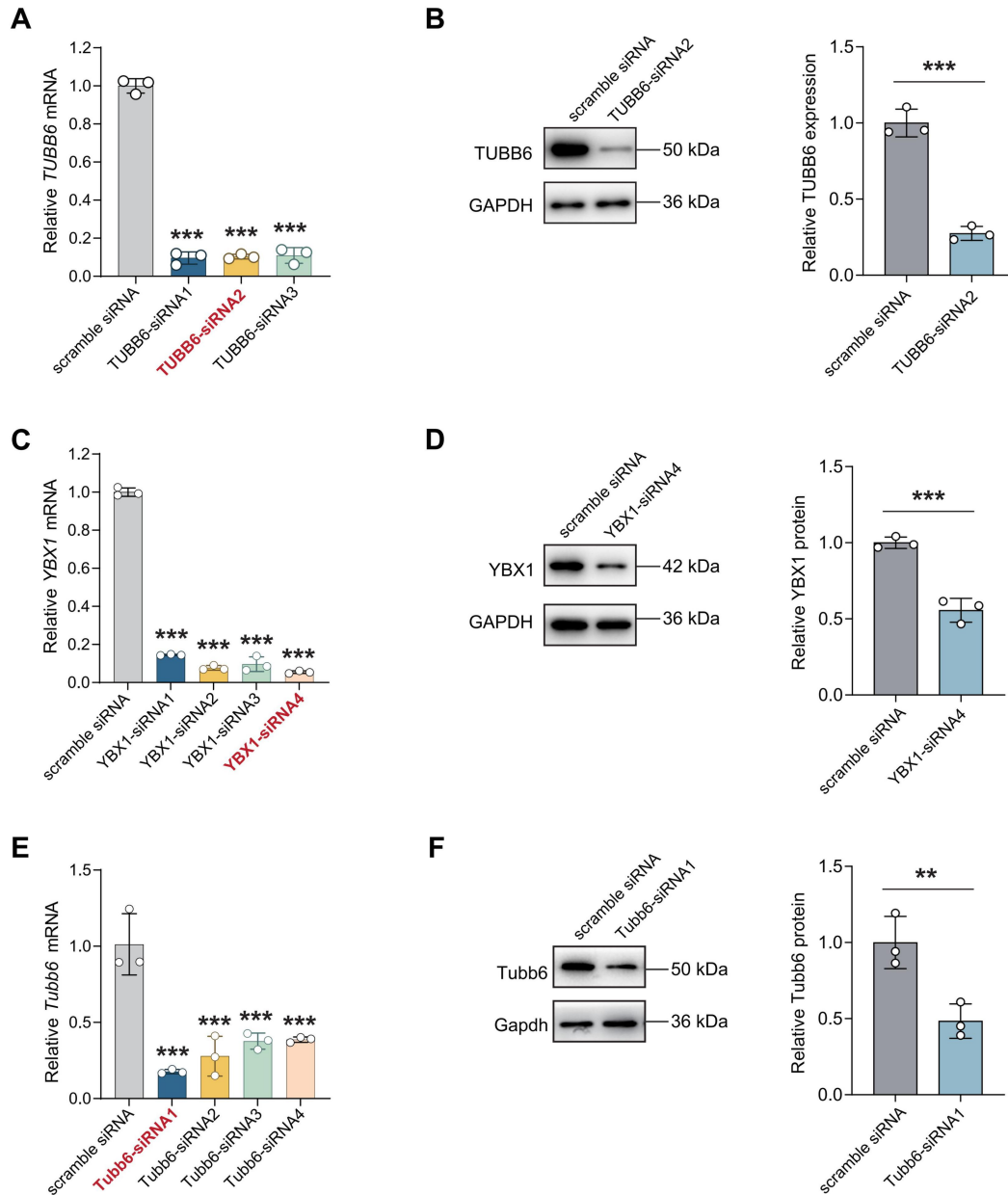


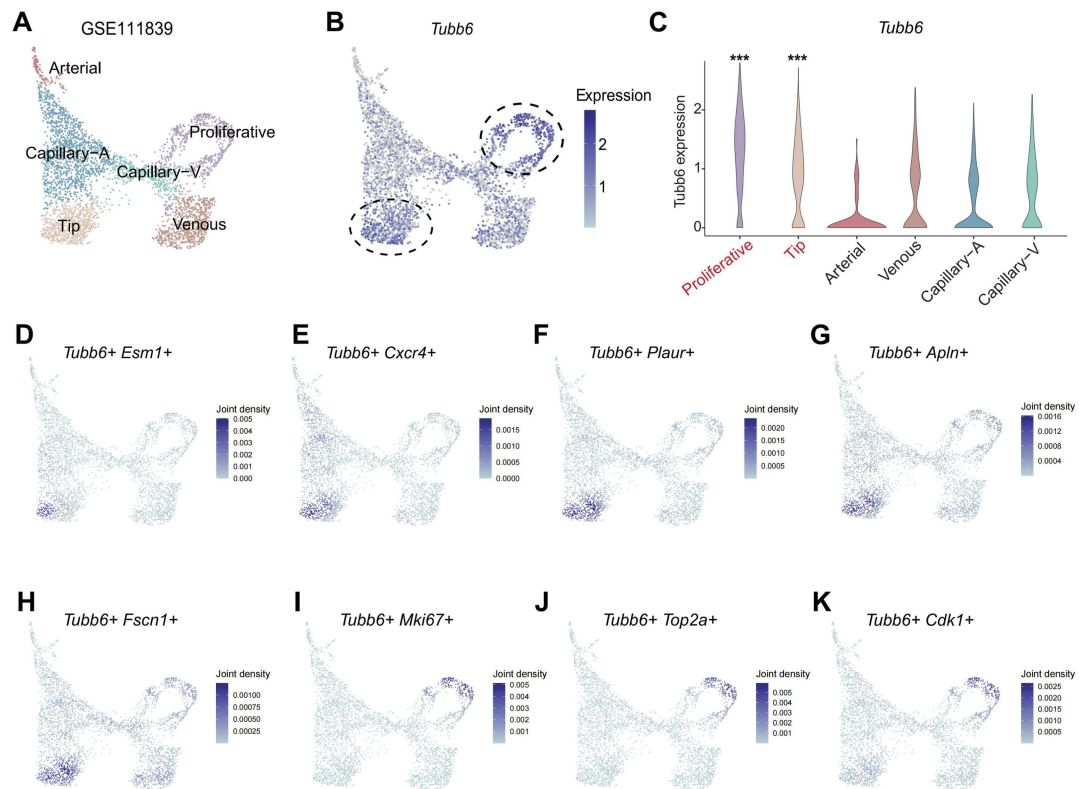
Supplementary Materials

***YBX1*-driven *TUBB6* upregulation facilitates ocular