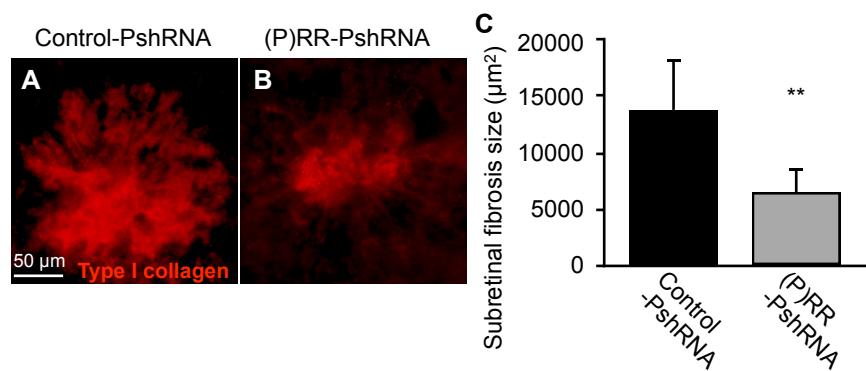


**Supplemental Information**

**Suppression of Choroidal Neovascularization  
and Fibrosis by a Novel RNAi Therapeutic  
Agent against (Pro)renin Receptor**

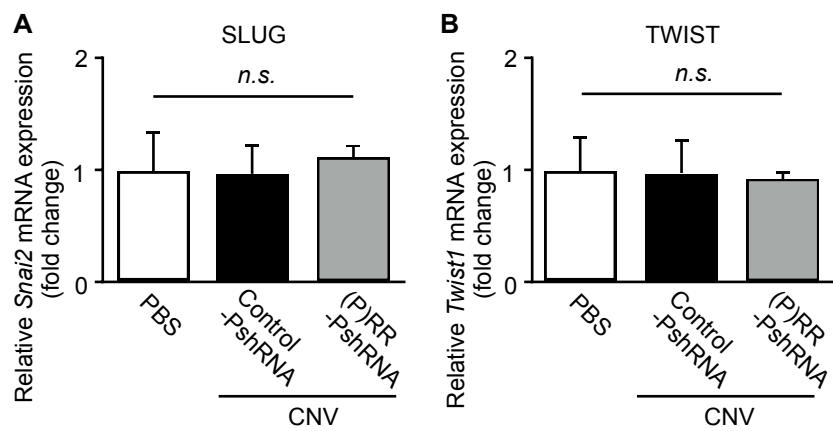
**Ye Liu, Atsuhiro Kanda, Di Wu, Erdal Tan Ishizuka, Satoru Kase, Kousuke Noda, Atsuhiro Ichihara, and Susumu Ishida**

