

**LCN2 deficiency mitigates the neuroinflammatory damage following acute
glaucoma**

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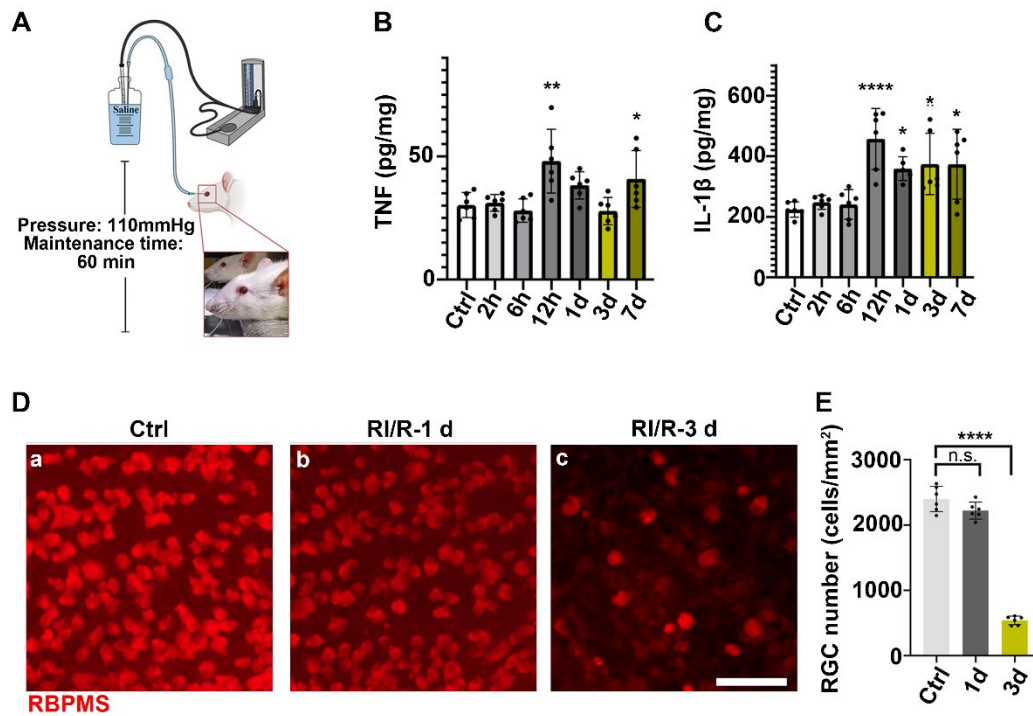


Figure S1. Inflammatory response occurred prior to RGC loss following RI/R injury. **A:** Diagram of RI/R model. **B, C:** ELISA showed the protein levels of TNF and IL-1 β in rat retinae following RI/R injury, compared to the control group (n = 6). **D:** Immunofluorescence images showed the number of RBPMS positive-RGCs in rat retinae following RI/R injury, compared to the control group. **E:** Bar graph depicted the number of RBPMS positive-RGCs per mm² in each group (n = 6). Data in **B, C, E** were represented as mean \pm SD; * p < 0.05, ** p < 0.01, **** p < 0.0001, n.s.: no significance, (compared with the control group using one-way analysis of variance). Bar= 100 μ m in **D**.

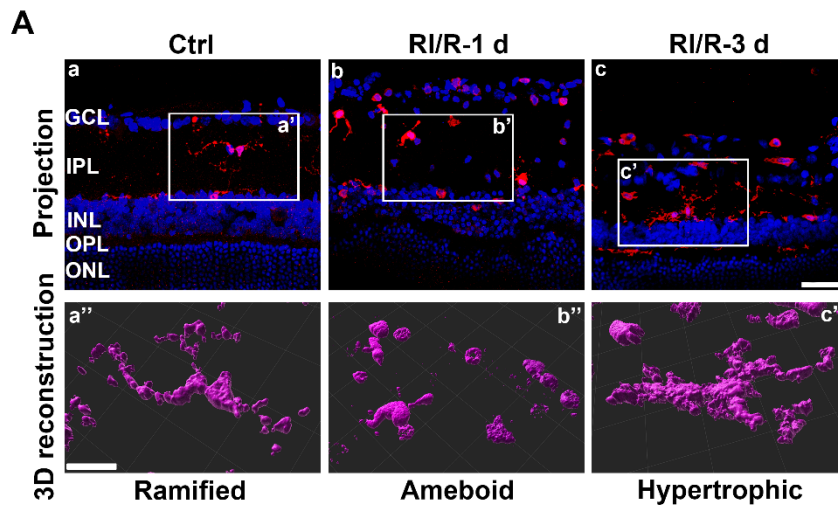


Figure S2. Dynamic changes in IBA1 positive cell morphology following RI/R. A: Representative 3D reconstruction of retinal IBA1-positive cell morphology following RI/R. Ramified shape characterized by the small cell body volume and long thin processes, which enabled microglia to monitor the surrounding environment. Ameboid morphology characterized by round sphere shapes and few processes, which was associated with microglia/macrophage phagocytosis and migration during disease and injury. Hypertrophic morphology characterized by enlarged cell body (compared to ramified shape) and fewer, shorter hypertrophic processes, which represented a post-injury morphology.