angiogenesis via
WNT3A-FZD8 pathway**

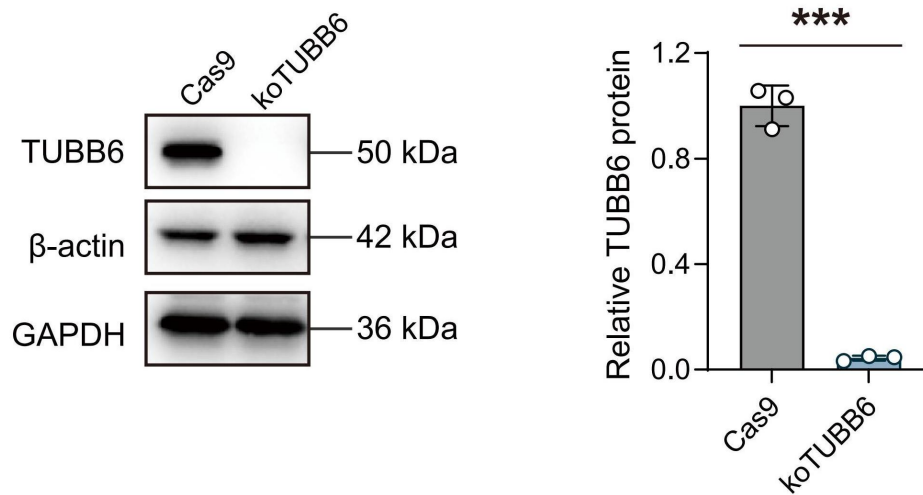
**Supplementary materials include 6 Supplementary Figures and 3
Supplementary Tables.**



