

1           **Endothelial epigenetic senescence driven microglial activation mediates**  
2                           **cardio-retinal neuroinflammation in heart failure**

3  
4 Mengdan Wang<sup>1#</sup>, Shuo Zhang<sup>1#,✉</sup>, Tursunjan Aziz<sup>2#</sup>, Shiyao Zhang<sup>1</sup>, Xiao Wu<sup>1</sup>, Gang  
5 Li<sup>1✉</sup>, Yan Wang<sup>1✉</sup>

6 <sup>#</sup>These authors contributed equally to this work.

7 <sup>✉</sup>Corresponding authors.

8  
9 **Author affiliations:**

10 <sup>1</sup>Xiamen Cardiovascular Hospital of Xiamen University, School of Medicine, Xiamen  
11 University, Xiamen, Fujian 361102, China

12 <sup>2</sup>State Key Laboratory of Membrane Biology, Tsinghua-Peking Center for Life Sciences,  
13 School of Life Sciences, Tsinghua University, Beijing 100084, China

14 **Correspondence to:** Shuo Zhang, Gang Li, or Yan Wang

15 ChengYi Building, School of Medicine, Xiamen University, Xiang'an South Road,  
16 Xiang'an District, Xiamen, Fujian, China

17 Email: shuozhang@xmu.edu.cn, ligang@xmu.edu.cn, or wy@medmail.com.cn

18

19 **Abstract**

20 **Rationale:** Heart failure (HF) is increasingly recognized as a systemic disorder that  
21 extends beyond the heart and affects neurovascular tissues, including the retina. However,  
22 the mechanisms by which circulating factors from HF trigger retinal neuroinflammation  
23 remain unclear.

24 **Methods:** HF was induced in adult mice by transverse aortic constriction (TAC). Retinal  
25 structure and function were evaluated using optical coherence tomography (OCT) and  
26 electroretinography (ERG). Parabiosis and plasma transfer experiments were performed  
27 to assess the role of circulating factors. Endothelial senescence, microglial activation, and  
28 inflammatory signaling were analyzed using immunofluorescence, qPCR, and molecular  
29 assays. The functional relevance of TGF $\beta$ 2 and microglia was tested using anti-TGF $\beta$ 2  
30 antibody administration and microglial depletion with PLX5622 treatment.

31 **Results:** TAC mice exhibited pronounced retinal thinning, diminished  
32 electroretinography (ERG) amplitudes, and reduced vascular density. Exposure of healthy  
33 mice to HF plasma reproduced these abnormalities, indicating that circulating mediators  
34 drive retinal injuries. TGF $\beta$ 2 levels were markedly elevated in the plasma of both patients  
35 with HF and TAC mice. Mechanistically, TGF $\beta$ 2 activated the pSMAD2/EP300 pathway  
36 in retinal endothelial cells, promoting H3K9 acetylation, P21 induction, and endothelial  
37 cell senescence. Senescent endothelial cells release proinflammatory factors that activate  
38 retinal microglia, leading to hypertrophic morphology, enhanced synaptic phagocytosis,

39 and upregulation of cytokines such as IL1 $\beta$ , TNF $\alpha$ , and IL6. Neutralization of TGF $\beta$ 2 or  
40 microglial depletion markedly reduced inflammation, preserved the retinal architecture,  
41 and restored visual function.

42 **Conclusions:** Elevated TGF $\beta$ 2 levels in heart failure drive retinal endothelial epigenetic  
43 senescence, which secondarily activates microglia and induces neuroinflammation.  
44 Endothelial-specific disruption of TGF $\beta$ 2 signaling is sufficient to protect the retina  
45 independently of primary cardiac recovery. Targeting the TGF $\beta$ 2–endothelial–microglia  
46 axis may represent a promising therapeutic strategy for preventing retinal neurovascular  
47 degeneration associated with systemic cardiac disease.

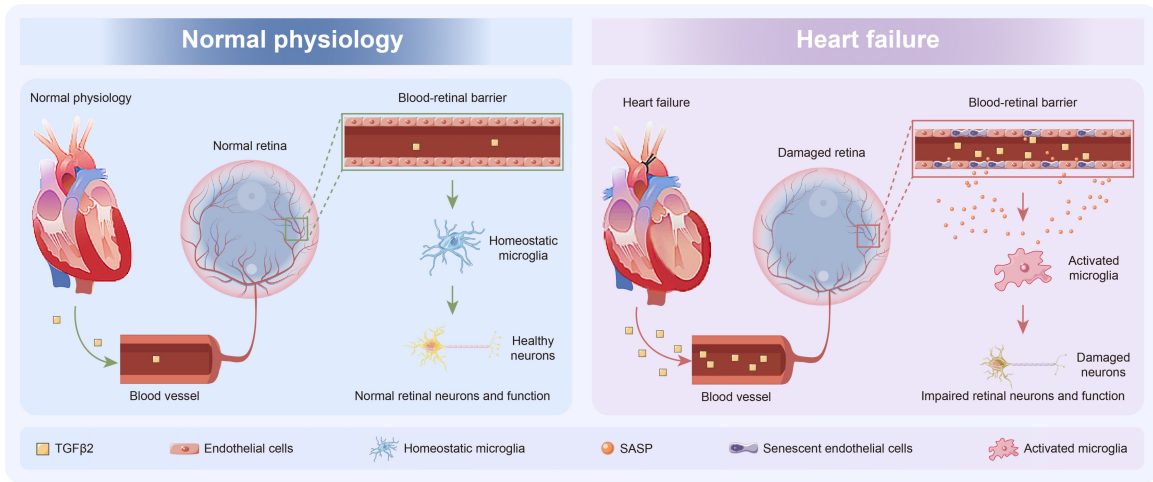
48

49 **Key Words:**

50 Heart failure; Microglia activation; Neuroinflammation; Epigenetic senescence; TGF $\beta$ 2

51

52 **Graphical Abstract**



53

54

55 **Introduction**

56 Heart failure (HF) is a major healthcare burden equally affecting an estimated 26 million  
57 people globally with an exponentially rising trend [1]. Even with emerging therapies, the  
58 five-year survival of HF patients is still <50% [2-6]. Heart failure is not a heart disease,  
59 but rather a disease of varying severity and severity of function/structure of multiple  
60 organs and tissues [7]. The cellular stress owing to inflammation and accelerated  
61 senescence leads to increased dysfunctional protein production, which enter the  
62 circulation and impact distant organs [8-10]. Although such findings highlight that HF  
63 has a systemic effect, the mechanisms of how circulating factors affect the remote organs,  
64 especially with respect to protein misfolding, are poorly understood.

65 Recent studies have demonstrated that HF affects respiratory and cardiovascular  
66 systems. It has also been shown to impair the central nervous system with functions  
67 [11-13]. The retina is part of the central nervous system. Its embryological origin and  
68 vascular supply are similar to the brain. The proper functioning of the immune system  
69 depends on the careful regulation of its internal environment [14]. The retina's density,  
70 vascular structure, and high metabolic demand make it susceptible to insult from any  
71 cardiovascular damage. Retinal microvascular changes are considered the first visible  
72 signs of systemic cardiovascular diseases [15, 16]. Past studies have shown that HF and  
73 atherosclerosis or ischemic heart disease harm retinal vasculature [15, 17-21]. Chronic  
74 inflammatory response and oxidative damage worsen blood-retinal barrier breakdown

75 and further aggravate vascular pathology [15, 17-21]. Even with these findings, the  
76 mechanisms which HF interacts with the retinal vasculature system and function is  
77 unclear.

78 In this study, we showed that HF severely damages the retinal and vascular networks  
79 in mice. It is worth noting that plasma from HF mice can do similar harm to healthy mice.  
80 Therefore, these findings suggest that blood-borne circulating factors are important for  
81 HF retinal injury. Further investigation revealed a significantly elevated level of TGF $\beta$ 2  
82 in the blood of heart failure patients and heart failure mice. TGF $\beta$ 2 is known for its  
83 important functions in regulating cell proliferation, differentiation and apoptosis [22, 23].  
84 Within the cardiovascular system, the dysregulated expression of TGF $\beta$ 2 is tightly  
85 associated with the main pathophysiological features of HF, including cardiac  
86 remodelling, fibrosis and inflammation [23, 24]. Results showed that TGF $\beta$ 2 activated  
87 the phosphorylated SMAD2 (pSMAD2)/EP300 pathway and enhanced the acetylation of  
88 histone H3 on lysine 9 (H3K9ac) to activate chromatin remodeling and upregulation of  
89 P21. The epigenetic senescence of endothelial cells is promoted by this cascade.  
90 Afterward, senescent endothelial cells release senescence-associated secretory phenotype  
91 (SASP) factors. This activates retinal microglia, increases their phagocytic activity,  
92 amplifies neuroinflammation. Consequently, retinal neurons get damaged in the process.  
93 Significantly, depletion of microglia interrupts this pathogenic mechanism, preventing  
94 damage to retinal structure and function caused by HF.

95 To probe the therapeutic utility of inhibiting TGF $\beta$ 2, HF mice were treated with  
96 anti-TGF $\beta$ 2 antibody by intravitreal injection. The findings displayed a decrease in the  
97 aging of endothelial cells, as well as improvement in functionality and structure of the  
98 retina. Apart from affecting the retina, our research has indicated that HF can lead to  
99 endothelial senescence in various organs, including the spleen, lungs, kidneys, and even  
100 the heart. The anti-TGF $\beta$ 2 antibody given intravenously was similarly effective in  
101 reducing endothelial cell senescence in these organs. The antibody significantly reduced  
102 cell senescence induced by plasma from HF patients, which was verified on primary  
103 human retinal microvascular endothelial cells.

104 In conclusion, these findings offer new insights into the mechanisms driving  
105 HF-induced damage to the retina and other organs. They further provide a theoretical  
106 basis for considering anti-TGF $\beta$ 2 antibodies as a potential therapeutic approach to  
107 address multi-organ damage in the context of heart failure.

108 **Materials and Methods**

109 **Animals**

110 C57BL/6J mice (stock no. 000664) were from Jackson Laboratory. B-NDG mice were  
111 from Biocytogen (Beijing). *Cdh5*-CreERT2 mice (C001330) were from Cyagen  
112 (Guangzhou) and *Tgfbr2*-flox mice (NM-CKO-200026) from Shanghai Southern Model  
113 Biotechnology. Endothelial-specific *Tgfbr2* conditional knockout (cKO) mice were  
114 generated by crossing *Tgfbr2*-flox with *Cdh5*-CreERT2 mice, followed by tamoxifen  
115 induction to activate Cre recombinase before TAC surgery. All mice were on a C57BL/6J  
116 background, randomly assigned to groups, and housed under a 12-h light/dark cycle at 22  
117  $\pm 2$  °C with ad libitum access to food and water. All animal procedures were approved by  
118 the IACUC of Xiamen University and complied with ARVO guidelines.

119 **Culture of human endothelial cells**

120 Human umbilical vein endothelial cells (HUVECs; Yizefeng Biotechnology, YPC-H001)  
121 and human primary retinal microvascular endothelial cells (HRMECs; Meisen Cell  
122 Technology, CTCC-150-HUM) were cultured in endothelial cell medium (ECM;  
123 ScienCell, 1001) supplemented with fetal bovine serum and growth factors at 37 °C in  
124 5% CO<sub>2</sub>. HRMEC medium was additionally supplemented with 100 U/mL penicillin and  
125 100  $\mu$ g/mL streptomycin. Cells were detached with accutase (Invitrogen, 00-4555-56).  
126 For treatments, cells were seeded at  $2 \times 10^5$  cells per well in 24-well plates, cultured  
127 overnight for attachment, and then exposed to the indicated treatments for 48 h.

128 **Collection and processing of blood samples**

129 Human peripheral blood was collected from 20 HF patients and 20 healthy controls at the  
130 Cardiovascular Hospital of Xiamen University with informed consent and ethics approval  
131 (No. XMH202449). HF diagnosis followed ACC/AHA criteria. Mouse blood was  
132 collected from the retroorbital sinus under isoflurane anesthesia into EDTA-containing  
133 tubes (Solarbio, YA1461). Human and mouse blood samples were centrifuged at 1,500 g  
134 for 10 min at 4 °C, and plasma was stored at -80 °C. Before use, plasma was dialyzed in  
135 saline using a 3.5 kDa D-tube dialyzer (EMD Millipore) at 4 °C with saline changes  
136 every 4 h over 48 h, sterilized by filtration through a 0.22 µm filter, and stored at -80 °C.

137 **Plasma treatment**

138 HUVECs and HRMECs were seeded and allowed to attach for 24 h. Dialyzed human or  
139 mouse plasma was added to the culture medium at 1% (v/v) for 48 h, after which cells  
140 were harvested for senescence assays. For EV depletion experiments, plasma was first  
141 subjected to ultracentrifugation as described below.

142 **Extracellular vesicle depletion**

143 To determine whether the senescence-inducing activity of HF plasma is associated with  
144 extracellular vesicles (EVs), plasma samples from normal/HF patients and Sham/TAC  
145 mice were subjected to EV depletion by differential ultracentrifugation. Briefly, plasma  
146 was centrifuged at 2,000 g for 20 min and then at 10,000 g for 30 min to remove residual  
147 debris and large vesicles. The resulting supernatant was subsequently ultracentrifuged at

148 100,000 g for 90 min at 4°C to pellet small EVs. The EV-depleted supernatant was  
149 collected, diluted 1:100 in culture medium, and applied to endothelial cells for 48 h.  
150 Following treatment, cells were harvested for senescence-related assays.

### 151 **Establishment of the TAC model and cardiac function monitoring**

152 Transverse aortic constriction was performed in 2-month-old C57BL/6J mice as  
153 previously described [25] to induce pressure-overload heart failure. Briefly, under  
154 isoflurane anesthesia, the aortic arch was exposed via thoracotomy and constricted  
155 between the innominate and left common carotid arteries using a 27-gauge needle and 6-0  
156 silk suture. The needle was removed immediately after ligation. Sham-operated mice  
157 underwent thoracotomy without aortic constriction. All mice received postoperative  
158 antibiotics and analgesics. Cardiac function was monitored by echocardiography (Echo)  
159 at baseline and monthly thereafter, as described [25]. LVEF%, LVFS%, LVID,d, LVID,s,  
160 LVEDV, and LVESV were measured.

### 161 **Retinal electroretinogram analysis**

162 Retinal function was assessed by electroretinography (ERG) as previously described [14].  
163 Briefly, mice were dark-adapted for  $\geq 12$  h, anesthetized with isoflurane, and pupil-dilated  
164 with tropicamide. Scotopic ERG responses were recorded at stimulus intensities of  
165 0.01–10 cd·s/m<sup>2</sup> using gold-ring corneal electrodes. After 30 min of light adaptation (30  
166 cd/m<sup>2</sup> background), photopic responses were recorded at 3–100 cd·s/m<sup>2</sup>. Fifteen  
167 waveforms per animal were averaged. a-wave amplitude was measured from baseline to

168 trough; b-wave amplitude was measured from the a-wave trough to the positive peak, or  
169 from baseline if no a-wave was present.

#### 170 **Optical coherence tomography**

171 Retinal structure was assessed by optical coherence tomography (4D-ISOCT;  
172 OPTOPROBE, UK). Retinal volumes were acquired from 200 vertical B-scans per eye  
173 (1,000 A-scans per B-scan). Retinal layer thickness was quantified using FIJI (NIH).  
174 Layers analyzed included GCL+IPL, IS+OS, and total retinal thickness.

#### 175 **Parabiosis**

176 Parabiosis was performed according to a previously established protocol [26]. Briefly,  
177 2-month-old WT mice were paired with age-matched Sham or TAC mice (surgery at 1.5  
178 months, 2-week recovery). Under sodium pentobarbital anesthesia, skin incisions were  
179 made along the flanks, and partners were sutured at the elbow, knee, and dorsal-ventral  
180 skin with 4-0 silk.

