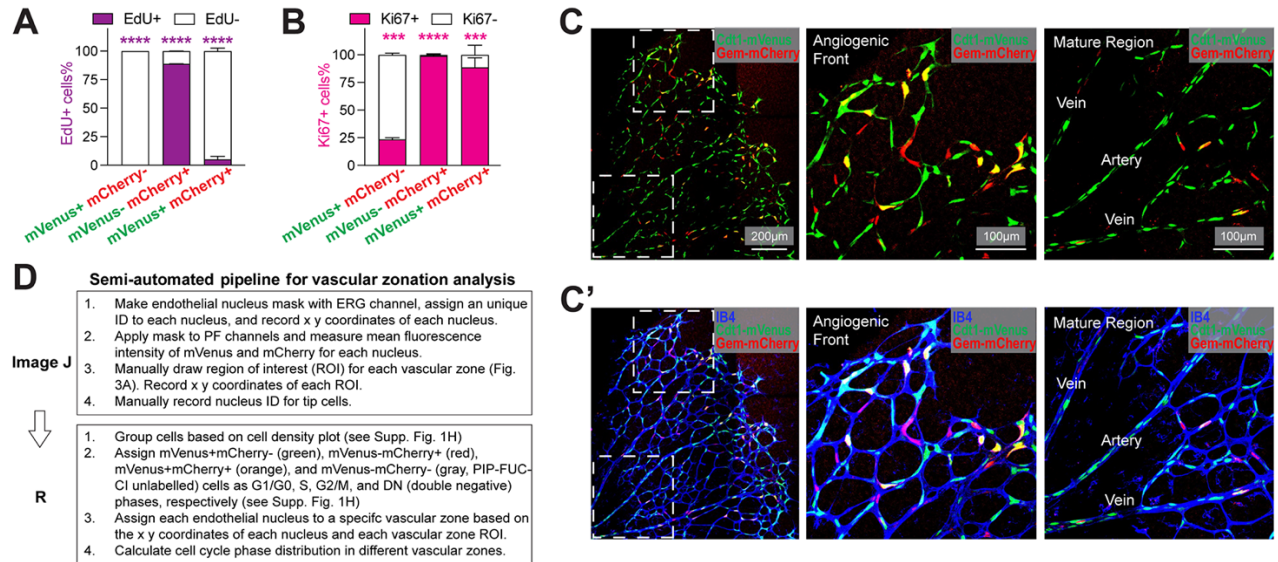
SUPPLEMENTAL FILE:

1. Supplemental Figures (2)
2. Supplemental Resource Tables (2)



Supplementary Figure 1. PIP-FUCCI reporter cells and mice and workflow for retina analysis. (A) Representative cell migration traces (green lines) for indicated cell cycle phases. Scale bar, 50 μ m. **(B)** Quantification of velocity per 2 hr segment for indicated cell cycle phases. (n=15 cells/phase, 3 replicate movies). **(C)** Quantification of total distance migrated per 2 hr segment for indicated cell cycle phases (n=15 cells/phase, 3 replicate movies). **** p < 0.0001, *** p < 0.001, by one-way ANOVA & Tukey's multiple comparisons test. **(D)** Construct used to generate the PIP-FUCCI reporter mouse line. **(E)** Location of genotyping primers. **(F)** Example of genotyping results on DNA agarose gel. **(G)** Example of ERG mask. **(H)** Workflow for whole retina

analysis with ERG mask. **(I)** Quantification of % PIP-FUCCI labeled cells in ERG+ endothelial cells from *PF/PF;Cdh5-Cre^{ERT2/+}* or *PF/+;Cdh5-Cre^{ERT2/+}* retinas (n= 2 pups for each genotype). **(J)** Accuracy of manual vs. algorithm-assigned cell cycle phases from the same subset of endothelial cells of the same retinal images (PIP-FUCCI labeled and stained for IB4 and ERG). n = 4 pups.



Supplementary Figure 2. Workflow for vascular zonation analysis of the retina.

(A) Quantification of retinal endothelial cells EdU labeled relative to PIP-FUCCI status (related to Fig. 2D-F). $n = 3$ pups. **** $p < 0.0001$ by two-way ANOVA & Sidak's multiple comparisons test. (B) Quantification of Ki67+ retinal endothelial cells relative to PIP-FUCCI status (related to Fig. 2G-I). $n = 2$ pups. **** $p < 0.0001$, *** $p < 0.001$ by two-way ANOVA & Sidak's multiple comparisons test. (C-C') Representative image of one leaflet of a *PF/PF; Cdh5-Cre^{ERT2/+}* retina stained for IB4 and ERG (same imaging fields as Fig. 2B, IB4 pseudo-colored blue in C'). Boxed areas in far left panel (scale bar, 200 μ m) magnified (scale bar, 100 μ m) in middle (Angiogenic Front) and far right (Mature Region) panels. (D) Workflow for semi-automated vascular zonation analysis of PIP-FUCCI retinal images with ERG mask.

SUPPLEMENTARY TABLES**Supplementary Table 1. PCR Primers**

Primer Name	Primer Sequence	Expected Products
Rosa26 WT-F	TGGAGTTGCAGATCACGAGG	WT = 225 bp
Rosa26 WT-R	TGTTTTGGAGGCAGGAAGCA	PF = no band
Rosa26 PF-F	GCTAACCATGTTTCATGCCTTC	WT = no band
Rosa26 PF-R	CGCCCTCGCCGGACACGCTGAAC	PF = 357 bp
GenCre F	GACCAGGTTTCGTTCACTCA	WT = no band
GenCre R	TAGCGCCGTAAATCAAT	Cre = 400 bp

Supplementary Table 2. Antibodies

Antibody	Species	Company	Catalog No.	Dilution	Final Conc
IsolectinB4-biotin	NA	ThermoFisher	I21414	1:100	10 µg/ml
Streptavidin-Alexa405	NA	ThermoFisher	S32351	1:50	20 µg/ml
Ki67-Alexa647	Rat	Biolegend	652407	1:20	25 µg/ml
ERG-Alexa647	Rabbit	Abcam	ab196149	1:100	5 µg/ml
DAPI	NA	Sigma	10236276001	1:1,000	10 ng/ml