## Supplemental information



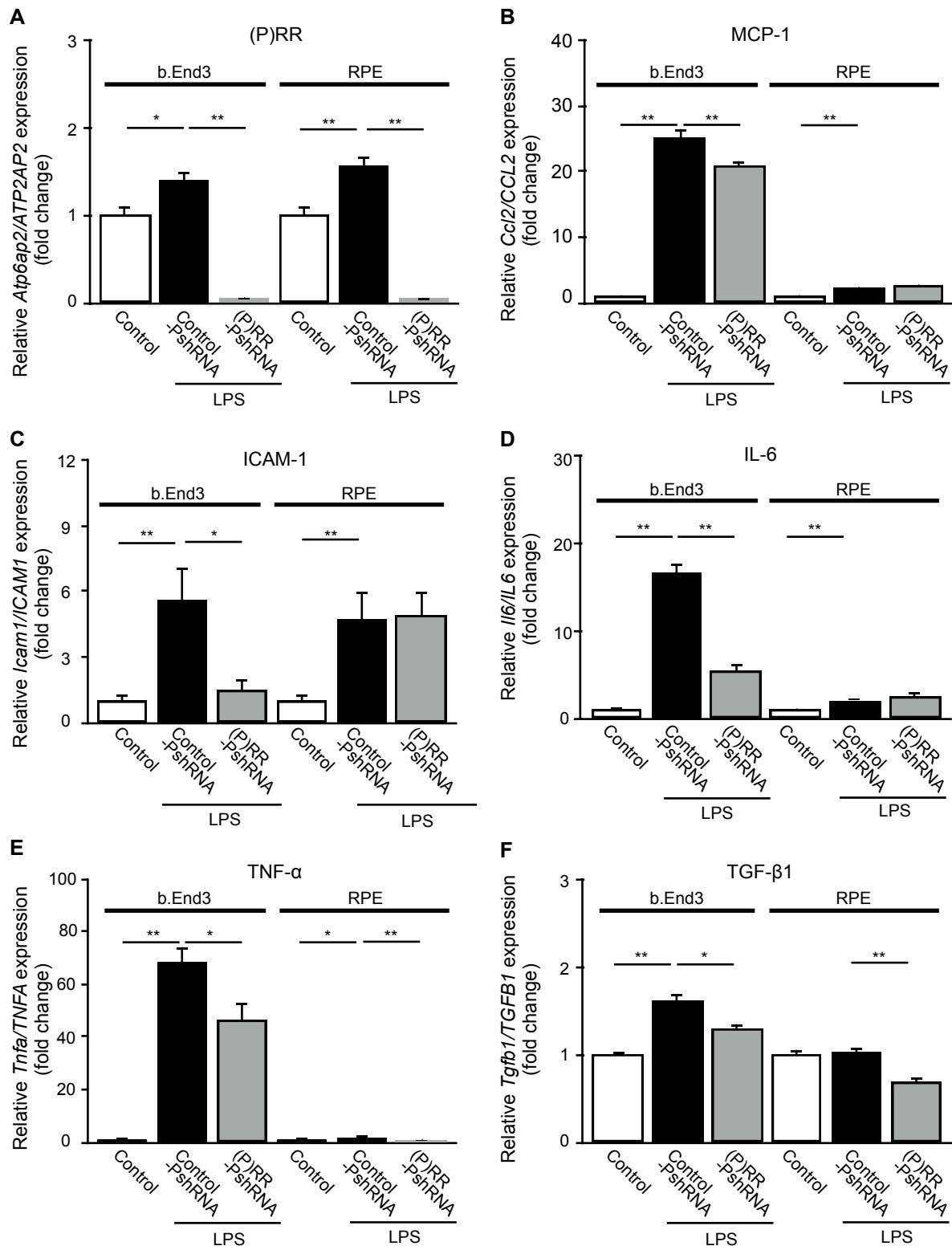
**Figure S1. Suppression of subretinal fibrosis at 21 days after laser photocoagulation by (P)RR-PshRNA**

(A, B) Representative micrographs of subretinal fibrosis lesions (type I collagen, *red*) in the RPE-choroid flat mounts at post-laser day 21 from mice treated with 100 pmol control-PshRNA (A) or (P)RR-PshRNA (B). Scale bar, 50 μm. (C) Quantification analysis of the size of subretinal fibrosis [control-PshRNA = 13,695 ± 4,566 μm<sup>2</sup>, (P)RR-PshRNA = 6,381 ± 2,236 μm<sup>2</sup>]. \*\**p* < 0.01 (*n* = 5).



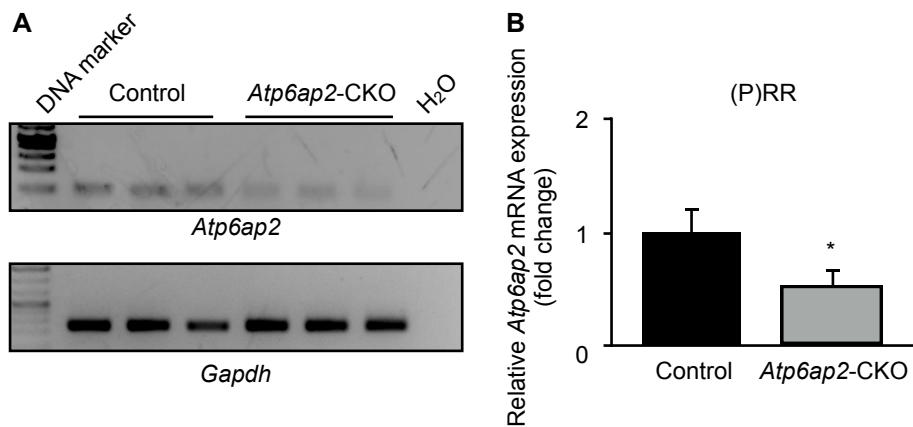
**Figure S2. Neither SNAI2 nor TWIST1 is associated with CNV induction in mice**

(A, B) Relative mRNA expression levels of *Snai2* (A) and *Twist1* (B) in the RPE-choroid complex from untreated normal mice (control) and CNV mice treated with 100 pmol control-PshRNA or (P)RR-PshRNA. n.s., not significant (n = 6).