Supplemental Figure S1. Knockdown efficiency of TUBB6 and YBX1 *in vitro* and *in vivo*. (A-B) qPCR (A) and immunoblotting (B) confirmed TUBB6 downregulation in TUBB6-siRNA-treated HUVECs. $n = 3$ per group. (C-D) qPCR (C) and immunoblotting (D) confirmed YBX1 downregulation in YBX1-siRNA-treated HUVECs. $n = 3$ per group. (E-F) qPCR (E) and immunoblotting (F) confirmed Tubb6 downregulation in retinas of mice injected with Tubb6-siRNA. $n = 3$ per group. Data represent different numbers (n) of biological replicates. Data are shown as mean \pm SD. Two-tailed Student's T test is used in A-F. ** $p < 0.01$ and *** $p < 0.001$.



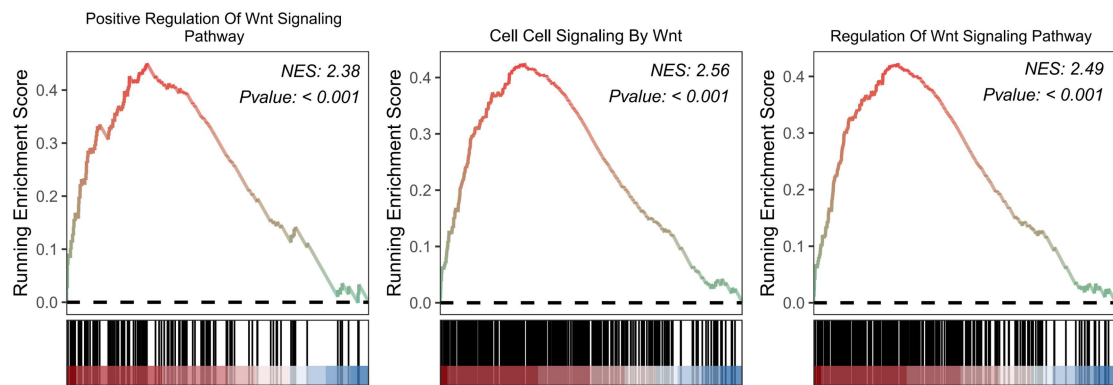
Supplemental Figure S2. *Tubb6* expression pattern in postnatal day 7 (P7) ECs from mice brain. (A) UMAP plot of EC subtypes identified in murine brain (GEO accession: GSE111839). **(B-C)** Feature plot **(B)** and violin plot **(C)** showed that *Tubb6* was highly expressed in proliferative ECs and tip cells compared to other EC subtypes. **(D-H)** *Tubb6* was co-expressed with various tip cell markers in the ECs of murine brain. **(I-K)** *Tubb6* was co-expressed with multiple proliferative markers in the ECs of murine brain. Non-parametric Wilcoxon rank sum test is used in **C**. *** $p < 0.001$.



Supplemental Figure S3. TUBB6-knockout HUVECs constructed with the CRISPR/Cas9 system. Immunoblotting confirmed TUBB6 knockout in HUVECs with Cas9-expressing construct plus the CRISPR/Cas9-TUBB6-KO construct. $n = 3$ per group. Data represent different numbers (n) of biological replicates. Data are shown as mean \pm SD. Two-tailed Student's T test is used. *** $p < 0.001$.