#### 181 **Primary microglia culture and synaptosome phagocytosis**

182 Primary mouse microglia were isolated from mixed glial cultures prepared from P1–P3  
183 mouse pups. Cultures were maintained in DMEM with 10% FBS, and microglial  
184 proliferation was promoted with GM-CSF (25 ng/mL; R&D Systems, 415-ML-050)  
185 added at day 3. Mature microglia were harvested by shaking (200 rpm, 30 min) at days  
186 10–12. Synaptosomes were purified from mouse hippocampi and labeled with pHrodo  
187 Red (Thermo Fisher, P36600) as previously described [27]. For the phagocytosis assay,

188 microglia were seeded at  $2 \times 10^5$  cells per well, cultured for 24 h, treated as indicated,  
189 and incubated with pHrodo-labeled synaptosomes (1.2  $\mu\text{g}/\mu\text{L}$ ) for the final 3 h before  
190 immunocytochemical analysis.

### 191 **Histology and staining**

192 Lectin staining. Enuclated eyes were fixed in 4% PFA for 1–2 h, and retinas were  
193 dissected, permeabilized in 0.2% Triton X-100 overnight at 4 °C, and incubated with  
194 fluorescent lectin (Thermo Fisher, I21413) overnight at 4 °C. Retinas were flat-mounted  
195 in four-leaf clover cuts and imaged on a Leica DMI4000B microscope. Sirius red staining.  
196 Frozen heart sections were fixed in 4% PFA, stained with 0.2% phosphomolybdic acid  
197 (2.5 min) followed by Picro-Sirius Red (Abcam, ab150681; 90 min), differentiated in  
198 0.5% acetic acid, dehydrated, and coverslipped. Wheat germ agglutinin (WGA) staining.  
199 Frozen tissue sections were fixed in cold acetone (–20 °C, 10 min), incubated with 5  
200  $\mu\text{g}/\text{mL}$  WGA (Sigma, L4895) and DAPI (1  $\mu\text{g}/\text{mL}$ ; CST, 4083) for 30 min, and imaged  
201 on a Leica SP8 DLS confocal microscope.

### 202 **RNA extraction and qPCR analysis**

203 Total RNA was extracted with TRIzol (Thermo Fisher, 15596026CN) and  
204 reverse-transcribed. qRT-PCR was performed with FastStart Universal SYBR Green  
205 Master (Roche, 04913850001) on a LightCycler 480II system (Roche). Target gene  
206 expression was calculated by the  $\Delta\Delta\text{Ct}$  method with  $\beta$ -actin as the internal control.

207 **RNA sequencing**

208 RNA integrity was assessed on an Agilent 2100 Bioanalyzer. Libraries were prepared  
209 with the NEBNext Ultra RNA Library Prep Kit (NEB, E7530) and sequenced on an  
210 Illumina NovaSeq 6000 (150-bp paired-end). Raw reads were quality-checked with  
211 FastQC (v0.11.5), adapter-trimmed with Trim Galore (v3.4), and mapped to the human  
212 transcriptome (hg38, GENCODE v41) with STAR (v2.7.10b). Gene counts were  
213 obtained with featureCounts (v2.0.6) and differential expression was analyzed with  
214 DESeq2 (v1.42.0).

215 **Single-cell sequencing data and RNA-seq data analysis**

216 Publicly available single-nucleus RNA-seq data (EGA: EGAS00001006374) and bulk  
217 RNA-seq data (GEO: GSE116250) from HF patients and healthy controls were analyzed.  
218 Single-nucleus data were processed with Seurat (v4.0): dimensionality reduction was  
219 performed by UMAP, differential expression was analyzed with DESeq2, and senescence  
220 scores were calculated with AUCell. GO enrichment was performed with clusterProfiler  
221 (v4.10.0), and heatmaps were generated with ComplexHeatmap.

222 **CUT&Tag**

223 CUT&Tag was performed with the Hyperactive In-Situ ChIP Library Prep Kit (TD901,  
224 Vazyme). Paired-end reads were aligned with Bowtie2 (v2.5.1; -end-to-end  
225 -very-sensitive -no-mixed -no-discordant -I 10 -X 700). GO and motif enrichment  
226 analyses were performed with clusterProfiler.

227 **Co-immunoprecipitation**

228 Endothelial cells were lysed in TNE buffer (20 mM Tris-HCl, pH 7.4; 100 mM NaCl; 1  
229 mM EDTA; 0.5% NP-40) with protease inhibitors (Roche, 4693159001). Lysates were  
230 incubated with anti-SMAD2/3 (CST, 8685; 1:200) or normal rabbit IgG (CST, 2729)  
231 overnight at 4 °C, followed by incubation with Dynabeads Protein G (Thermo Fisher,  
232 10004D) for 4 h. Bound proteins were eluted, resolved by SDS-PAGE, and  
233 immunoblotted with anti-EP300 (Santa Cruz, sc-48343; 1:1000).

234 **Chromatin immunoprecipitation**

235 ChIP was performed as previously described [28] with modifications. Endothelial cells  
236 were cross-linked with 1% formaldehyde, quenched with 125 mM glycine, lysed, and  
237 sonicated to 150–350 bp fragments. Chromatin was incubated overnight at 4 °C with  
238 antibodies against SMAD2/3 (CST, 8685), EP300 (CST, 57625), or H3K9ac (CST, 9649),  
239 followed by incubation with Dynabeads Protein G for 4 h. After sequential washes and  
240 elution, DNA was purified and analyzed by qPCR on a LightCycler 480 system.

241 **RNA interference**

242 siRNA transfections were performed with Lipofectamine RNAiMAX (Thermo Fisher,  
243 13778100) according to the manufacturer's protocol. siRNAs targeting human SMAD2  
244 (siSmad2-2) and EP300 (siEp300-3) and a non-targeting control (siN0000001-1-5) were  
245 from RiboBio. The siRNA sequences targeting human Smad2 and Ep300 were as  
246 follows:

247 si*Smad2-2*, sense: 5'-GUCCCAUGAAAAGACUAAA-3',  
248 si*Smad2-2*, antisense: 5'-UUAAGUCUUUUCAUGGGAC-3';  
249 si*Ep300-3*, sense: 5'-CAAUAGAGCGGAAUACUAU-3',  
250 si*Ep300-3*, antisense: 5'-AUAGUAUCCGCUCUAUUG-3'.

### 251 **Tail vein injections of TGF $\beta$ 2**

252 Recombinant TGF $\beta$ 2 (6 ng per mouse in PBS) or PBS was injected via the tail vein every  
253 2 days for 2 months, starting at 2 months of age. The dose was calculated to approximate  
254 the elevated plasma TGF $\beta$ 2 concentration observed in HF patients (~6 ng/mL vs. ~1  
255 ng/mL in healthy controls), assuming a total blood volume of ~2 mL per mouse. Mice  
256 were assessed 3 days after the final injection.

### 257 **Anti-TGF $\beta$ 2 antibody administration**

258 Intravitreal injection. TAC or Sham mice received six intravitreal injections of  
259 anti-TGF $\beta$ 2 antibody (Invitrogen, MA5-37505; 1 mg/kg in 1  $\mu$ L saline) or isotype control  
260 IgG (Abclonal, AC011) between 3 and 5 months of age. Injections were performed under  
261 isoflurane anesthesia using a 32 G needle under microscopic guidance. Mice were  
262 assessed at 5 months after a 1-week recovery. Tail vein injection. The same antibody and  
263 dosing schedule (1 mg/kg, six injections over 2 months) were administered via the tail  
264 vein for multi-organ senescence experiments. Cell treatment. Endothelial cells at 70–80%  
265 confluence were treated with anti-TGF $\beta$ 2 antibody (1  $\mu$ g/mL) or vehicle for 48 h before  
266 downstream analysis.

267 **ELISA**

268 Plasma TGF $\beta$ 2 was measured with a human TGF $\beta$ 2 ELISA kit (Elabscience,  
269 E-EL-H1587) following the manufacturer's instructions. Plasma samples were  
270 acid-activated before assay to measure total TGF $\beta$ 2. Absorbance was read at 450 nm, and  
271 TGF $\beta$ 2 concentrations were calculated from a standard curve. Each sample was assayed  
272 in triplicate.

273 **Immunohistochemistry and Immunocytochemistry**

274 Immunohistochemistry (IHC). Tissue sections were prepared after perfusion fixation and  
275 stained as previously described [29]. Primary antibodies: anti-Iba1 (Wako, 019-19741;  
276 1:500), anti-CD31 (CST, 3528S; 1:100), and anti- $\beta$ -galactosidase (CST, 2372S; 1:100).  
277 Confocal imaging was performed on a Leica SP8 DLS, and quantification used ImageJ  
278 (NIH). Immunocytochemistry (ICC). Cells were fixed in 4% PFA (30 min),  
279 permeabilized with 0.2% Triton X-100 (5 min), and blocked with 5% BSA (30 min).  
280 Primary antibodies (overnight, 4 °C): anti-Iba1 (Wako, 019-19741; 1:500; or SYSY,  
281 234308; 1:500), anti-CD31 (CST, 3528S; 1:100), anti-p16INK4a (CST, 18769; 1:200),  
282 anti- $\beta$ -galactosidase (CST, 2372S; 1:200), anti-p21 (CST, 2947S; 1:500), and  
283 anti-H3K9ac (CST, 5327S; 1:200). Secondary antibodies and DAPI were applied for 30  
284 min at room temperature. Imaging and analysis used Leica SP8 DLS, ImageJ, and Imaris  
285 (Bitplane).

286 **Immunoblot analysis**

287 Immunoblotting analysis was performed according to previously established protocols  
288 [30]. Protein extracts were resolved by SDS-PAGE and transferred to PVDF membranes.  
289 Membranes were blocked with 5% non-fat dry milk (1 h, room temperature) and  
290 incubated overnight at 4 °C with primary antibodies: TGFβ2 (Proteintech, 19999-1-AP;  
291 1:1000; or Invitrogen, MA5-37505; 1:1000), p-SMAD2 (CST, 3108; 1:1000), SMAD2  
292 (CST, 3103; 1:1000), SMAD2/3 (CST, 8685; 1:1000), p-SMAD1/5/9 (CST, 13820;  
293 1:1000), SMAD5 (CST, 12534; 1:1000), EP300 (Invitrogen, MA1-16608; 1:1000; or  
294 Santa Cruz, sc-48343; 1:1000), Histone H3 (CST, 14269; 1:1000), H3K9ac (CST, 9649;  
295 1:1000), p21 (CST, 2947; 1:1000), and GAPDH (CST, 5174; 1:1000; or Rockland,  
296 200-901-BJ5; 1:1000). HRP-conjugated secondary antibodies were applied for 1 h, and  
297 signals were detected by ECL (Meilunbio, MA0186). For mouse plasma, albumin was  
298 depleted with the Minute Albumin Depletion Kit (Invent Biotechnologies, WA-013).  
299 Bands were quantified with ImageJ and normalized to GAPDH.

300 **Flow Cytometric Sorting**

301 Retinas were dissected and digested in HBSS containing 1.5 mg/mL collagenase II and  
302 100 U/mL DNase I (37 °C, 15–20 min). For endothelial cell isolation, single-cell  
303 suspensions were stained with anti-CD31-APC and anti-CD45-FITC (1:200) and DAPI.  
304 Live CD31<sup>+</sup>CD45<sup>-</sup> cells were sorted as retinal vascular endothelial cells. For microglia  
305 isolation, retinas were digested with 1 mg/mL collagenase D and 100 U/mL DNase I,

306 stained with anti-CD11b-APC and anti-CD45-PE, and live CD11b<sup>+</sup>CD45<sup>int</sup> cells were  
307 sorted as retinal microglia. Sorting was performed on a flow cytometer.

### 308 **Statistical analysis**

309 All experiments were independently repeated at least thrice. Immunofluorescence data  
310 were analyzed using ImageJ software, whereas other experimental data were analyzed  
311 using GraphPad Prism software (version 8.0). Data are presented as mean  $\pm$  standard  
312 error of the mean (SEM). Differences between the two groups were analyzed using a  
313 two-tailed Student's *t*-test. One-way or two-way analysis of variance (ANOVA) was used  
314 for comparisons involving more than two groups. \**P* < 0.05; \*\**P* < 0.01; \*\*\**P* < 0.001;  
315 \*\*\*\**P* < 0.0001 were considered statistically significant.

316 **Results**

317 **Heart failure induces retinal and vascular network damage**

318 To assess the impact of HF on retina, we subjected 2-month-old mice to transverse aortic  
319 constriction (TAC) surgery and monitored their cardiac function monthly. At 5 months,  
320 retinal structure and function were evaluated using optical coherence tomography (OCT)  
321 and electroretinography (ERG) (Figure 1A). Compared to the sham-operated (Sham)  
322 group, the TAC group exhibited pronounced myocardial hypertrophy and fibrosis  
323 (Figures 1B-C). Echocardiographic analysis further demonstrated progressive left  
324 ventricular dysfunction, as evidenced by a decline in the left ventricular ejection fraction  
325 (LVEF%) and fractional shortening (LVFS%) (Figures 1D-G), as well as increases in  
326 HW/TL, the left ventricular internal diameter during diastole (LVID,d), systole (LVID,s),  
327 left ventricular end-diastolic volume (LVEDV) and end-systolic volume (LVESV)  
328 (Figures 1H-K). These findings confirmed the successful establishment of a HF model.

329 The ERG tests were conducted under both scotopic (dark-adapted) and photopic  
330 (light-adapted) conditions to evaluate retinal function. Under scotopic conditions, TAC  
331 mice showed reduced a-wave and b-wave amplitudes at light intensities of 0.01, 3, and 10  
332 cd·s/m<sup>2</sup>, indicating impaired rod photoreceptor function and bipolar cell signaling  
333 (Figures 1L-M). Similarly, under photopic conditions, TAC mice exhibited decreases in  
334 b-wave amplitudes at 3, 10, 30, and 100 cd·s/m<sup>2</sup>, reflecting compromised cone  
335 photoreceptor and bipolar cell signaling function (Figures 1N-O).

336 OCT scans revealed a reduction in total retinal thickness in the TAC group, with  
337 notable thinning observed in the ganglion cell layer and inner plexiform layer (GCL+IPL),  
338 as well as in the inner and outer photoreceptor segments (IS+OS) (Figures 1P-Q).  
339 Furthermore, lectin staining highlighted a decrease in retinal vascular density and vessel  
340 diameter in TAC mice (Figures 1R-S), suggesting that heart failure not only disrupts  
341 retinal structure and function but also induces damage to the retinal vascular network.

#### 342 **Heart failure plasma disrupts the retinal and vascular network**

343 To explore the involvement of blood-borne factors in HF-induced retinal and vascular  
344 damage, we conducted a parabiosis experiment, allowing blood exchange between the  
345 two mice. This approach enabled us to evaluate the effect of HF plasma on the retinal  
346 structure and function of healthy wild-type (WT) mice. In this experiment, 2-month-old  
347 WT mice were paired with either the Sham or TAC mice. Retinal structure and function  
348 were assessed at 5 months of age using OCT and ERG (Figure 2A).

349 Our findings showed that exposure to HF plasma adversely affected retinal function  
350 in WT mice. Under dark-adapted conditions, the WT (TAC) group exhibited reduced  
351 a-wave and b-wave amplitudes, indicating impaired signal transduction function of rod  
352 photoreceptors and bipolar cells (Figures 2B-C). Similarly, under light-adapted  
353 conditions, the WT (TAC) group displayed lower b-wave amplitudes across different  
354 light intensities, reflecting diminished cone photoreceptor and bipolar cell activity  
355 (Figures 2D-E). Additionally, OCT scans showed a reduction in retinal thickness in the

356 WT (TAC) group, with noticeable thinning in the GCL+IPL and IS+OS (Figures 2F-G).  
357 Lectin staining further revealed that HF plasma decreased retinal vascular density and  
358 reduced vessel diameter (Figures 2H-I). These observations indicate that components of  
359 HF plasma disrupt both the retinal structure and its vascular network through systemic  
360 circulation.

361 In summary, HF exerts systemic effects that extend beyond the heart and release  
362 circulating factors that contribute to retinal and vascular damage. These findings  
363 underscore the potential of HF to influence distant organs such as the retina via  
364 blood-mediated mechanisms.