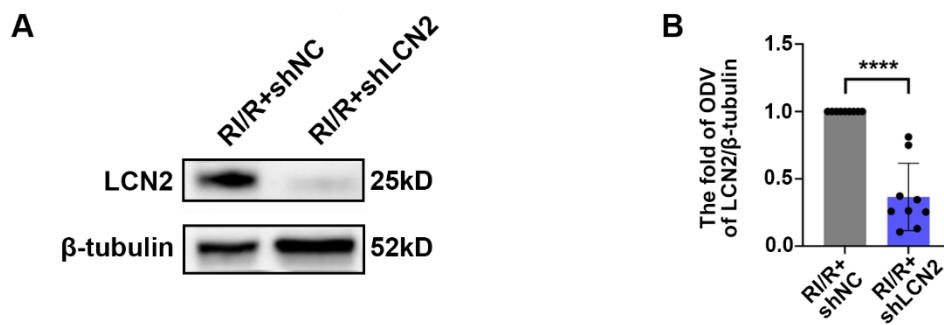


Figure S3. LCN2 shRNA-AAV knocked down retinal LCN2 expression following RI/R. **A:** Western blotting showed the change of LCN2 expression in rat retinae with the LCN2 shRNA-AAV treatment following RI/R. **B:** Bar graphs depicted the fold of ODV of LCN2 compared to the RI/R+shNC group (n = 9). Data in **B** were represented as mean \pm SD; **** $p < 0.0001$ (compared with the RI/R+shNC group using Student t-test).