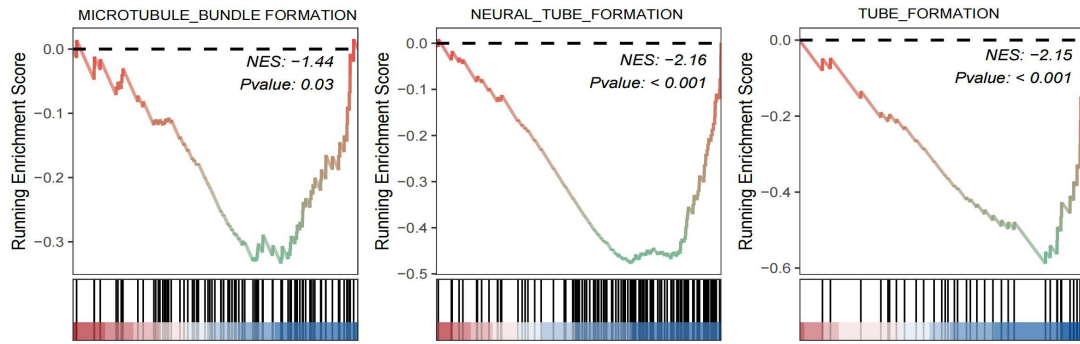
	1	10	20	30	40	50	60	70
Homo_sapiens	MREIVHIIQAGOCGNQIGTKFWEVTSDEHGIDQAGGYVGDSSALQLERISVYYNESSSKKYVPRRAALVDLEP							
Mus_musculus	MREIVHIIQAGOCGNQIGTKFWEVTSDEHGIDQAGGYVGDSSALQLERISVYYNESSSKKYVPRRAALVDLEP							
Rattus_norvegicus	MREIVHIIQAGOCGNQIGTKFWEVTSDEHGIDPAGGYVGDSSALQLERISVYYNESSSKKYVPRRAALVDLEP							
consensus> 70	MREIVHIIQAGOCGNQIGTKFWEVTSDEHGID . AGGYVGDSSALQLERISVYYNESSSKKYVPRRAALVDLEP							
	80	90	100	110	120	130	140	
Homo_sapiens	GTMDSVRSQPFQQLFRPDNFI FGQTGAGNNWAKGHYTEGAELVDSVLDVVRKECEHCDCLOGFQI LTHSLG							
Mus_musculus	GTMDSVRSQPFQQLFRPDNFI FGQTGAGNNWAKGHYTEGAELVDSVLDVVRKECEHCDCLOGFQI LTHSLG							
Rattus_norvegicus	GTMDSVRSQPFQQLFRPDNFI FGQTGAGNNWAKGHYTEGAELVDSVLDVVRKECEHCDCLOGFQI LTHSLG							
consensus> 70	GTMDSVRSQPFQQLFRPDNFI FGQTGAGNNWAKGHYTEGAELVDSVLDVVRKECEHCDCLOGFQI LTHSLG							
	150	160	170	180	190	200	210	
Homo_sapiens	GGTGSQMGCTLLISKIREEYPDRIMNTFSVMPSPKVSQDTVVEPYNATLSVHQLVENTDETYCIDNEALYDI							
Mus_musculus	GGTGSQMGCTLLISKIREEYPDRIMNTFSVMPSPKVSQDTVVEPYNATLSVHQLVENTDETYCIDNEALYDI							
Rattus_norvegicus	GGTGSQMGCTLLISKIREEYPDRIMNTFSVMPSPKVSQDTVVEPYNATLSVHQLVENTDETYCIDNEALYDI							
consensus> 70	GGTGSQMGCTLLISKIREEYPDRIMNTFSVMPSPKVSQDTVVEPYNATLSVHQLVENTDETYCIDNEALYDI							
	220	230	240	250	260	270	280	
Homo_sapiens	CFRTLKLTTPPTYGDLNHLVSATMSGVTTSLRFPQQLNADLRKLA VNMVPPRLHFFMPGFAPLTARGSQQ							
Mus_musculus	CFRTLKLTTPPTYGDLNHLVSATMSGVTTSLRFPQQLNADLRKLA VNMVPPRLHFFMPGFAPLTARGSQQ							
Rattus_norvegicus	CFRTLKLTTPPTYGDLNHLVSATMSGVTTSLRFPQQLNADLRKLA VNMVPPRLHFFMPGFAPLTARGSQQ							
consensus> 70	CFRTLKLTTPPTYGDLNHLVSATMSGVTTSLRFPQQLNADLRKLA VNMVPPRLHFFMPGFAPLTARGSQQ							
	290	300	310	320	330	340	350	
Homo_sapiens	YRALTVPELTQQMFDANKNMAACDPRHGRYLTVA TVFRGPM SMKEVDEQMLAIQNKNSYFVEWIPNNVK							
Mus_musculus	YRALTVPELTQQMFDANKNMAACDPRHGRYLTVA TVFRGPM SMKEVDEQMLAIQNKNSYFVEWIPNNVK							
Rattus_norvegicus	YRALTVPELTQQMFDANKNMAACDPRHGRYLTVA TVFRGPM SMKEVDEQMLAIQNKNSYFVEWIPNNVK							
consensus> 70	YRALTVPELTQQMFDANKNMAACDPRHGRYLTVA TVFRGPM SMKEVDEQMLAIQNKNSYFVEWIPNNVK							
	360	370	380	390	400	410	420	
Homo_sapiens	VAVCDIPPRGLKMASTFIGNSTAIQELFKRISEQFSAMFRRKAF LHWFTGEGMDEMFE TEAESNMNDLVS							
Mus_musculus	VAVCDIPPRGLKMASTFIGNSTAIQELFKRISEQFSAMFRRKAF LHWFTGEGMDEMFE TEAESNMNDLVS							
Rattus_norvegicus	VAVCDIPPRGLKMASTFIGNSTAIQELFKRISEQFSAMFRRKAF LHWFTGEGMDEMFE TEAESNMNDLVS							
consensus> 70	VAVCDIPPRGLKMASTFIGNSTAIQELFKRISEQFSAMFRRKAF LHWFTGEGMDEMFE TEAESNMNDLVS							
	430	440						
Homo_sapiens	EYQQYQDATVNDGEEAFEDDEEEINE							
Mus_musculus	EYQQYQDATVNDGEEAFEDDEEEINE							
Rattus_norvegicus	EYQQYQDATVNDGEEAFEDDEEEINE							
consensus> 70	EYQQYQDATVNDGEEAFEDDEEEINE							