### 365 **Heart failure plasma induces endothelial cell senescence**

366 Following the confirmation that factors present in HF plasma contribute to retinal damage  
367 in mice, we sought to investigate the underlying molecular mechanisms. An analysis of  
368 single-nucleus RNA sequencing data (accession number: EGAS00001006374) from HF  
369 patient hearts revealed distinct differences in cell distribution compared with healthy  
370 controls (Figure 3A, Figure S1A-B). GO enrichment analysis indicated that pathways  
371 related to immune responses, inflammation, and cellular senescence were upregulated in  
372 patients with HF, while pathways involved in metabolic processes and vascular  
373 development were downregulated (Figure 3B). Further analysis using UMAP projections  
374 showed an elevated senescence score in endothelial cells from HF patients (Figures

375 3C-D). This suggests that endothelial cell senescence may impair vascular function and  
376 contribute to the progression of HF, with potential implications for distant organ systems.

377 We further assessed senescence-associated  $\beta$ -galactosidase (SA- $\beta$ -gal) expression in  
378 retinal endothelial cells from the mice of the Sham and TAC groups. SA- $\beta$ -gal levels  
379 were higher in the TAC group, indicating that HF promoted endothelial cell senescence  
380 in tissues beyond the heart (Figures 3E-F). To explore this further, human umbilical vein  
381 endothelial cells (HUVECs) were exposed to plasma from HF patients and TAC mice  
382 (Figure 3G). SA- $\beta$ -gal expression increased in endothelial cells treated with HF patient  
383 plasma, while no such effect was observed in normal plasma treated cells (Figures 3H-I).  
384 Likewise, endothelial cells treated with plasma from TAC mice displayed elevated  
385 SA- $\beta$ -gal levels compared to cells exposed to Sham plasma (Figures 3J-K), supporting  
386 the conclusion that HF plasma contains factors that induce endothelial cell senescence.

387 RNA sequencing of endothelial cells treated with TAC plasma revealed widespread  
388 alterations in gene expression (Figure 3L). GO enrichment network analysis identified  
389 upregulation in pathways related to tumor necrosis factor (TNF) signaling, chemokine  
390 signaling, innate immune responses, and inflammation, whereas pathways associated  
391 with the cell cycle and DNA replication were downregulated (Figure 3M). These findings  
392 were corroborated by Gene Set Enrichment Analysis (GSEA), which highlighted the  
393 enrichment of these inflammatory and senescence-related pathways in endothelial cells  
394 exposed to TAC plasma (Figure 3N). In conclusion, these data indicate that circulating

395 factors in the HF plasma contribute to retinal and vascular damage by inducing  
396 endothelial cell senescence. Senescent endothelial cells activate immune and  
397 inflammatory pathways, reinforcing the SASP.

398 To further investigate whether the senescence-inducing activity of HF plasma is  
399 mediated by free soluble factors or extracellular vesicles (EVs), we depleted EVs from  
400 both normal/HF patient plasma and Sham/TAC mouse plasma by ultracentrifugation  
401 (100,000g, 90 min, 4°C) and treated endothelial cells with the EV-depleted supernatant  
402 for 48 h. EV-depleted HF plasma still significantly increased SA- $\beta$ -gal expression in  
403 endothelial cells compared to EV-depleted normal plasma (Figure S1C-E). Similarly,  
404 EV-depleted TAC plasma induced higher SA- $\beta$ -gal levels than EV-depleted Sham plasma  
405 (Figure S1F-G). These results indicate that the senescence-inducing factor, TGF $\beta$ 2, exists  
406 predominantly as a free soluble protein in HF plasma rather than being encapsulated  
407 within extracellular vesicles.

#### 408 **TGF $\beta$ 2 mediates endothelial cell senescence and impairs retinal and vascular** 409 **networks**

410 To investigate the mechanisms by which HF plasma induces endothelial cell senescence,  
411 we conducted an in-depth analysis of single-nucleus RNA-seq data from HF patients. The  
412 analysis revealed an upregulation of genes related to inflammation in cardiac tissue from  
413 HF patients, which correlated with an increase in markers of the SASP, including TNF  
414 family cytokines and various interleukins, indicating their involvement in inflammatory

415 responses and cellular senescence (Figure 4A). In parallel, TGF $\beta$ 2 expression was  
416 notably elevated in the HF samples (Figure 4A). To identify the cardiac cellular origin of  
417 elevated TGF $\beta$ 2, we further analyzed the single-nucleus RNA sequencing data from HF  
418 patients. UMAP visualization revealed a widespread increase in TGF $\beta$ 2 expression across  
419 multiple cardiac cell populations in HF samples compared to normal controls (Figure  
420 S2A). Quantitative analysis demonstrated that TGF $\beta$ 2 was significantly upregulated in  
421 cardiac fibroblasts, cardiac neurons, cardiomyocytes, and adipocytes (Figure S2B),  
422 indicating that the elevated circulating TGF $\beta$ 2 in HF is derived from multiple cardiac cell  
423 types rather than a single cellular source. TGF $\beta$ 2, a multifunctional cytokine involved in  
424 regulating cell proliferation, differentiation, and apoptosis, plays a key role in cardiac  
425 remodeling, fibrosis, and inflammation, all of which are characteristics of HF pathology  
426 [23, 24].

427 Further analysis of an additional RNA-Seq dataset (accession number: GSE116250)  
428 related to HF demonstrated that TGF $\beta$ 2 levels were elevated in the blood of patients with  
429 dilated cardiomyopathy (DCM) and ischemic cardiomyopathy (ICM) compared to  
430 healthy controls (Figure 4B). Enzyme-linked immunosorbent assay (ELISA) confirmed  
431 that TGF $\beta$ 2 concentrations in the plasma were higher in HF patients than in controls  
432 (Figures 4C), and this trend was also observed in TAC mice (Figures 4D-E).

433 To directly examine the role of TGF $\beta$ 2 in inducing endothelial cell senescence,  
434 HUVECs were treated with TGF $\beta$ 2. This treatment led to the upregulation of senescence

435 markers, including SA- $\beta$ -gal and P21, confirming that TGF $\beta$ 2 promotes endothelial cell  
436 senescence (Figures 4F-I). To further evaluate the impact of TGF $\beta$ 2 on retinal function,  
437 TGF $\beta$ 2 protein was administered to mice via tail vein injection (Figure 4J). Under  
438 dark-adapted conditions, TGF $\beta$ 2-treated mice exhibited reductions in both a- and b-wave  
439 amplitudes, reflecting impaired rod-bipolar cell signal transduction function (Figures  
440 4K-L). Under light-adapted conditions, reduced b-wave amplitudes further indicated  
441 compromised cone-bipolar cell signal transduction function (Figures 4M-N). Furthermore,  
442 retinal endothelial cells from TGF $\beta$ 2-treated mice exhibited higher levels of SA- $\beta$ -gal  
443 expression than PBS-treated controls, providing additional evidence that TGF $\beta$ 2 induces  
444 endothelial cell senescence (Figures 4O-P).

445 In summary, these findings demonstrated that elevated TGF $\beta$ 2 levels in HF plasma  
446 contribute to endothelial cell senescence and impair retinal function, underscoring its role  
447 in HF-related retinal damage.

#### 448 **TGF $\beta$ 2 induces endothelial cell senescence via the pSMAD2/EP300-H3K9ac-P21** 449 **pathway**

450 The TGF $\beta$  signaling pathway is categorized into two distinct pathways: the classical  
451 SMAD-dependent pathway and the non-canonical SMAD-independent pathway [31, 32].  
452 In the SMAD-dependent pathway, TGF $\beta$ 2 binds to its receptor, triggering the activation  
453 of the SMAD2 protein, which undergoes phosphorylation and subsequently translocates  
454 to the nucleus [31-33]. TGF $\beta$ 2 also activates SMAD5, which is typically associated with

455 bone morphogenetic protein signaling [31-33]. TGF $\beta$ 2 specifically increased SMAD2  
456 phosphorylation without influencing SMAD5, indicating that its primary effects were  
457 mediated through the pSMAD2 pathway (Figure 5A and 5B).

458 Aging is associated with widespread gene dysregulation, primarily driven by  
459 alterations in the chromatin structure [34-36]. TGF $\beta$ 2 treatment increased chromatin  
460 accessibility in endothelial cells, as indicated by the loss of the heterochromatin marker  
461 H3K9me3 (Figure 5C and 5D) [37, 38]. As histone acetylation generally enhances  
462 chromatin accessibility and EP300 is a histone acetyltransferase involved in this process,  
463 we explored the interaction between pSMAD2 and EP300 [38-40].  
464 Co-immunoprecipitation showed pSMAD2 binding EP300 (Figure 5E). Silencing *Smad2*  
465 or *Ep300* reduced SA- $\beta$ -gal and P21 levels (Figure 5F through 5I). ChIP-qPCR  
466 confirmed that TGF $\beta$ 2 enhanced SMAD2/3, EP300, and H3K9ac occupancy at the  
467 *Cdkn1a* (P21) locus, which was abrogated by either knockdown (Figure 5J through 5L).

468 To determine whether the enhanced TGF $\beta$  signaling in endothelial cells under HF  
469 conditions is driven by receptor upregulation, we examined TGFBR2 expression across  
470 multiple systems. RNA-seq analysis of HUVECs treated with TAC versus Sham plasma  
471 did not detect a significant change in *Tgfb2* transcript levels (Table S1). This was further  
472 confirmed by qPCR in HUVECs treated with TAC mouse plasma or HF patient plasma,  
473 as well as in flow cytometry-sorted retinal vascular endothelial cells from TAC mice, all  
474 of which showed no significant alteration in *Tgfb2* expression (Figure S3A-C). These

475 results indicate that the pathological activation of TGF $\beta$  signaling in retinal endothelial  
476 cells is primarily driven by elevated circulating TGF $\beta$ 2 ligand rather than by receptor  
477 upregulation. To further confirm the functional requirement of TGFBR2 in this signaling  
478 cascade, we generated endothelial-specific *Tgfbr2* conditional knockout (cKO) mice.

479 Western blot analysis of flow cytometry-sorted retinal vascular endothelial cells  
480 showed increased expression of pSMAD2, H3K9ac, and P21 in the WT-TAC group,  
481 whereas in the *Tgfbr2* cKO-TAC group, these proteins were restored to normal levels  
482 (Figure 5M and 5N). CHIP-qPCR analysis further confirmed that SMAD2/3, EP300, and  
483 H3K9ac enrichment in the first exon of *Cdkn1a* was elevated in the WT-TAC group but  
484 returned to normal levels in the *Tgfbr2* cKO-TAC group (Figure 5O). CUT & Tag  
485 demonstrated increased pSMAD2 peaks around the transcription start sites (Figure 5P).  
486 GO and motif analyses linked pSMAD2 targets to inflammatory, chemokine, and cell  
487 cycle arrest pathways (Figure 5Q and 5R). Collectively, TGF $\beta$ 2 promotes endothelial  
488 epigenetic senescence by recruiting EP300 to acetylate H3K9 at senescence-associated  
489 genes; blockade of SMAD2 or EP300 prevents this cascade.

#### 490 **Endothelial-specific *Tgfbr2* deletion protects against TAC-induced retinal injury**

491 To determine whether endothelial *Tgfbr2* mediates TAC-induced retinal injury *in vivo*,  
492 we generated endothelial-specific *Tgfbr2* conditional knockout mice by crossing  
493 *Tgfbr2*-flox mice with *Cdh5*-CreERT2 mice followed by tamoxifen induction (Figure  
494 6A). Quantitative PCR analysis of flow cytometry-sorted retinal microvascular

495 endothelial cells confirmed efficient deletion of *Tgfb2* in *Tgfb2* cKO mice (Figure  
496 S4A).

497 We next assessed the effect of endothelial-specific *Tgfb2* deletion on retinal  
498 structure and function after TAC. Under scotopic conditions, TAC-induced reductions in  
499 both a-wave and b-wave amplitudes were markedly attenuated in *Tgfb2* cKO mice  
500 (Figure 6B-C). Similarly, under photopic conditions, the decrease in b-wave amplitudes  
501 observed in WT-TAC mice was significantly rescued by endothelial-specific *Tgfb2*  
502 deletion (Figure 6D-E). OCT analysis further showed that *Tgfb2* cKO preserved total  
503 retinal thickness as well as GCL+IPL and IS+OS thicknesses in TAC mice (Figure 6F-G).  
504 In parallel, SA- $\beta$ -gal staining revealed that TAC-induced endothelial senescence in retinal  
505 microvascular endothelial cells was markedly reduced in *Tgfb2* cKO mice (Figure 6H-I).  
506 Moreover, endothelial-specific *Tgfb2* deletion restored retinal vascular density and  
507 improved vascular diameter distribution in TAC mice (Figure 6J-K).

508 To further determine whether these protective effects were secondary to alterations in  
509 the overall cardiac phenotype, we evaluated cardiac structure and function in WT and  
510 *Tgfb2* cKO mice after TAC. Although *Tgfb2* deletion efficiently reduced endothelial  
511 *Tgfb2* expression, it did not significantly rescue TAC-induced cardiac hypertrophy or  
512 systolic dysfunction, as reflected by HW/TL, LVEF, LVFS, LVID,d, LVID,s, LVEDV,  
513 and LVESV (Figure S4B-I). These findings suggest that the retinal protection conferred  
514 by endothelial-specific *Tgfb2* deletion is unlikely to be merely secondary to improved

515 cardiac performance, but rather reflects a direct role of endothelial *Tgfr2* signaling in  
516 mediating the cardio-retinal pathogenic axis.

517 **Endothelial cell senescence-induced microglial activation leads to retinal neuronal**  
518 **damage**

519 The retina, as part of the central nervous system, is protected by the blood-retinal barrier,  
520 which prevents the entry of circulating factors, such as TGF $\beta$ 2 [41]. We hypothesized  
521 that elevated levels of TGF $\beta$ 2 in the bloodstream of patients with HF may induce  
522 endothelial cell senescence, leading to the release of SASP factors that alter the retinal  
523 microenvironment. Microglia, the primary immune cells within the central nervous  
524 system, are highly responsive to inflammatory and immune signals [42]. Once activated,  
525 microglia exhibit enhanced phagocytic activity and initiate inflammatory cascades, which  
526 results in retinal neuronal damage.

527 Immunofluorescence staining showed an increase in both the number and soma size  
528 of Iba1<sup>+</sup> microglia in the retinas of TAC mice, accompanied by a decrease in the number  
529 of total processes, indicating a transition to an activated state (Figures 7A-B). Further  
530 analysis identified a correlation between the level of microglial activation and SA- $\beta$ -gal  
531 expression in retinal vascular endothelial cells, suggesting that microglial activation is  
532 associated with endothelial cell senescence (Figure 7C).

533 To determine whether microglial activation was mediated by SASP factors,  
534 endothelial cells were incubated with plasma from the Sham and TAC groups for 48 h. 4

535 hours prior to the conclusion of this incubation, the medium was replaced with a  
536 microglia-specific medium to allow conditioning with endothelial cell-derived factors.  
537 The conditioned media was then collected and applied to microglial cultures for an  
538 additional 24-hour incubation. During the final 3 h, microglia were exposed to pHrodo  
539 green-labelled synaptosomes to assess their phagocytic activity. Microglia treated with  
540 conditioned media from endothelial cells exposed to TAC plasma showed an increase in  
541 phagocytic function, as evidenced by the elevated pHrodo green fluorescence intensity  
542 (Figures 7D-F). This increase was diminished following the addition of an anti-TGF $\beta$ 2  
543 antibody (Figures 7G-I). Furthermore, quantitative PCR analysis demonstrated higher  
544 levels of inflammatory markers, including IL-6, TNF- $\alpha$ , IL-1 $\beta$ , and IFN- $\gamma$ , in microglia  
545 treated with conditioned medium from TAC plasma-exposed endothelial cells (Figure  
546 S3D). The addition of anti-TGF $\beta$ 2 antibodies reduced the expression of these  
547 proinflammatory factors (Figure S3E).