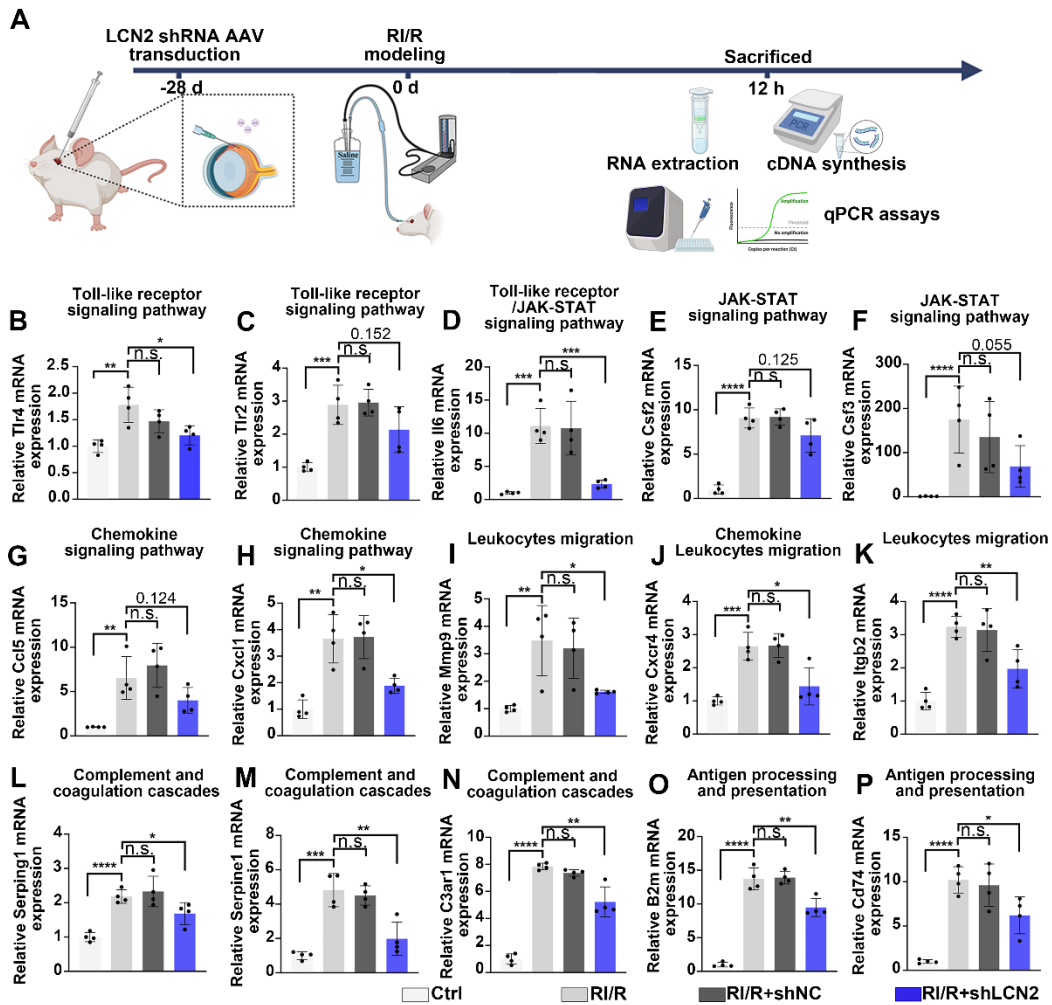


Figure S4. qPCR verification of key genes affected by *Lcn2* knockdown following RI/R. **A:** Diagram represented workflow of qPCR analysis. **B-P:** Bar graphs depicted the relative expression of target mRNA in each group (n = 4). Data in **B-P** were represented as mean \pm SD; * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$, n.s.: no significance, (compared with the RI/R group using Student t-test).

Table S1. Primer sequences for qPCR analysis

Gene	Forward	Reverse
<i>Tlr4</i>	5'-TTGCTGCCAACATCATCCAGGAAG-3'	5'-CAGAGCGGCTACTCAGAAACTGC-3'
<i>Tlr2</i>	5'-TCTGGAGTCTGCTGTGCCCTTC-3'	5'-GGAGCCACGCCACATCATTTC-3'
<i>Il6</i>	5'-ACTTCCAGCCAGTTGCCTTCTTG-3'	5'-TGGTCTGTTGTGGGTGGTATCCTC-3'
<i>Csf2</i>	5'-AATGACATGCGTGTCTGGAGAAC-3'	5'-TGGTGAGGTTGCCCCGTAGAC-3'
<i>Csf3</i>	5'-GGCTCTTCCTCTACCAAGGTCTCC-3'	5'-AGATGGTGGTGGCAAAGTTGTTCG-3'
<i>Ccl5</i>	5'-GACACCACTCCCTGCTGCTTTG-3'	5'-CTCTGGGTTGGCACACACTTGG-3'
<i>Cxcl1</i>	5'-GCAGACAGTGGCAGGGATTAC-3'	5'-AGTGTGGCTATGACTTCGGTTTGG-3'
<i>Mmp9</i>	5'-CTCCTGGTGCTCCTGGCTCTAG-3'	5'-GCTGTGTGTCCGTGAGGTTGG-3'
<i>Cxcr4</i>	5'-AAGCAAGGATGTGAGTTCGAGAGC-3'	5'-CCGAGGAAGGCGTAGAGGATGG-3'
<i>Itgb2</i>	5'-TTGCAGCAGAAGGACGGAAACG-3'	5'-ATGACCAGGAGGAGGACACCAATC-3'
<i>Serping1</i>	5'-GGCGGAGAACACCAACCACAAG-3'	5'-TTGGCACTCAAGTAGACGGCATTG-3'
<i>Serpine1</i>	5'-GCGTCTTCCTCCACAGCCATTC-3'	5'-TGTCTCTGTTGGATTGTGCCGAAC-3'
<i>C3ar1</i>	5'-GCCTTGAGAGAAGTTGTGCTACCC-3'	5'-AATGCCCTTTACGGACTGTCTTGC-3'
<i>B2m</i>	5'-ACCGTGATCTTTCTGGTGCTTGTC-3'	5'-ACACGTAGCAGTTGAGGAAGTTGG-3'
<i>Cd74</i>	5'-TCCTGGTGGCTCTGCTCTTGG-3'	5'-TCATGCGAAGGTTCTCCAGTTGC-3'
<i>Gapdh</i>	5'-AGTGCCAGCCTCGTCTCATA-3'	5'-GACTGTGCCGTTGAACTTGC-3'

Table S2. Demographic data and clinical characteristics of patients.

	non-glaucomatous	glaucoma
n	16	16
median age; years	65 (49-74)	69 (51-77)
female sex	56.3% (9)	62.5% (10)
disease type	age-related cataract	acute primary angle closure glaucoma
diabetes	-	-
hypertension	-	-