Supplemental Figure S4. Conservation analysis of TUBB6 among distinct species. TUBB6 protein sequence is highly conserved among human, mouse and rats.



Supplemental Figure S5. GSEA analyses of ECs from PDR patients.

GSEA analyses showing activated WNT-related pathways in the CD31-enriched ECs of PDR patients.



Supplemental Figure S6. GSEA analyses of si-TUBB6-treated ECs. GSEA analyses showing disturbed tubulin-related pathways in HUVECs after TUBB6 knockdown.

Supplementary Table S1. Sequences of siRNAs and sgRNA	
Targeted genes	Sequence (5'→3')
<i>In vitro</i>	
scramble siRNA	sense:UUCUCCGAACGUGUCACGUTT
	antisense: ACGUGACACGUUCGGAGAATT
TUBB6-siRNA-1	sense:GUGAGGGCAUGGAUGAAAU
	antisense: CACUCCCGUACCUACUUUA
TUBB6-siRNA-2	sense:UGGAGAGAAUCAACGUCUA
	antisense: ACCUCUCUUAGUUGCAGAU
TUBB6-siRNA-3	sense:UGCUGGCCAUCCAGAGUAA
	antisense: ACGACCGGUAGGUCUCAUU
YBX1-siRNA-1	sense:GGACGGCAAUGAAGAAGAUTT
	antisense: AUCUUCUUCAUUGCCGUCCTT
YBX1-siRNA-2	sense:CCACGCAAUUACCAGCAAATT
	antisense: UUUGCUGGUAAUUGCGUGGTT
YBX1-siRNA-3	sense:CUGUGGAGUUUGAUGUUGUTT
	antisense: ACAACAUCAAACUCCACAGTT
YBX1-siRNA-4	sense:GUCGACCACAGUAUCCAATT
	antisense: UUGGAUACUGUGGUCGACTT
TUBB6-sgRNA	CUGAUCACUCCCAAACUU
<i>In vivo</i>	
scramble siRNA	sense:UUCUCCGAACGUGUCACGUTT
	antisense: ACGUGACACGUUCGGAGAATT
Tubb6-siRNA-1	sense:CAUUGUCAGUGCACCAGCUTT
	antisense: AGCUGGUGCACUGACAAUGTT
Tubb6-siRNA-2	sense:GGGAACCAGAUCCGUACCATT
	antisense: UGGUACCGAUCUGGUUCCCTT
Tubb6-siRNA-3	sense:UUCGGACAGACGGGUGCCGTT
	antisense: CGGCACCCGUCUGUCCGAATT
Tubb6-siRNA-4	sense:CCGGGGCCCAUGUCCAUGTT
	antisense: CAUGGACAUGGGGCCCCGGTT