548 These findings indicate that TGF $\beta$ 2 in the circulation of patients with HF induces  
549 endothelial cell senescence, which promotes the release of SASP factors. These factors  
550 activate microglia, leading to increased phagocytic activity and the initiation of  
551 inflammatory processes, ultimately contributing to retinal neuronal damage.

552 **Microglial depletion mitigates retinal structural and functional damage in heart**  
553 **failure**

554 To investigate the role of microglial activation in heart failure-induced retinal damage,  
555 we administered PLX5622, a small molecule inhibitor of colony-stimulating factor-1  
556 receptor, to eliminate microglia in Sham or TAC mice (Figure 7J). In dark-adapted ERG  
557 tests, TAC mice showed reduced a-wave and b-wave amplitudes, indicating impaired  
558 retinal function (Figures 7K-L). After PLX5622 treatment, the ERG responses in TAC  
559 mice showed improvement (Figures 7K-L). Under light-adapted conditions, PLX5622  
560 treatment also increased b-wave amplitudes across different light intensities, indicating a  
561 protective effect on retinal function (Figures 7M-N). OCT revealed that TAC mice  
562 exhibited reduced total retinal thickness, GCL+IPL thickness, and photoreceptor layer  
563 thickness, reflecting structural damage to the retina (Figures 7O-P). Treatment with  
564 PLX5622 restored these retinal layers, further indicating the involvement of microglia in  
565 the structural damage observed in heart failure (Figures 7O-P). In conclusion, this study  
566 demonstrates that endothelial cell senescence triggered by HF contributes to both  
567 structural and functional retinal damage mediated by microglial activation. Microglial  
568 depletion using PLX5622 interrupts this pathological process, leading to improved retinal  
569 integrity and function.

570 **Anti-TGFβ2 antibody reduces endothelial cell senescence induced by HF plasma**

571 Our findings indicate that elevated TGFβ2 levels in the plasma of patients with HF  
572 promote endothelial cell senescence by activating the pSMAD2/EP300-H3K9ac-P21  
573 signaling pathway. To further delineate the role of TGFβ2 in this process and assess the  
574 effect of an anti-TGFβ2 antibody on endothelial cell senescence, we treated endothelial  
575 cells with plasma from both Sham and TAC mice and co-administered either IgG control  
576 or anti-TGFβ2 antibody. After 48 h of treatment, analysis of senescence-associated  
577 markers demonstrated that TAC plasma increased the expression of SA-β-gal, P16<sup>Ink4a</sup>,  
578 and P21 in endothelial cells, while anti-TGFβ2 antibody treatment effectively reduced  
579 these markers (Figures 8A-F). Additionally, the decline in H3K9me3 expression  
580 observed in TAC plasma-treated cells was mitigated by the anti-TGFβ2 antibody (Figures  
581 8G-H).

582 To expand our understanding of the impact of the anti-TGFβ2 antibody on gene  
583 expression in endothelial cells, RNA-seq analysis was conducted. Principal component  
584 analysis (PCA) demonstrated that antibody treatment modified the gene expression  
585 profile altered by TAC plasma (Figure S5A). GO enrichment analysis indicated that  
586 antibody treatment enhanced pathways associated with DNA repair, cell cycle regulation,  
587 and DNA damage response, while downregulating pathways linked to cellular senescence,  
588 such as stress response, TP53 signaling, and oxidative stress response (Figure S5B).

589 These findings suggest that the anti-TGF $\beta$ 2 antibody effectively reduces endothelial cell  
590 senescence induced by TAC plasma.

591 **Intravitreal injections of the anti-TGF $\beta$ 2 antibody mitigate retinal and vascular**  
592 **network damage in HF mice**

593 To assess the therapeutic potential of anti-TGF $\beta$ 2 antibody on retinal and vascular  
594 damage induced by HF, we performed Sham or TAC surgery in 2-month-old mice. After  
595 a 1-month recovery period, from months 3 to 5, the mice received 6 intravitreal injections  
596 of anti-TGF $\beta$ 2 antibody. OCT and ERG were performed at 5 months of age (Figure 8I).  
597 ERG results showed that the TAC group exhibited reduced a- and b-wave amplitudes  
598 across various light intensities under dark-adapted conditions (Figure 8J-K). Treatment  
599 with anti-TGF $\beta$ 2 antibody improved these amplitudes (Figure 8J-K). Similarly, under  
600 light-adapted conditions, b-wave amplitudes in the TAC group were reduced at different  
601 light intensities, and these amplitudes improved following anti-TGF $\beta$ 2 antibody treatment  
602 (Figure 8L-M), indicating restoration of retinal function.

603 OCT scans further revealed that the TAC group had a reduced total retinal thickness,  
604 with thinning observed in the GCL+IPL and the IS+OS. Treatment with the anti-TGF $\beta$ 2  
605 antibody restored these retinal layer thicknesses (Figure 8N-O). Additionally, SA- $\beta$ -gal  
606 staining indicated an increased expression of SA- $\beta$ -gal in retinal vascular endothelial cells  
607 in the TAC group (Figure 8P-Q). This expression was reduced following treatment with

608 the anti-TGF $\beta$ 2 antibody, suggesting that the antibody effectively mitigated retinal  
609 vascular endothelial cell senescence (Figure 8P-Q).

610 In conclusion, our findings demonstrate that anti-TGF $\beta$ 2 antibodies reduce  
611 HF-induced retinal and vascular damage. In vitro experiments showed that the antibody  
612 attenuated endothelial cell senescence caused by HF plasma by inhibiting the  
613 pSMAD2/EP300-H3K9ac-P21 pathway and normalizing associated gene expression. In  
614 vivo, intravitreal injection of anti-TGF $\beta$ 2 antibody reduced endothelial cell senescence  
615 and improved retinal structure and function in HF mice. These results support the  
616 potential clinical application of the anti-TGF $\beta$ 2 antibody in preventing and treating  
617 retinal damage associated with heart failure.

618 **Anti-TGF $\beta$ 2 antibody reduces endothelial cell senescence in multiple organs of heart**  
619 **failure mice**

620 To assess the impact of the anti-TGF $\beta$ 2 antibody on endothelial cell senescence across  
621 multiple organs induced by HF, we conducted Sham or TAC surgery on 2-month-old  
622 mice. 1 month after surgery, anti-TGF $\beta$ 2 antibody treatment was initiated via tail vein  
623 injection and administered at regular intervals until the mice reached 5 months of age,  
624 with a total of 6 injections (Figure 9A). Immunofluorescence staining was used to  
625 examine the expression levels of the senescence marker SA- $\beta$ -gal in endothelial cells  
626 from the spleen, lungs, kidneys, and heart of TAC mice. Compared with the Sham group,  
627 the TAC surgery led to an increase in SA- $\beta$ -gal levels in endothelial cells, indicating

628 accelerated senescence in these organs due to HF (Figures 9B-I). In contrast, the group  
629 treated with the anti-TGF $\beta$ 2 antibody showed reduced SA- $\beta$ -gal expression across all  
630 examined organs, with levels comparable to those observed in the Sham group (Figures  
631 9B-I). These findings suggest that anti-TGF $\beta$ 2 antibody administration reduces  
632 endothelial cell senescence induced by TAC across multiple organs, including the spleen,  
633 lungs, kidneys, and heart, demonstrating its potential protective role against HF-related  
634 systemic endothelial damage.

635 **Anti-TGF $\beta$ 2 antibody mitigates senescence of primary human retinal microvascular**  
636 **endothelial cells induced by plasma from HF patients**

637 To explore the role of TGF $\beta$ 2 in human retinal microvascular endothelial cell senescence  
638 and to evaluate the potential therapeutic effect of an anti-TGF $\beta$ 2 antibody, we selected  
639 primary human retinal microvascular endothelial cells (HRMECs) for experiments. This  
640 model was chosen to closely replicate physiological conditions in humans. HRMECs  
641 were treated with plasma derived from either healthy individuals or HF patients for 48 h.  
642 Simultaneously, an IgG control or anti-TGF $\beta$ 2 antibody was administered to investigate  
643 its ability to inhibit senescence induced by HF plasma (Figure 9J). The results indicated  
644 that plasma from HF patients, compared to that from healthy controls, elevated the  
645 expression of SA- $\beta$ -gal, P16<sup>Ink4a</sup>, and P21 in HRMECs, suggesting accelerated cellular  
646 senescence (Figures 9K-P). The anti-TGF $\beta$ 2 antibody reduced the expression of these

647 senescence markers, demonstrating its inhibitory effect on endothelial cell senescence  
648 triggered by HF plasma (Figures 9K-P).

649 In conclusion, this study illustrates that the anti-TGF $\beta$ 2 antibody confers a protective  
650 effect against multi-organ endothelial cell senescence associated with HF. Specifically, in  
651 HRMECs, the antibody reduced senescence induced by HF plasma. These findings  
652 underscore the potential clinical relevance of targeting TGF $\beta$ 2 to prevent or treat retinal  
653 microvascular damage in HF, highlighting its therapeutic potential.

654 **Discussion**

655 Our research indicated that TGF $\beta$ 2 has a new but detrimental role in HF-associated  
656 retinal damages. Our results showed that plasma from HF mice damaged the retina and  
657 vascular tissues of healthy mice, indicating that HF has systemic effects on distant organs  
658 via blood. Under HF conditions, TGF $\beta$ 2 levels were found to be high which induced  
659 endothelial cell epigenetic senescence via the pSMAD2/EP300-H3K9ac-P21 signalling.  
660 SASP factor release activates retinal microglia in senescence, enhancing phagocytosis,  
661 causing neuroinflammatory response, and retinal neuronal damage. Earlier studies have  
662 established the importance of TGF $\beta$ 2 in inflammation and tissue fibrosis. However, ours  
663 is the first to discover its major role in the cardio-retinal axis.

664 Based on clinical studies that have been done so far, HF has an impact on the  
665 diameter of the retinal vessels likely related to systemic inflammation and metabolic  
666 disturbance. However, most are observational with a lack of clear mechanistic insight [15,  
667 17-21]. Our study was aimed at clarifying how the TGF $\beta$ 2 contributes to HF-induced  
668 retinal pathology, with a specific emphasis on the TGF $\beta$ 2-mediated epigenetic regulation  
669 of endothelial cell senescence. This fills an important gap in our knowledge of HF-related  
670 retinal dysfunction along with the concept of the cardio-retinal axis. The retina,  
671 contributing to the central nervous system, works closely with brain activity and cardiac  
672 health [15]. Hypertensive retinal damage is caused by hemodynamic and metabolic  
673 imbalance but also by the direct damage to retinal vascular endothelial cells by

674 circulating factors like TGF $\beta$ 2. This mechanism presents a novel explanation for the  
675 impact of HF on the retina.

676 To increase the translational value of our data, we validated our findings using  
677 primary human retinal microvascular endothelial cells treated with plasma from patients  
678 with heart failure. The similarity of the study system to human physiological conditions  
679 increased the utility of our findings. According to the evidence, treatment with  
680 anti-TGF $\beta$ 2 antibody reduced the expression of senescence markers. This indicates that  
681 anti-TGF $\beta$ 2 antibody can mitigate pathogenic factors in human heart failure plasma that  
682 induce endothelial cell senescence. The validated approach in human cells and patient  
683 samples bolsters the clinical potential to target TGF $\beta$ 2 to prevent or treat retinal  
684 microvascular damage associated with heart failure. The insights into mechanism  
685 aforementioned from our mouse models and support for the clinical application of  
686 anti-TGF $\beta$ 2 approach findings are consistent with the.

687 Significantly, TAC-induced retinal dysfunction, vascular degeneration, and  
688 endothelial senescence were greatly attenuated by endothelial-specific *Tgfb2* deletion,  
689 although global cardiac remodelling and systolic function were not significantly  
690 improved. This separation supports our main conclusion that the cardio-retinal axis  
691 observed in this study represents an actively driven inter organ pathogenic pathway  
692 driven by the circulation of TGF $\beta$ 2 rather than a passive peripheral result of cardiac  
693 dysfunction. As represented in Figure 9, our results show that the systemic neutralization

694 of TGF $\beta$ 2 attenuated cardiac endothelial cell senescence confirming the role of  
695 circulating TGF $\beta$ 2 in cardiac endothelial injury. Nevertheless, findings indicate that  
696 endothelial senescence is not the primary driver of cardiac dysfunction in the TAC model,  
697 thus blockade of this pathway alone is not enough to achieve significant recovery of  
698 overall cardiac function. As the main site of disease, the heart experiences prolonged  
699 mechanical pressure overload. The pressure overload itself directly drives myocardial  
700 hypertrophy, fibrotic remodeling, and cardiomyocyte injury. Meanwhile, blockade of  
701 endothelial TGF $\beta$ 2 signaling does not erase this mechanical stress. In contrast, the retina  
702 being a distal target organ gets damaged mostly by circulating TGF $\beta$ 2 and the blockade  
703 of this signal is sufficient to effectively break the local pathological cascade. The above  
704 observations open up possibilities that selective protection of distant neurovascular units  
705 may confer meaningful therapeutic benefit in retinal injury due to heart failure even in the  
706 presence of incomplete correction of major cardiac lesion.

707 Additional research should examine the potential function of TGF- $\beta$ 2 in other  
708 conditions. Particular attention should be paid to any potential effects in central nervous  
709 system disease, including Alzheimer's disease and multiple sclerosis where TGF- $\beta$ 2  
710 actions are not well understood [43, 44]. Moreover, it would be of interest to assess the  
711 interaction of TGF $\beta$ 2 with other signaling pathways, such as Wnt/ $\beta$ -catenin and PI3K/Akt,  
712 to better comprehend the regulating networks associated with systemic organ damage  
713 [45-48]. Therapeutic strategies targeting TGF $\beta$ 2 and its downstream pathways may help

714 decrease damage from heart failure to the retina and other distant organ systems. The  
715 development of these studies will not only bolster the theoretical framework for novel  
716 treatment strategies but also offer new avenues for addressing heart failure issues.

717 In conclusion, our findings provide an alternative strategy for targeting TGF $\beta$ 2 in  
718 retinal damage caused by HF. When TGF $\beta$ 2 is targeted early, it can block EC senescence,  
719 microglial activation and retinal degeneration. This therapeutic approach can offer better  
720 outcomes for patients and decrease systemic damage caused by heart failure. Thus, it  
721 presents an important opportunity for clinical use.

722 **Abbreviations**

723 cKO: conditional knockout; CUT&Tag: cleavage under targets and tagmentation; EP300:  
724 E1A binding protein p300; ERG: electroretinography; EV: extracellular vesicle; GCL:  
725 ganglion cell layer; GO: Gene Ontology; GSEA: Gene Set Enrichment Analysis; H3K9ac:  
726 histone H3 lysine 9 acetylation; HF: heart failure; HRMEC: human retinal microvascular  
727 endothelial cell; HUVEC: human umbilical vein endothelial cell; IFN- $\gamma$ : interferon  
728 gamma; IgG: immunoglobulin G; IL-1 $\beta$ : interleukin-1 beta; IL-6: interleukin-6; INL:  
729 inner nuclear layer; IPL: inner plexiform layer; LVEF: left ventricular ejection fraction;  
730 LVFS: left ventricular fractional shortening; NVU: neurovascular unit; OCT: optical  
731 coherence tomography; ONL: outer nuclear layer; OPL: outer plexiform layer; PBS:  
732 phosphate-buffered saline; PCA: principal component analysis; pSMAD2:  
733 phosphorylated SMAD2; SA- $\beta$ -gal: senescence-associated  $\beta$ -galactosidase; SASP:  
734 senescence-associated secretory phenotype; snRNA-seq: single-nucleus RNA sequencing;  
735 TAC: transverse aortic constriction; TGF $\beta$ 2: transforming growth factor beta 2; TGFBR2:  
736 transforming growth factor beta receptor 2; TNF- $\alpha$ : tumor necrosis factor alpha; UMAP:  
737 Uniform Manifold Approximation and Projection.