**Figure S3. Blockade of inflammatory responses by (P)RR-PshRNA in LPS-stimulated endothelial cells and RPE cells**

(A-E) Gene expression levels of inflammatory molecules *Atp6ap2/ATP6AP2* (A), *Ccl2/CCL2* (B), *Icam1/ICAM1* (C), *Il6/IL6* (D), *Tnfa/TNFA* (E) and profibrotic cytokine *Tgfb1/TGFB1* (F) in 1 nM control-PshRNA or (P)RR-PshRNA transfected mouse microvascular endothelial cells stimulated with 10 ng/ml LPS for 8 hours and 1 nM control-PshRNA or (P)RR-PshRNA transfected human RPE cells stimulated with 10 ng/ml LPS for 12 hours. \* $p < 0.05$ , \*\* $p < 0.01$  ( $n = 6$ ).



**Figure S4. Gene expression of (P)RR/*Atp6ap2* in the RPE-choroid complex of (P)RR/*Atp6ap2*-CKO mice**

(A) Gene expression analysis of (P)RR/*Atp6ap2* in the RPE-choroid complex of control and (P)RR/*Atp6ap2*-CKO mice. *Gapdh* was used as an internal control. (B) Quantification analysis of the relative expression of (P)RR/*Atp6ap2*. \* $p < 0.05$  ( $n = 3$ ).

**Table S1. Primer sequences used in quantitative RT-PCR**

Target gene	Sequence
Mouse	
(P)RR/Atp6ap2	forward 5'- CCTCATTAGGAAGACAAGGACTATCC -3' reverse 5'- GGGTTCTCGCTTGTTCGC -3'
Ccl2	forward 5'- TTGGCTCAGCCAGATGCA -3' reverse 5'- CCTACTCATTGGGATCATCTTGC -3'
Icam1	forward 5'- CCTGTTCTGGCTCTGAAG -3' reverse 5'- GTCTGCTGAGACCCCTTG -3'
Il6	forward 5'- CACAGAGGATACCACCTCCAAACA -3' reverse 5'- TCCACGATTCCCAGAGAAACA -3'
Tnfa	forward 5'- GGTGCCTATGTCAGCCTCTT -3' reverse 5'- CGATCACCCCCGAAGTCAGTA -3'
Emr1	forward 5'- CTTGGCTATGGCTTCCAGTC -3' reverse 5'- GCAAGGAGGACAGAGTTATCGTG -3'
Tgfb1	forward 5'- CAGTGGCTAACCAAGGAGAC -3' reverse 5'- ATCCCGTTGATTCCACGTG -3'
Acta2	forward 5'- TCTGTAAGGCCGGCTTG -3' reverse 5'- TGTCCCATTCCCACCATCA -3'
Colla1	forward 5'- TGACTGGAAGAGCGGAGAGT -3' reverse 5'- GACGGCTGAGTAGGGAACAC -3'
Fn1	forward 5'- GTCAGTGTCTCCAGTGTCTAC -3' reverse 5'- TGGCTTGCTGCCAATCAGT -3'
Snai1	forward 5'- CACACGCTGCCTTGTGTCT -3' reverse 5'- GGTCAGCAAAGCACGGTT -3'
Snai2	forward 5'- CAGCGAACTGGACACACACA -3' reverse 5'- ATAGGGCTGTATGCTCCCGAG -3'
Twist1	forward 5'- GGACAAGCTGAGCAAGATTCA -3' reverse 5'- CGGAGAAGGCGTAGCTGAG -3'
Gapdh	forward 5'- AGGTCGGTGTGAACGGATTG -3' reverse 5'- TGTAGACCATGTAGTTGAGGTCA -3'
Human	
(P)RR/ATP6AP2	forward 5'- AGGCAGTGTCAATTGTCACC -3' reverse 5'- GCCTTCCCTACCATATACACTC -3'
CCL2	forward 5'- CGCCTCCAGCATGAAAGTCT -3' reverse 5'- ATGAAGGTGGCTGCTATG -3'
ICAM1	forward 5'- GCAAGCTCCCAGTCAAATGCAAAC -3' reverse 5'- TGTCTACTGACCCCAACCTTGATG -3'
IL6	forward 5'- CCACTCACCTCTCAGAACG -3' reverse 5'- CATCTTGAAAGGTTCAGGTTG -3'
TNFA	forward 5'- ACTTTGGAGTGATCGGCC -3' reverse 5'- GCTTGAGGGTTGCTACAAC -3'
TGFB1	forward 5'- GCCCTGGACACCAACTATTG -3' reverse 5'- CGTGTCCAGGCTCAAATG -3'
GAPDH	forward 5'- CCTGGCCAAGGTCAATCCATG -3' reverse 5'- GGAAGGCCATGCCAGTGAGC -3'