Supplementary Table S2. Primers used in this study.		
Gene/RNA	Forward primer (5'→3')	Reverse primer (5'→3')
<i>In vitro</i>		
β-ACTIN	ACCTTCTACAATGAGCTGCG	CCTGGATAGCAACGTACATGG
TUBB6	AAGAGGCCCTAAGAACAGCC	AACTTGGTGCCGATCTGGT
ESM1	ACAGCAGTGAGTGCAAAAGCA	GCGGTAGCAAGTTTCTCCCC
PLUAR	TGTAAGACCAACGGGGATTGC	AGCCAGTCCGATAGCTCAGG
CXCR4	ACTACACCGAGGAAATGGGCT	CCACAATGCCAGTTAAGAAGA
APLN	GTCTCCTCCATAGATTGGTCTGC	GGAATCATCCAAACTACAGCCAG
FSCN1	CTGCTACTTTGACATCGAGTGG	GGGCGGTTGATGAGCTTCA
TOP2A	ACCATTGCAGCCTGTAAATGA	GGGCGGAGCAAATATGTTCC
MKI67	GACTTTGGGTGCGACTTGAC	ACAACCTTCCACTGGGACG
CDK1	CCTATGGAGTTGTGTATAAGGGT	AGCACATCCTGAAGACTGACT
YBX1	GCGGGGACAAGAAGGTCATC	CGAAGGTA CTTCCTGGGGTTA
FZD8	ACCCAGCCCCTTTTCCTCCATT	GTCCACCCTCCTCAGCCAAC
WNT3A	CGTGCTGGACAAAGCTACCA	CCAAACTCGATGTCCTCGCT
<i>in vivo</i>		
Gapdh	TGCACCACCAACTGCTTAG	GGATGCAGGGATGATGTTT
Tubb6	CGGCCTGACAACTTCATCTTCG	CCTGAAGACAGTCGCAATGCTC

Supplementary Table S3. Antibodies used in this study.				
Anti-protein	Host	Dilution and Application	Supplier	Catalog
<i>In vitro</i>				
GAPDH	Mouse	1:10000, Immunoblotting	Proteintech	60004-1-Ig
TUBB6	Mouse	1:100, Immunofluorescence; 1:2000, Immunoblotting	Proteintech	66362-1-Ig
YBX1	Rabbit	1:100, Immunofluorescence	Proteintech	20339-1-AP
WNT3A	Rabbit	1:5000, Immunoblotting	Proteintech	26744-1-AP
FRIZZLED 8	Mouse	1:5000, Immunoblotting	Proteintech	55093-1-AP
<i>in vivo</i>				
Gapdh	Mouse	1:10000, Immunoblotting	Proteintech	60004-1-Ig
Tubb6	Rabbit	1:200, Immunofluorescence; 1:2000, Immunoblotting	Merck	ZRB2285
Ve-Cadherin	Rat	1:75, Immunofluorescence	Abcam	ab282277
Ng2	Rabbit	1:100, Immunofluorescence	Merck	AB5320
IB4	/	1:200, Immunofluorescence	Vector Laboratories	FL-1201-.5