738 **Supplementary Material**

739 Supplementary figures and methods are available at Theranostics online.

740 **Acknowledgements**

741 The authors declare that no AI tools were used in the preparation of this manuscript.

742 **Funding**

743 This study was supported by grants from the National Natural Science Foundation of  
744 China (Grant No. 82270417 to G.L.), the Natural Science Foundation of Fujian Province  
745 (Grant No. 2023J01677 to G.L.), and the Natural Science Foundation of Xiamen (Grant  
746 Nos. 3502Z20191103 to G.L. and 3502Z20209007, 3502Z20224032, 3502Z20241002 to  
747 Y.W.).

748 **Author Contributions**

749 M.W., S.Z., and T.A. are the co-first authors. Y.W., S.Z., and G.L. are the  
750 co-corresponding authors. S.Z. and M.W. conceived the study and designed the  
751 experiments. M.W. and S.Z. drafted the manuscript. M.W., S.Z., G.L. and Y.W. edited  
752 the manuscript. M.W., S.Z., T.A., SY.Z., and X.W. performed most of the experiments  
753 and data analyses. G.L. and Y.W. contributed to the reagents, materials, and analytical  
754 tools. All the authors have reviewed and approved the final manuscript.

755 **Data Availability**

756 The single-nucleus RNA sequencing data used in this study are available from the  
757 European Genome-Phenome Archive (EGAS00001006374). The RNA-seq data are  
758 available from the Gene Expression Omnibus (GSE116250). All other data supporting  
759 the findings of this study are included in the manuscript and supplementary materials or  
760 are available from the corresponding author upon reasonable request.

761 **Competing Interests**

762 The authors have declared that no competing interests exist.

763 **References**

- 764 1. Savarese G, Lund LH. Global Public Health Burden of Heart Failure. *Card Fail Rev.*  
765 2017; 3: 7–11.
- 766 2. Savarese G, Becher PM, Lund LH, Seferovic P, Rosano GMC, Coats AJS. Global  
767 burden of heart failure: a comprehensive and updated review of epidemiology.  
768 *Cardiovasc Res.* 2023; 118: 3272–87.
- 769 3. Henkel DM, Redfield MM, Weston SA, Gerber Y, Roger VL. Death in heart failure:  
770 a community perspective. *Circ Heart Fail.* 2008; 1: 91–7.
- 771 4. Tanai E, Frantz S. Pathophysiology of Heart Failure. *Compr Physiol.* 2015; 6:  
772 187–214.
- 773 5. Hunter WG, Kelly JP, McGarrah RW, 3rd, Kraus WE, Shah SH. Metabolic  
774 Dysfunction in Heart Failure: Diagnostic, Prognostic, and Pathophysiologic Insights  
775 From Metabolomic Profiling. *Curr Heart Fail Rep.* 2016; 13: 119–31.
- 776 6. Doehner W, Frenneaux M, Anker SD. Metabolic impairment in heart failure: the  
777 myocardial and systemic perspective. *J Am Coll Cardiol.* 2014; 64: 1388–400.
- 778 7. Ciccarelli M, Dawson D, Falcao-Pires I, Giacca M, Hamdani N, Heymans S, et al.  
779 Reciprocal organ interactions during heart failure: a position paper from the ESC  
780 Working Group on Myocardial Function. *Cardiovasc Res.* 2021; 117: 2416–33.

- 781 8. Muller OJ, Heckmann MB, Ding L, Rapti K, Rangrez AY, Gerken T, et al.  
782 Comprehensive plasma and tissue profiling reveals systemic metabolic alterations in  
783 cardiac hypertrophy and failure. *Cardiovasc Res.* 2019; 115: 1296–305.
- 784 9. He X, Du T, Long T, Liao X, Dong Y, Huang ZP. Signaling cascades in the failing  
785 heart and emerging therapeutic strategies. *Signal Transduct Target Ther.* 2022; 7: 134.
- 786 10. Li H, Hastings MH, Rhee J, Trager LE, Roh JD, Rosenzweig A. Targeting  
787 Age-Related Pathways in Heart Failure. *Circ Res.* 2020; 126: 533–51.
- 788 11. Sbolli M, Fiuzat M, Cani D, O'Connor CM. Depression and heart failure: the lonely  
789 comorbidity. *Eur J Heart Fail.* 2020; 22: 2007–17.
- 790 12. Frey A, Sell R, Homola GA, Malsch C, Kraft P, Gunreben I, et al. Cognitive Deficits  
791 and Related Brain Lesions in Patients With Chronic Heart Failure. *JACC Heart Fail.* 2018;  
792 6: 583–92.
- 793 13. Dridi H, Liu Y, Reiken S, Liu X, Argyrousi EK, Yuan Q, et al. Heart failure-induced  
794 cognitive dysfunction is mediated by intracellular Ca(2+) leak through ryanodine receptor  
795 type 2. *Nat Neurosci.* 2023; 26: 1365–78.
- 796 14. Wang MD, Zhang S, Liu XY, Wang PP, Zhu YF, Zhu JR, et al. Salvianolic acid B  
797 ameliorates retinal deficits in an early-stage Alzheimer's disease mouse model through  
798 downregulating BACE1 and Abeta generation. *Acta Pharmacol Sin.* 2023; 44: 2151–68.
- 799 15. Nagele MP, Barthelmes J, Ludovici V, Cantatore S, von Eckardstein A, Enseleit F, et  
800 al. Retinal microvascular dysfunction in heart failure. *Eur Heart J.* 2018; 39: 47–56.

- 801 16. Allon R, Aronov M, Belkin M, Maor E, Shechter M, Fabian ID. Retinal  
802 Microvascular Signs as Screening and Prognostic Factors for Cardiac Disease: A  
803 Systematic Review of Current Evidence. *Am J Med.* 2021; 134: 36–47 e7.
- 804 17. Horton WB, Barrett EJ. Microvascular Dysfunction in Diabetes Mellitus and  
805 Cardiometabolic Disease. *Endocr Rev.* 2021; 42: 29–55.
- 806 18. Seferovic PM, Polovina M. The eyes are the mirror of the heart: role of retinal  
807 microvascular abnormalities in predicting long-term risk of heart failure. *Eur J Heart Fail.*  
808 2019; 21: 1216–8.
- 809 19. Barthelmes J, Nagele MP, Cantatore S, Novruzov E, Ludovici V, von Eckardstein A,  
810 et al. Retinal microvascular dysfunction in patients with coronary artery disease with and  
811 without heart failure: a continuum? *Eur J Heart Fail.* 2019; 21: 988–97.
- 812 20. Hanssen H, Streese L, Vilser W. Retinal vessel diameters and function in  
813 cardiovascular risk and disease. *Prog Retin Eye Res.* 2022; 91: 101095.
- 814 21. Chaikijurajai T, Ehlers JP, Tang WHW. Retinal Microvasculature: A Potential  
815 Window Into Heart Failure Prevention. *JACC Heart Fail.* 2022; 10: 785–91.
- 816 22. Morikawa M, Derynck R, Miyazono K. TGF-beta and the TGF-beta Family:  
817 Context-Dependent Roles in Cell and Tissue Physiology. *Cold Spring Harb Perspect Biol.*  
818 2016; 8.
- 819 23. Goumans MJ, Ten Dijke P. TGF-beta Signaling in Control of Cardiovascular  
820 Function. *Cold Spring Harb Perspect Biol.* 2018; 10: a022210.

821 24. Yue Y, Meng K, Pu Y, Zhang X. Transforming growth factor beta (TGF-beta)  
822 mediates cardiac fibrosis and induces diabetic cardiomyopathy. *Diabetes Res Clin Pract.*  
823 2017; 133: 124–30.

824 25. Schiattarella GG, Altamirano F, Tong D, French KM, Villalobos E, Kim SY, et al.  
825 Nitrosative stress drives heart failure with preserved ejection fraction. *Nature.* 2019; 568:  
826 351–6.

827 26. Gao Y, Hong Y, Huang L, Zheng S, Zhang H, Wang S, et al. beta2-microglobulin  
828 functions as an endogenous NMDAR antagonist to impair synaptic function. *Cell.* 2023;  
829 186: 1026–38 e20.

830 27. Zheng Q, Li G, Wang S, Zhou Y, Liu K, Gao Y, et al. Trisomy 21-induced  
831 dysregulation of microglial homeostasis in Alzheimer's brains is mediated by USP25. *Sci*  
832 *Adv.* 2021; 7: eabe1340.

833 28. Guan JS, Haggarty SJ, Giacometti E, Dannenberg JH, Joseph N, Gao J, et al.  
834 HDAC2 negatively regulates memory formation and synaptic plasticity. *Nature.* 2009;  
835 459: 55–60.

836 29. Zheng Q, Song B, Li G, Cai F, Wu M, Zhao Y, et al. USP25 inhibition ameliorates  
837 Alzheimer's pathology through the regulation of APP processing and Abeta generation. *J*  
838 *Clin Invest.* 2022; 132.

- 839 30. Zhao Y, Zheng Q, Hong Y, Gao Y, Hu J, Lang M, et al. beta(2)-Microglobulin  
840 coaggregates with Abeta and contributes to amyloid pathology and cognitive deficits in  
841 Alzheimer's disease model mice. *Nat Neurosci.* 2023; 26: 1170–84.
- 842 31. Meng XM, Nikolic-Paterson DJ, Lan HY. TGF-beta: the master regulator of fibrosis.  
843 *Nat Rev Nephrol.* 2016; 12: 325–38.
- 844 32. Deng Z, Fan T, Xiao C, Tian H, Zheng Y, Li C, et al. TGF-beta signaling in health,  
845 disease, and therapeutics. *Signal Transduct Target Ther.* 2024; 9: 61.
- 846 33. Massague J, Sheppard D. TGF-beta signaling in health and disease. *Cell.* 2023; 186:  
847 4007–37.
- 848 34. Cai Y, Song W, Li J, Jing Y, Liang C, Zhang L, et al. The landscape of aging. *Sci*  
849 *China Life Sci.* 2022; 65: 2354–454.
- 850 35. Lopez-Otin C, Blasco MA, Partridge L, Serrano M, Kroemer G. Hallmarks of aging:  
851 An expanding universe. *Cell.* 2023; 186: 243–78.
- 852 36. Meharena HS, Marco A, Dileep V, Lockshin ER, Akatsu GY, Mullahoo J, et al.  
853 Down-syndrome-induced senescence disrupts the nuclear architecture of neural  
854 progenitors. *Cell Stem Cell.* 2022; 29: 116–30 e7.
- 855 37. Zhao H, Ji Q, Wu Z, Wang S, Ren J, Yan K, et al. Destabilizing heterochromatin by  
856 APOE mediates senescence. *Nat Aging.* 2022; 2: 303–16.
- 857 38. Hwang JY, Aromolaran KA, Zukin RS. The emerging field of epigenetics in  
858 neurodegeneration and neuroprotection. *Nat Rev Neurosci.* 2017; 18: 347–61.

- 859 39. White J, Derheimer FA, Jensen-Pergakes K, O'Connell S, Sharma S, Spiegel N, et al.  
860 Histone lysine acetyltransferase inhibitors: an emerging class of drugs for cancer therapy.  
861 Trends Pharmacol Sci. 2024; 45: 243–54.
- 862 40. Zhang W, Qu J, Liu GH, Belmonte JCI. The ageing epigenome and its rejuvenation.  
863 Nat Rev Mol Cell Biol. 2020; 21: 137–50.
- 864 41. Rudraraju M, Narayanan SP, Somanath PR. Regulation of blood-retinal barrier  
865 cell-junctions in diabetic retinopathy. Pharmacol Res. 2020; 161: 105115.
- 866 42. Prinz M, Jung S, Priller J. Microglia Biology: One Century of Evolving Concepts.  
867 Cell. 2019; 179: 292–311.
- 868 43. Vidovic N, Spittau B. Microglial Transforming Growth Factor-beta Signaling in  
869 Alzheimer's Disease. Int J Mol Sci. 2024; 25: 3090.
- 870 44. Meyers EA, Kessler JA. TGF-beta Family Signaling in Neural and Neuronal  
871 Differentiation, Development, and Function. Cold Spring Harb Perspect Biol. 2017; 9:  
872 a022244.
- 873 45. Zhang L, Zhou F, ten Dijke P. Signaling interplay between transforming growth  
874 factor-beta receptor and PI3K/AKT pathways in cancer. Trends Biochem Sci. 2013; 38:  
875 612–20.
- 876 46. Yu JS, Ramasamy TS, Murphy N, Holt MK, Czapiewski R, Wei SK, et al.  
877 PI3K/mTORC2 regulates TGF-beta/Activin signalling by modulating Smad2/3 activity  
878 via linker phosphorylation. Nat Commun. 2015; 6: 7212.

- 879 47. Dzialo E, Czepiel M, Tkacz K, Siedlar M, Kania G, Blyszczuk P. WNT/beta-Catenin  
880 Signaling Promotes TGF-beta-Mediated Activation of Human Cardiac Fibroblasts by  
881 Enhancing IL-11 Production. *Int J Mol Sci.* 2021; 22: 10072.
- 882 48. Carethers JM. Intersection of transforming growth factor-beta and Wnt signaling  
883 pathways in colorectal cancer and metastasis. *Gastroenterology.* 2009; 137: 33–6.
- 884



887 **Figure 1. Retinal and vascular networks were damaged in the HF mouse model**

888 (A) Schematic representation of the experimental timeline illustrating progression from  
889 Sham or TAC surgery to the end of the study. Surgeries were performed at 2 months of  
890 age using monthly echocardiographic assessments. OCT and ERG were conducted at five  
891 months of age, followed by tissue collection.

892 (B-C) Representative histological images showing cross-sections of cardiomyocytes and  
893 fibrosis in the Sham and TAC groups, along with quantification of cardiomyocyte area  
894 and fibrosis percentage, n = 16 mice per group. Scale bar, 50  $\mu\text{m}$  in the upper panels and  
895 25  $\mu\text{m}$  in the lower panels.

896 (D) Echocardiographic images showing left ventricular function from 2 to 5 months of  
897 age in mice. Horizontal scale bar, 300 ms; vertical scale bar, 6 mm.

898 (E) Heart weight (HW) to tibia length (TL) ratio at 5 months in Sham and TAC mice, n =  
899 8 mice per group.

900 (F-K) Temporal analysis of LVEF%, LVFS%, LVID,d, LVID,s, LVEDV, and LVESV in  
901 the Sham and TAC groups, n = 8 mice per group.

902 (L-M) ERG waveform responses and quantification of a-wave and b-wave amplitudes  
903 under scotopic conditions in Sham and TAC mice, n = 24 to 30 eyes from 12 to 15 mice  
904 per group.

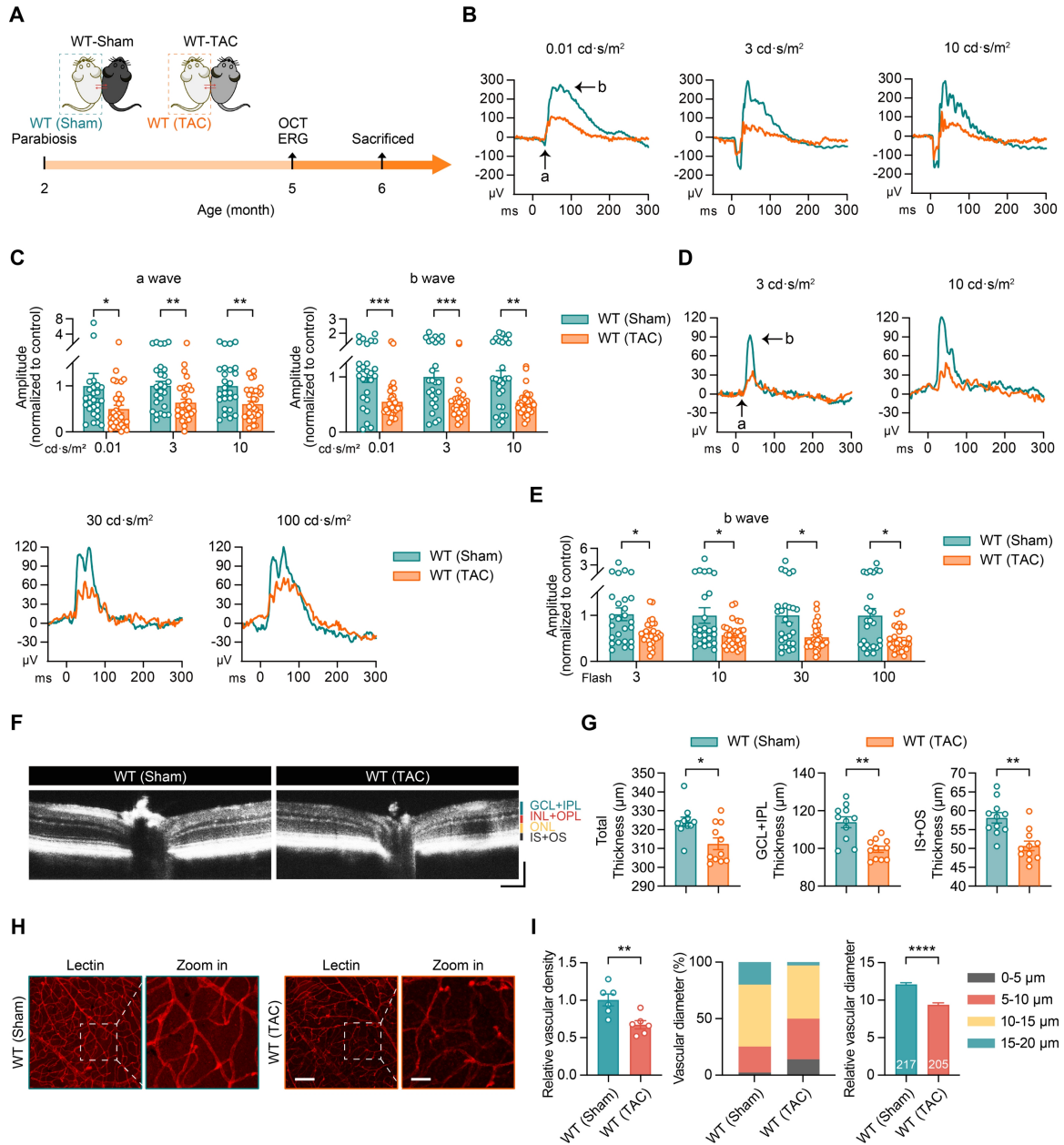
905 (N-O) ERG b-wave responses and amplitude comparisons under photopic conditions  
906 between Sham and TAC mice, n = 24 to 28 eyes from 12 to 14 mice per group.

907 (P-Q) OCT cross-sectional images of retinas from Sham and TAC mice, with  
908 quantifications of total retinal thickness, GCL+IPL, and IS+OS thicknesses, n = 13 to 14  
909 mice per group. Horizontal scale bar, 130  $\mu\text{m}$ ; vertical scale bar, 150  $\mu\text{m}$ .

910 (R-S) Lectin-stained images showing retinal vasculature, including magnified views, 3D  
911 reconstructions, and projected vascular diameters, with quantification of retinal vascular  
912 density, the proportion of vessels with different diameters, and vessel diameter analysis, n  
913 = 239 to 253 vessels from 6 mice per group. Scale bar, 100  $\mu\text{m}$  in the lectin images and  
914 30  $\mu\text{m}$  in the zoom-in images.

915 All data are presented as the mean  $\pm$  SEM. *P*-values were determined using Student's  
916 t-test in (C), (E), (Q), and (S); repeated measures ANOVA with Bonferroni's post hoc  
917 analysis in (F-K); and one-way ANOVA with Tukey's post hoc analysis in (M) and (O).  
918 \**p* < 0.05, \*\**p* < 0.01, \*\*\**p* < 0.001, \*\*\*\**p* < 0.0001.

919



920

921 **Figure 2. Parabiosis with blood of HF mice leads to retinal and vascular damage**

922 (A) Schematic representation of experimental design. WT (Sham) refers to WT mice in

923 parabiotic pairings with sham-operated mice and WT (TAC) refers to WT mice in

924 parabiotic pairings with TAC-operated mice.

925 (B-C) ERG waveform responses and quantification of a-wave and b-wave amplitudes  
926 under scotopic conditions in WT (sham) and WT (TAC) mice, n = 26 to 30 eyes from 13  
927 to 15 mice per group.

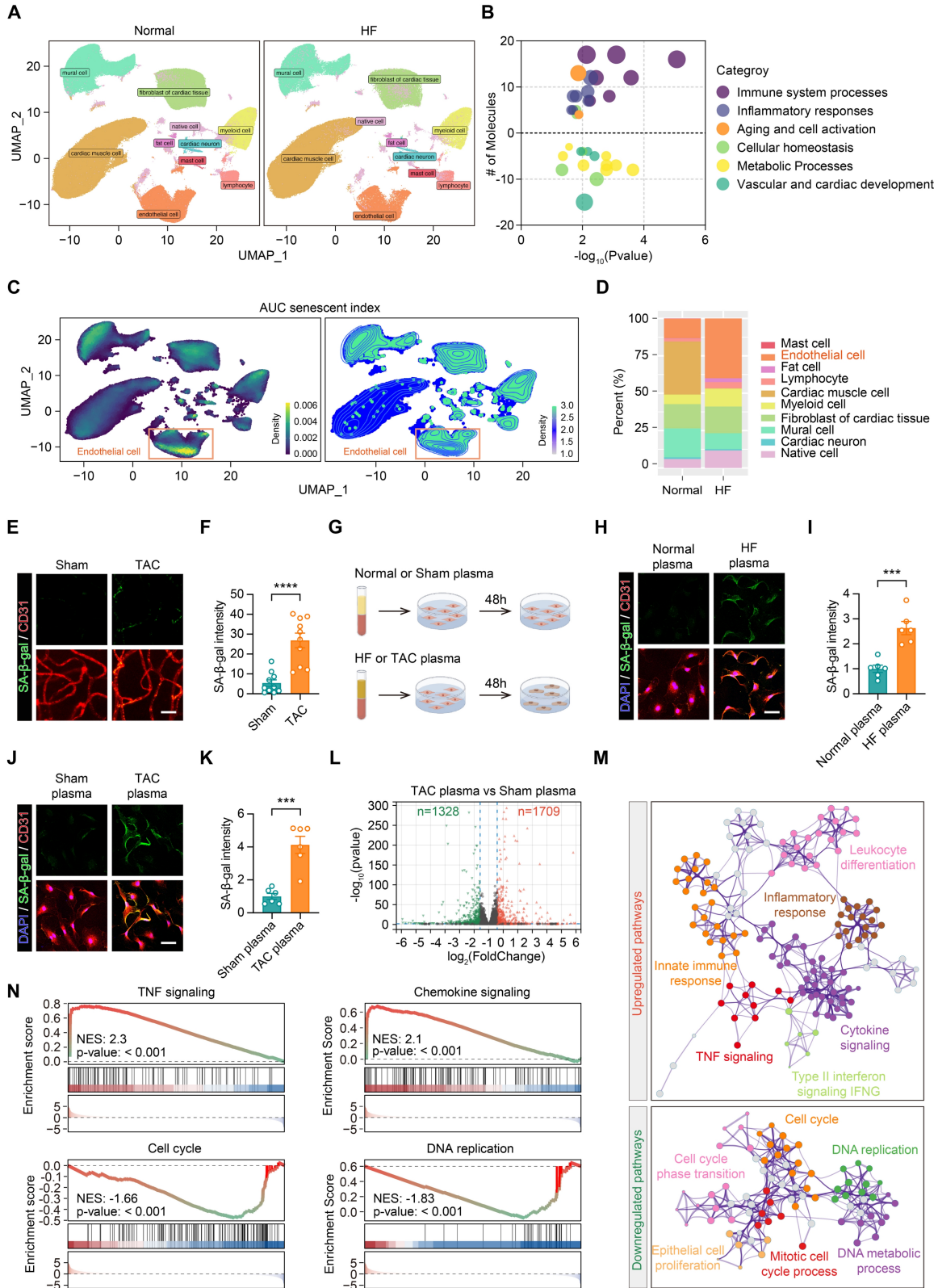
928 (D-E) ERG waveform responses and quantification of b-wave amplitudes under photopic  
929 conditions in WT (Sham) and WT (TAC) mice, n = 26 to 30 eyes from 13 to 15 mice per  
930 group.

931 (F-G) OCT images of the retinas from WT (Sham) and WT (TAC) mice, with  
932 quantification of total retinal thickness, GCL+IPL, and IS+OS thicknesses, n = 11 mice  
933 per group. Horizontal scale bar, 130  $\mu\text{m}$ ; vertical scale bar, 150  $\mu\text{m}$ .

934 (H-I) Lectin-stained images showing retinal vascular networks in WT (Sham) and WT  
935 (TAC) mice, with quantification of vascular density, the proportion of vessels with  
936 different diameters, and vessel diameter analysis, n = 205 to 217 vessels from 6 mice per  
937 group. Scale bar, 100  $\mu\text{m}$  in the lectin images and 30  $\mu\text{m}$  in the zoom-in images.

938 All data are presented as mean  $\pm$  SEM. *P*-values were determined using one-way  
939 ANOVA with Tukey's post hoc analysis in (C) and (E); Student's t-test in (G) and (I). \**p*  
940 < 0.05, \*\**p* < 0.01, \*\*\**p* < 0.001, \*\*\*\**p* < 0.0001.

941



942

943 **Figure 3. Heart failure blood induces endothelial cell senescence**

944 (A) UMAP projection showing the distribution of cardiac cells in samples from normal  
945 and HF patients. The single-cell sequencing data was from the European  
946 Genome-Phenome Archive (accession number: EGAS00001006374).

947 (B) GO enrichment analysis illustrating pathway enrichment of differentially expressed  
948 genes between HF patients and normal controls. Compared to normal controls, pathways  
949 above the dashed line are upregulated in heart failure patients, while pathways below the  
950 dashed line are downregulated.

951 (C-D) UMAP projections displaying the AUC senescent index for each cell type in heart  
952 failure samples, with a comparison of the proportion of senescent cells between normal  
953 and heart failure samples.

954 (E-F) Representative images and statistical analysis showing changes in SA- $\beta$ -gal  
955 staining in retinal vascular endothelial cells from Sham and TAC mice, n = 10 mice per  
956 group. Scale bar, 35  $\mu$ m.

957 (G) Schematic diagram of the experimental design, showing treatment of endothelial cells  
958 with plasma from normal or heart failure patients, as well as plasma from Sham or TAC  
959 mice for 48 h.

960 (H-I) Representative images and statistical analysis showing changes in SA- $\beta$ -gal staining  
961 in endothelial cells after treatment with plasma from normal and heart failure patients, n =  
962 6 independent experiments per group. Scale bar, 50  $\mu$ m.

963 (J-K) Representative images and statistical analysis showing changes in SA- $\beta$ -gal  
964 staining in endothelial cells after treatment with plasma from Sham and TAC mice,  $n = 6$   
965 independent experiments per group. Scale bar, 50  $\mu\text{m}$ .

966 (L) Volcano plot showing differentially expressed genes in endothelial cells treated with  
967 plasma from Sham and TAC mice,  $n = 3$  independent experiments per group.

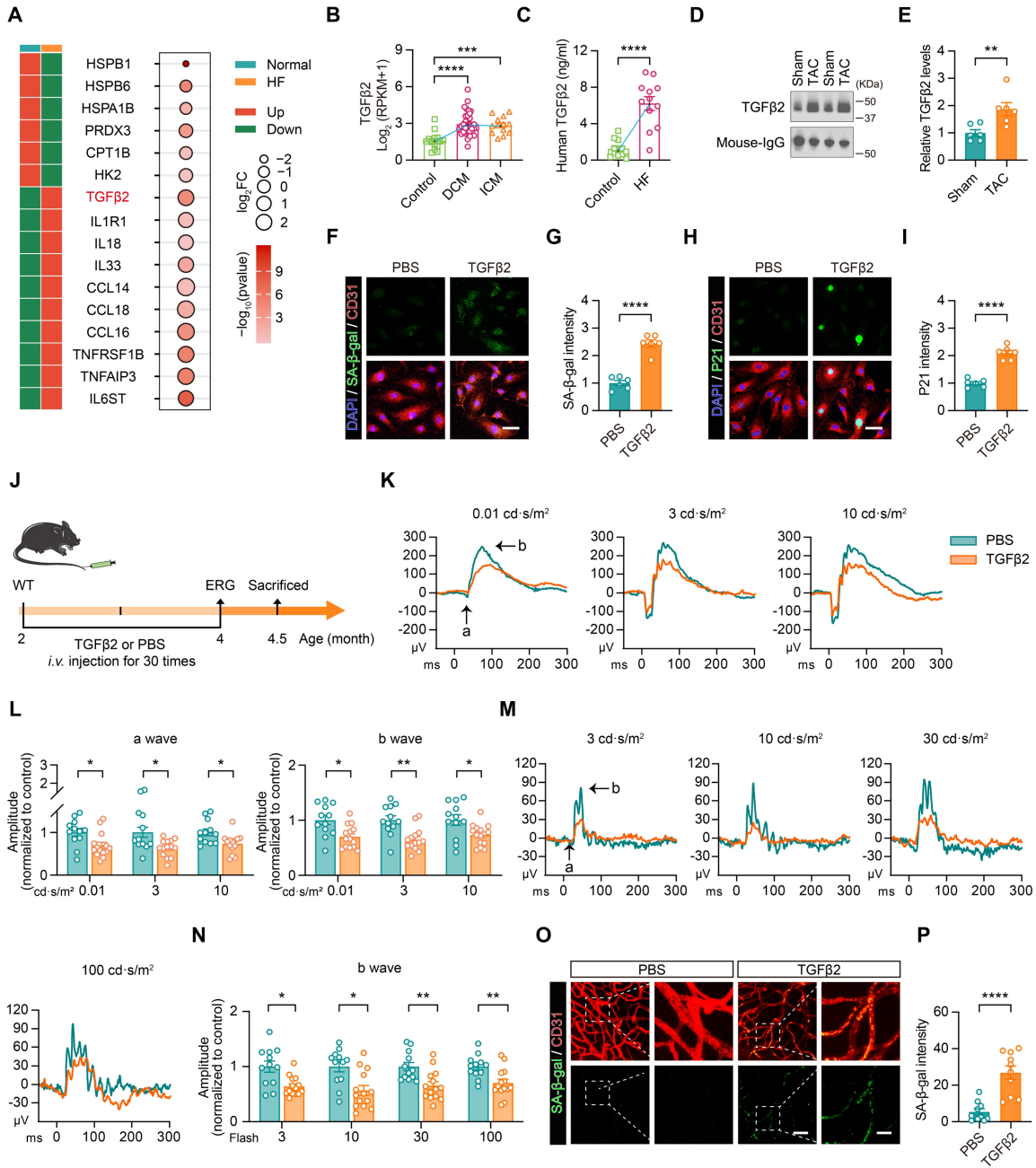
968 (M) Network diagram showing significantly upregulated and downregulated  
969 GO-enriched pathways in endothelial cells treated with plasma from the Sham and TAC  
970 mice.

971 (N) Gene set enrichment analysis (GSEA) showing significant enrichment of TNF  
972 signaling, chemokine signaling, cell cycle, and DNA replication-related gene sets in  
973 endothelial cells treated with TAC plasma.

974 All data are presented as mean  $\pm$  SEM.  $P$ -values were determined using Student's  $t$ -test in

975 (F), (I), and (K). \*\*\* $p < 0.001$ , \*\*\*\* $p < 0.0001$ .

976



977

978 **Figure 4. Elevated TGFβ2 in heart failure blood induces endothelial cell senescence**

979 **and retinal damage**

980 (A) Heatmap of RNA-seq data from GEO (accession number: GSE116250) showing  
981 differentially expressed genes significantly upregulated and downregulated in heart  
982 samples from HF patients compared to the normal controls.

983 (B) RNA-Seq dataset GSE116250 showing TGF $\beta$ 2 expression levels in normal, DCM,  
984 and ICM heart samples, n = 13 to 37 cases per group.

985 (C) ELISA showing differences in plasma TGF $\beta$ 2 concentrations between normal and  
986 heart failure patients, n = 12 cases per group.

987 (D-E) Western blotting and quantification showing TGF $\beta$ 2 protein levels in plasma from  
988 Sham and TAC mice, n = 6 mice per group.

989 (F-G) Representative images and quantification of SA- $\beta$ -gal staining in endothelial cells  
990 treated with PBS or TGF $\beta$ 2 (6 ng/mL) for 48 h, n = 6 independent experiments per group.  
991 Scale bar, 50  $\mu$ m.

992 (H-I) Representative images and quantification showing changes in P21 expression levels  
993 in endothelial cells treated with PBS or TGF $\beta$ 2 for 48 h, n = 6 independent experiments  
994 per group. Scale bar, 50  $\mu$ m.

995 (J) Schematic representation of the experimental timeline showing TGF $\beta$ 2 protein  
996 injection into the tail vein of mice, starting at 2 months of age and administered every two  
997 days for two months. ERG was conducted at 4 months of age.

998 (K-L) ERG waveform responses and quantification of a-wave and b-wave amplitudes  
999 under scotopic conditions in mice injected with PBS and TGFβ2 protein, n = 12 to 14  
1000 eyes from 6 to 7 mice per group.

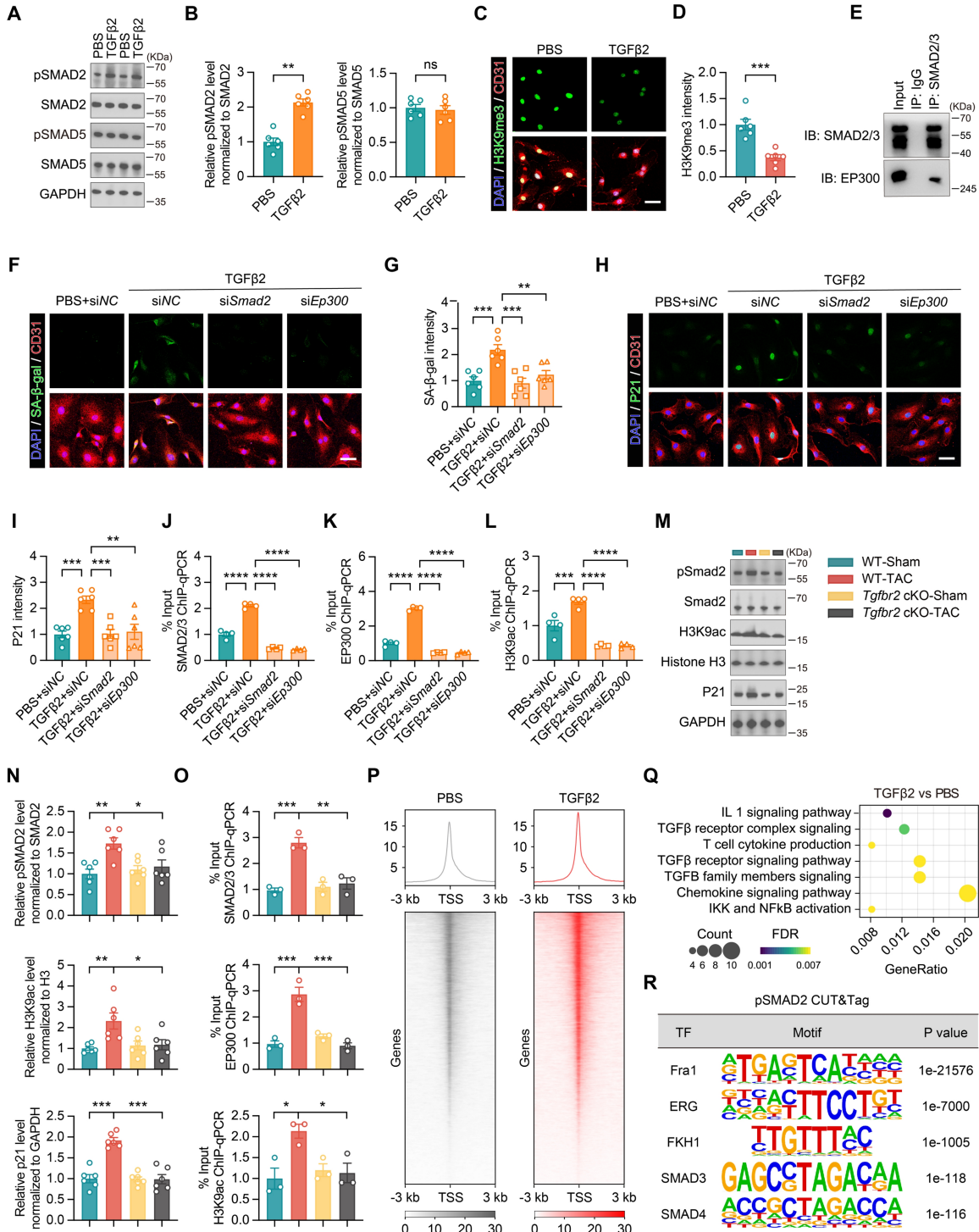
1001 (M-N) ERG b-wave responses and amplitude comparisons under photopic conditions on  
1002 mice injected with PBS and TGFβ2 protein, n = 12 to 14 eyes from 6 to 7 mice per group.

1003 (O-P) Representative images and quantification of changes in SA-β-gal staining in retinal  
1004 vascular endothelial cells from PBS and TGFβ2 treated mice, n = 10 mice per group.

1005 Scale bar, 50 μm in the overview images and 15 μm in the zoom in images.

1006 All data are presented as mean ± SEM. *P*-values were determined by one-way ANOVA  
1007 with Tukey's post hoc analysis in (B), (L), and (N), and by Student's *t*-test in (C), (E),  
1008 (G), (I), and (P). \**p* < 0.05, \*\**p* < 0.01, \*\*\**p* < 0.001, \*\*\*\**p* < 0.0001.

1009



1010

1011 **Figure 5. TGFβ2 induces endothelial cell senescence by activating the**

1012 **pSMAD2/EP300-H3K9ac-P21 pathway**

1013 (A-B) Western blotting and quantification showing changes in pSMAD2, SMAD2,  
1014 pSMAD5, and SMAD5 protein levels in endothelial cells treated with PBS or TGFβ2, n =  
1015 6 independent experiments per group.

1016 (C-D) Representative images and quantification of H3K9me3 staining in endothelial cells  
1017 after 48 h of treatment with PBS or TGFβ2, n = 6 independent experiments per group.  
1018 Scale bar, 50 μm.

1019 (E) Co-IP showing the interaction between SMAD2/3 and EP300 in endothelial cells.

1020 (F-G) Representative images and quantification of SA-β-gal staining in endothelial cells  
1021 treated with PBS or TGFβ2 (siNC, si*Smad2*, or si*Ep300*), n = 6 independent experiments  
1022 per group. Scale bar, 50 μm.

1023 (H-I) Representative images and quantification of P21 staining in endothelial cells under  
1024 different treatment conditions, n = 6 independent experiments per group. Scale bar, 50  
1025 μm.

1026 (J-L) ChIP-qPCR analysis showing enrichment of SMAD2/3, EP300, and H3K9ac at the  
1027 first exon of the *Cdkn1a* (P21) gene in endothelial cells under different treatments, n = 4  
1028 independent experiments per group.

1029 (M-N) Western blot analysis and quantification showing changes in pSMAD2, SMAD2,  
1030 H3K9ac, Histone H3, and P21 protein levels in flow cytometry-sorted retinal vascular  
1031 endothelial cells from different experimental groups. 3 mice per group were pooled for  
1032 each sample, n = 6 per group.

1033 (O) ChIP-qPCR analysis demonstrating the enrichment of SMAD2/3, EP300, and  
1034 H3K9ac at the first exon of Cdkn1a in flow cytometry-sorted retinal vascular endothelial  
1035 cells from different experimental groups. 3 mice per group were pooled for each sample;  
1036 n = 6 per group.

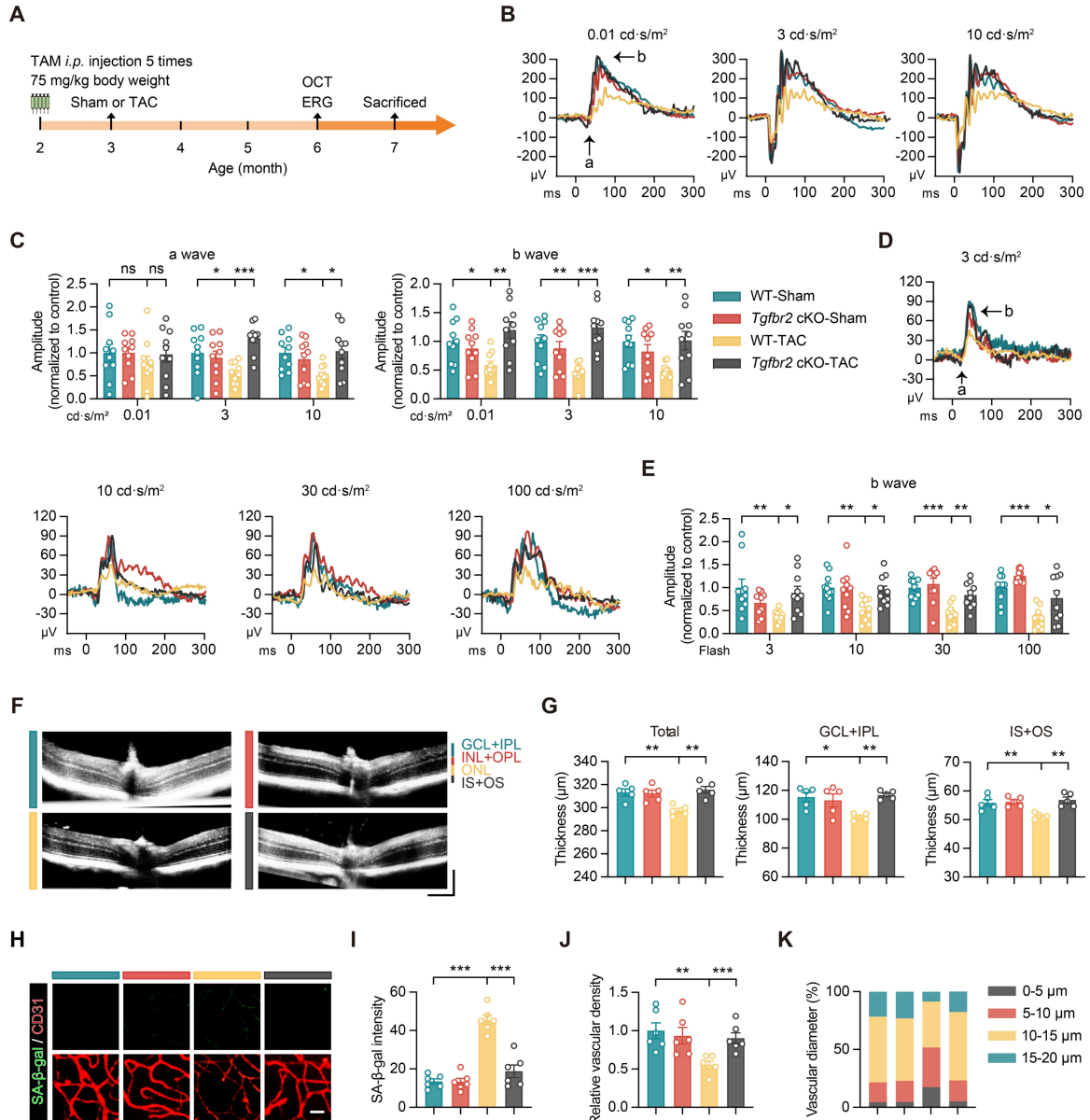
1037 (P) CUT&Tag analysis showing pSMAD2 enrichment near the transcription start site  
1038 (TSS) in endothelial cells, comparing PBS and TGF $\beta$ 2 treatments, n = 3 independent  
1039 experiments per group.

1040 (Q) GO enrichment analysis showing the significantly upregulated pathways following  
1041 TGF $\beta$ 2 treatment.

1042 (R) pSMAD2 CUT&Tag analysis showing the binding of Fra1, ERG, FKH1, SMAD3,  
1043 and SMAD4 to pSMAD2 sites.

1044 All data are presented as the mean  $\pm$  SEM. P-values were determined by Student's t-test  
1045 in (B) and (D), and by one-way ANOVA with Dunnett's post hoc analysis in (G), (I-L),  
1046 and (N-O). ns, not significant. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ , \*\*\*\* $p < 0.0001$ .

1047



1048  
1049 **Figure 6. Endothelial-specific *Tgfr2* deletion protects against TAC-induced retinal**  
1050 **dysfunction, vascular degeneration, and endothelial senescence**

1051 (A) Schematic representation of the experimental timeline. Tamoxifen was administered  
1052 by intraperitoneal injection before Sham or TAC surgery to induce endothelial-specific  
1053 *Tgfr2* deletion. OCT and ERG were performed at 6 months of age, followed by tissue  
1054 collection at 7 months of age.

1055 (B-C) Representative scotopic ERG waveforms and quantification of a-wave and b-wave  
1056 amplitudes at flash intensities of 0.01, 3, and 10 cd·s/m<sup>2</sup> in WT-Sham, *Tgfr2* cKO-Sham,  
1057 WT-TAC, and *Tgfr2* cKO-TAC mice.

1058 (D-E) Representative photopic ERG waveforms and quantification of b-wave amplitudes  
1059 at flash intensities of 3, 10, 30, and 100 cd·s/m<sup>2</sup> across the indicated groups.

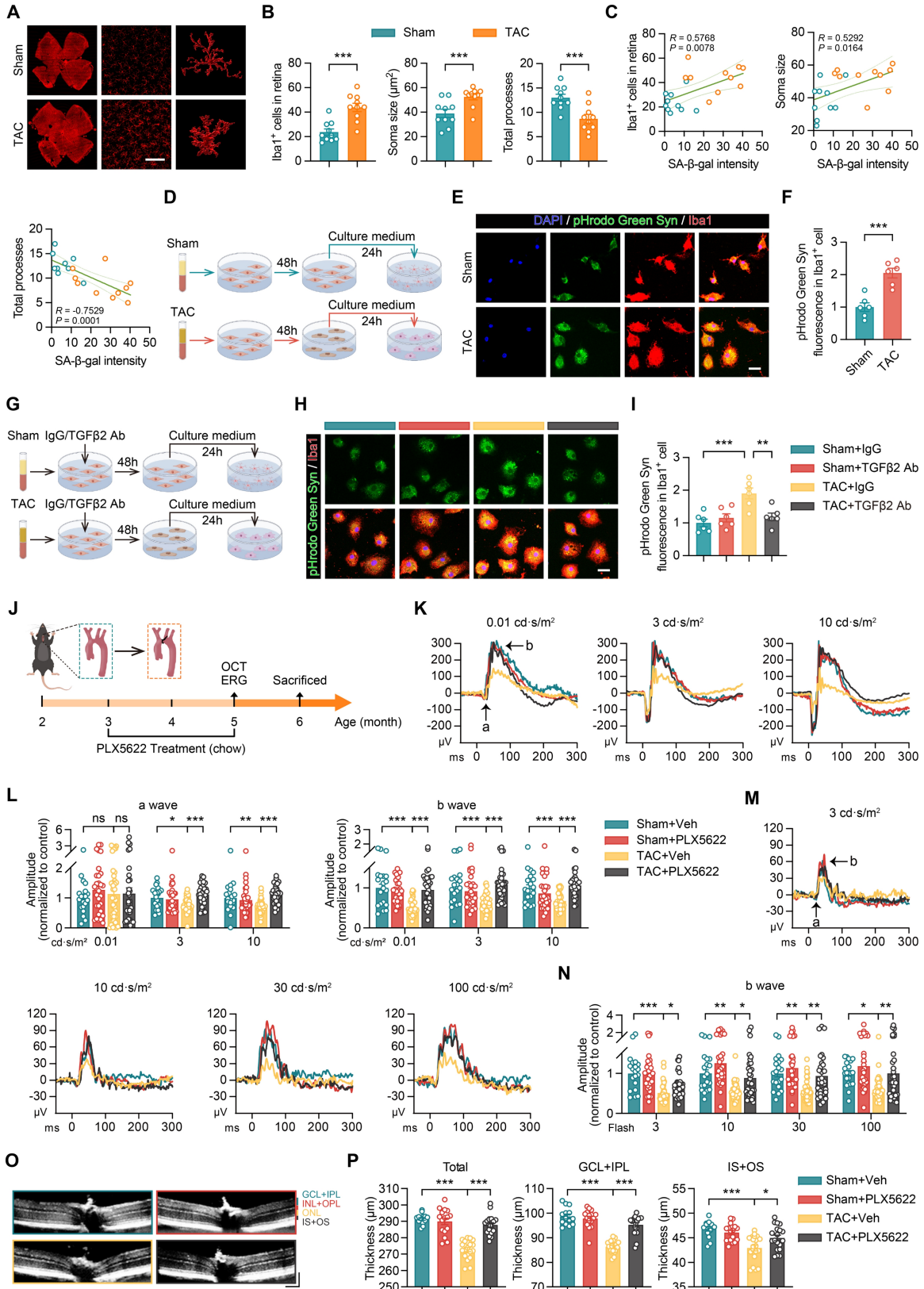
1060 (F-G) Representative OCT images and quantification of total retinal thickness, GCL+IPL  
1061 thickness, and IS+OS thickness in the indicated groups. Horizontal scale bar, 130 μm;  
1062 vertical scale bar, 150 μm.

1063 (H-I) Representative images and quantification of SA-β-gal staining in CD31<sup>+</sup> retinal  
1064 microvascular endothelial cells from the indicated groups. Scale bar, 35 μm.

1065 (J-K) Quantification of relative retinal vascular density and the distribution of retinal  
1066 vascular diameters across the indicated groups.

1067 All data are presented as mean ± SEM. P-values were determined by one-way ANOVA  
1068 with Tukey's post hoc analysis in (C) and (E), and by one-way ANOVA with Dunnett's  
1069 post hoc analysis in (G), (I), and (J). ns, not significant. \**p* < 0.05, \*\**p* < 0.01, \*\*\**p* <  
1070 0.001.

1071



1073 **Figure 7. Activation of microglia by senescent endothelial cells leads to retinal**  
1074 **structural and functional damage**

1075 (A-B) Representative images and quantitative analysis showing the number of Iba1<sup>+</sup> cells,  
1076 cell area, and total number of processes in the retinas of Sham and TAC mice, n = 10  
1077 mice per group. Scale bar, 150 μm.

1078 (C) Correlation between the number of Iba1<sup>+</sup> cells, soma area, total number of processes,  
1079 and SA-β-gal intensity in retinal endothelial cells (Figure 3E-F), n = 10 mice per group.

1080 (D) Schematic representation of the experimental timeline.

1081 (E-F) Representative images and quantitative analysis showing the fluorescence intensity  
1082 of pHrodo Green Syn in Iba1<sup>+</sup> microglia after plasma treatment in the Sham and TAC  
1083 groups, n = 6 independent experiments per group. Scale bar, 25 μm.

1084 (G) Schematic representation of the experimental timeline.

1085 (H-I) Representative images and quantitative analysis showing changes in the  
1086 fluorescence intensity of pHrodo Green Syn in Iba1<sup>+</sup> microglia and the fluorescence  
1087 intensity of Iba1<sup>+</sup> cells in different treatment groups, n = 6 independent experiments per  
1088 group. Scale bar, 25 μm.

1089 (J) Schematic showing the experimental timeline for Sham or TAC surgery performed on  
1090 2-month-old WT mice, followed by PLX5622 treatment (1.2 g/kg, provided ad libitum)  
1091 from 3 to 5 months. OCT and ERG measurements were performed at 5 months, and  
1092 tissue analysis was conducted at 6 months of age.

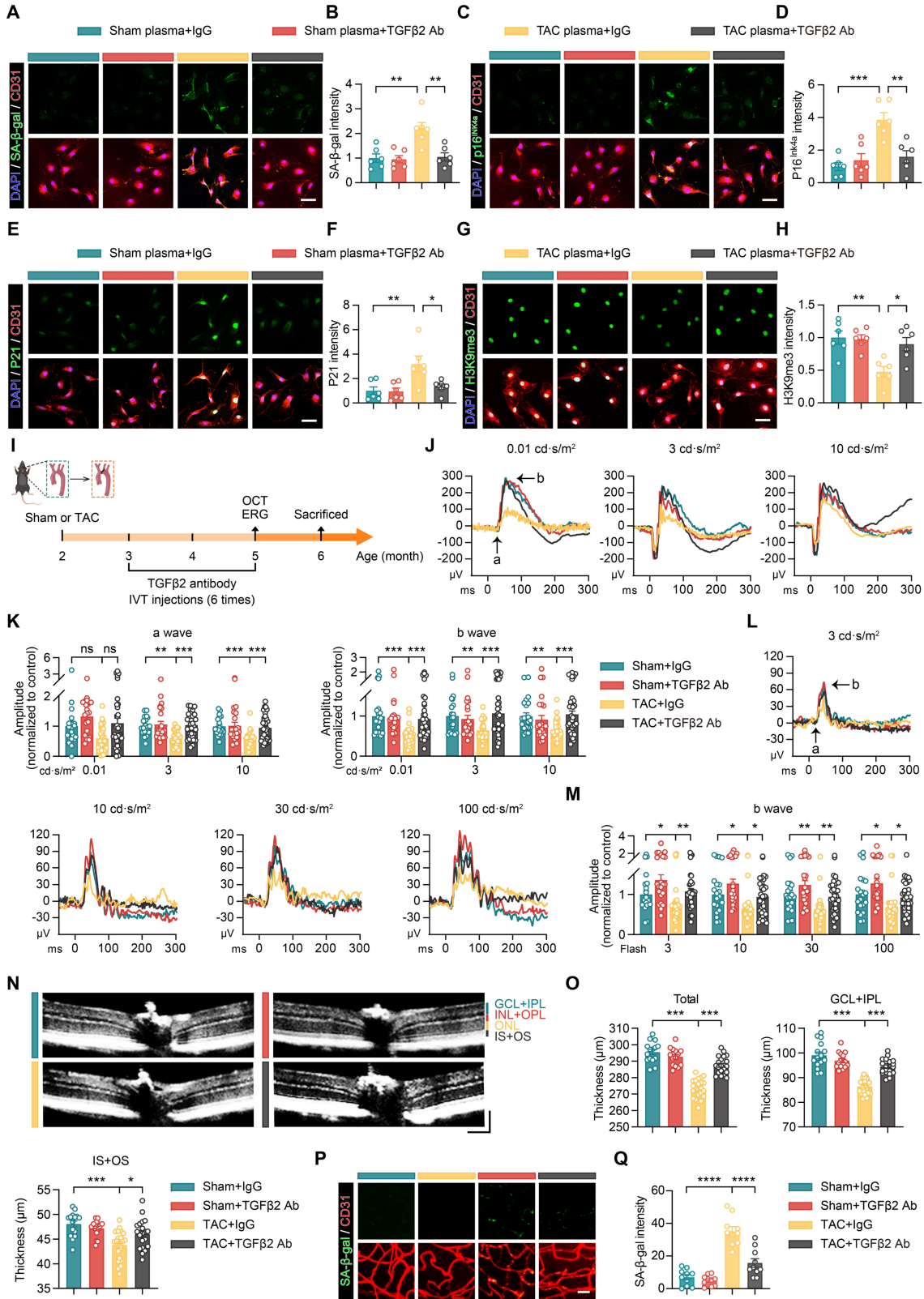
1093 (K-L) Scotopic ERG responses showing waveform plots and quantitative analysis of  
1094 a-wave and b-wave amplitudes across groups, n = 20 to 40 eyes from 10 to 20 mice per  
1095 group.

1096 (M-N) Photopic ERG responses showing waveform plots and quantitative analysis of  
1097 b-wave amplitudes across groups, n = 20 to 40 eyes from 10 to 20 mice per group.

1098 (O-P) OCT cross-sectional images showing retinal structural changes across groups, with  
1099 quantitative analysis, n = 14 to 20 mice per group. Horizontal scale bar, 130  $\mu\text{m}$ ; vertical  
1100 scale bar, 150  $\mu\text{m}$ .

1101 All data are presented as mean  $\pm$  SEM. *P*-values were determined using Student's t-test in  
1102 (B) and (F), one-way ANOVA with Dunnett's post hoc analysis in (I) and (P), and  
1103 one-way ANOVA with Tukey's post hoc analysis in (L) and (N). ns, not significant. \**p* <  
1104 0.05, \*\**p* < 0.01, \*\*\**p* < 0.001.

1105



1107 **Figure 8. Anti-TGF $\beta$ 2 antibody treatment improves retinal and vascular network**  
1108 **damage in a mouse model of heart failure**

1109 (A-H) Representative images and quantitative analysis showing the results after treating  
1110 endothelial cells with Sham or TAC plasma and co-treatment with IgG or anti-TGF $\beta$ 2  
1111 antibody for 48 h. The expression levels of SA- $\beta$ -gal (A-B), P16<sup>Ink4a</sup> (C-D), P21 (E-F),  
1112 and H3K9me3 (G-H) were assessed, n = 6 independent experiments per group. Scale bar,  
1113 50  $\mu$ m.

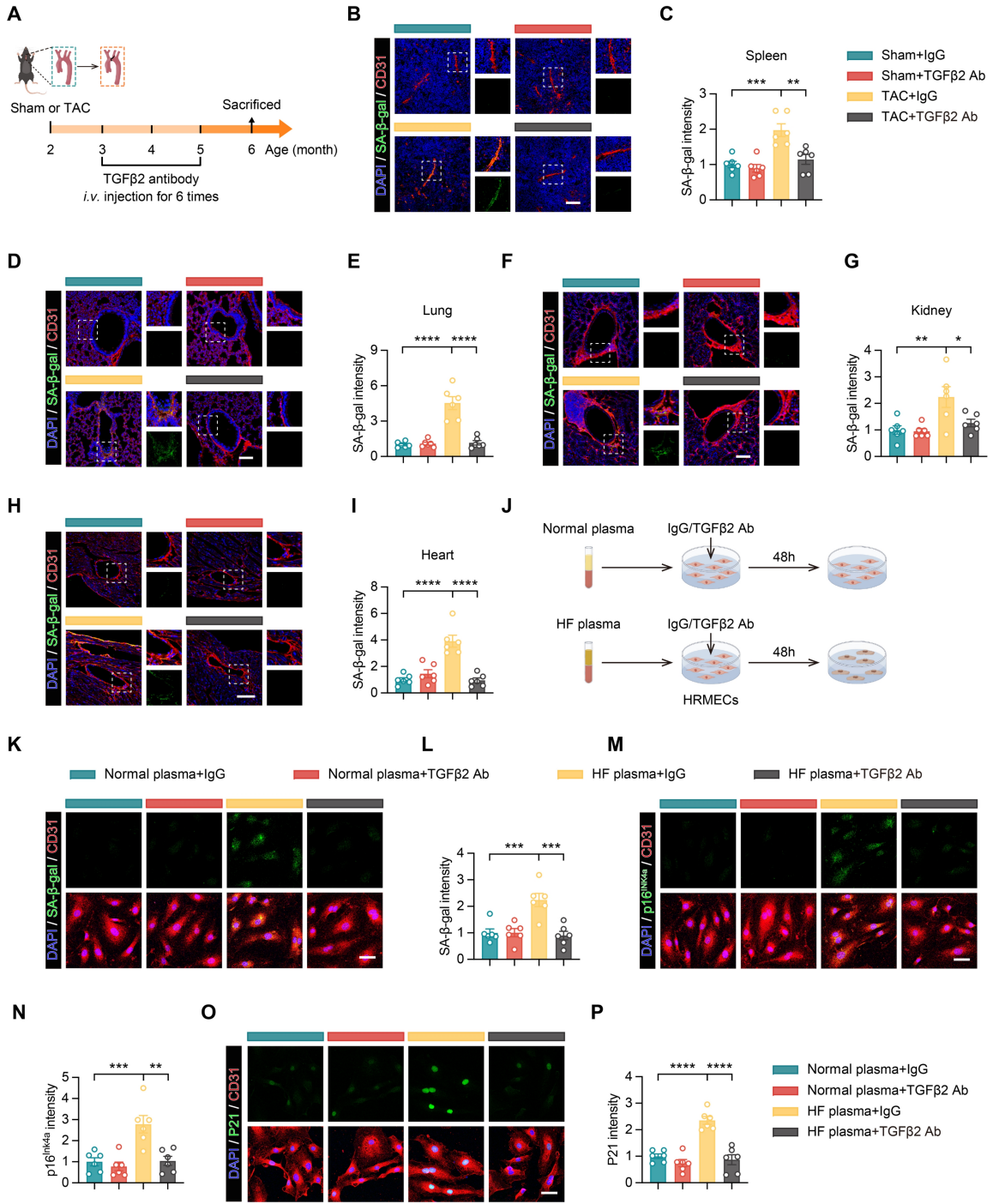
1114 (I) Schematic of the experimental timeline. Sham or TAC surgery was performed on  
1115 2-month-old WT mice, followed by a 1-month recovery period. From 3 to 5 months of  
1116 age, the mice received 6 intravitreal injections of the anti-TGF $\beta$ 2 antibody. OCT and  
1117 ERG assessments were conducted at 5 months of age, followed by histological analysis at  
1118 6 months of age.

1119 (J-K) Scotopic ERG responses showing ERG waveforms and quantitative analysis of  
1120 a-wave and b-wave amplitudes in each group, n = 20 to 34 eyes from 10 to 17 mice per  
1121 group.

1122 (L-M) Photopic ERG responses showing ERG waveforms and quantitative analysis of  
1123 b-wave amplitudes in each group, n = 20 to 34 eyes from 10 to 17 mice per group.

1124 (N-O) OCT cross-sectional images showing structural changes in the retina across groups,  
1125 with quantitative analysis, n = 15 to 22 mice per group. Horizontal scale bar, 130  $\mu$ m;  
1126 vertical scale bar, 150  $\mu$ m.

1127 (P-Q) Representative images and quantitative analysis of SA- $\beta$ -gal expression in retinal  
1128 vascular endothelial cells across groups, n = 10 mice per group. Scale bar, 35  $\mu$ m.  
1129 All data are presented as mean  $\pm$  SEM. *P*-values were determined by one-way ANOVA  
1130 with Dunnett's post hoc analysis in (B), (D), (F), (H), (O), and (Q) and by one-way  
1131 ANOVA with Tukey's post hoc analysis in (K) and (M). ns, not significant. \**p* < 0.05,  
1132 \*\**p* < 0.01, \*\*\**p* < 0.001, \*\*\*\**p* < 0.0001.  
1133



1134

1135 **Figure 9. TGFβ2 antibody ameliorates multi-organ endothelial cell senescence**  
1136 **caused by HF and prevents HRMECs senescence induced by HF patient plasma**

1137 (A) Schematic timeline showing the administration of the TGFβ2 antibody from 2 to 5  
1138 months of age in Sham and TAC mice.

1139 (B-I) Representative immunofluorescence images showing the intensity of SA-β-gal  
1140 staining in CD31-labeled endothelial cells from the spleen, lungs, kidneys, and heart, n =  
1141 6 mice per group. Scale bar, 100 μm.

1142 (J) Experimental workflow diagram.

1143 (K-P) Representative images and quantitative analyses of HRMECs treated with normal  
1144 or HF plasma, along with IgG or TGFβ2 antibody for 48 h. The expression levels of  
1145 SA-β-gal (K-L), P16<sup>Ink4a</sup> (M-N), and P21 (O-P) were measured in the endothelial cells, n  
1146 = 6 independent experiments per group. Scale bar, 50 μm.

1147 All data are presented as mean ± SEM. *P*-values were determined by one-way ANOVA  
1148 with Dunnett's post hoc analysis for (C), (E), (G), (I), (L), (N), and (P). \**p* < 0.05, \*\**p* <  
1149 0.01, \*\*\**p* < 0.001, \*\*\*\**p* < 0.0001.