

Supplemental Information

Leptin and IL-6 Family Cytokines Synergize to Stimulate Müller Glia Reprogramming and Retina Regeneration

Xiao-Feng Zhao, Jin Wan, Curtis Powell, Rajesh Ramachandran, Martin G. Myers Jr and Daniel Goldman

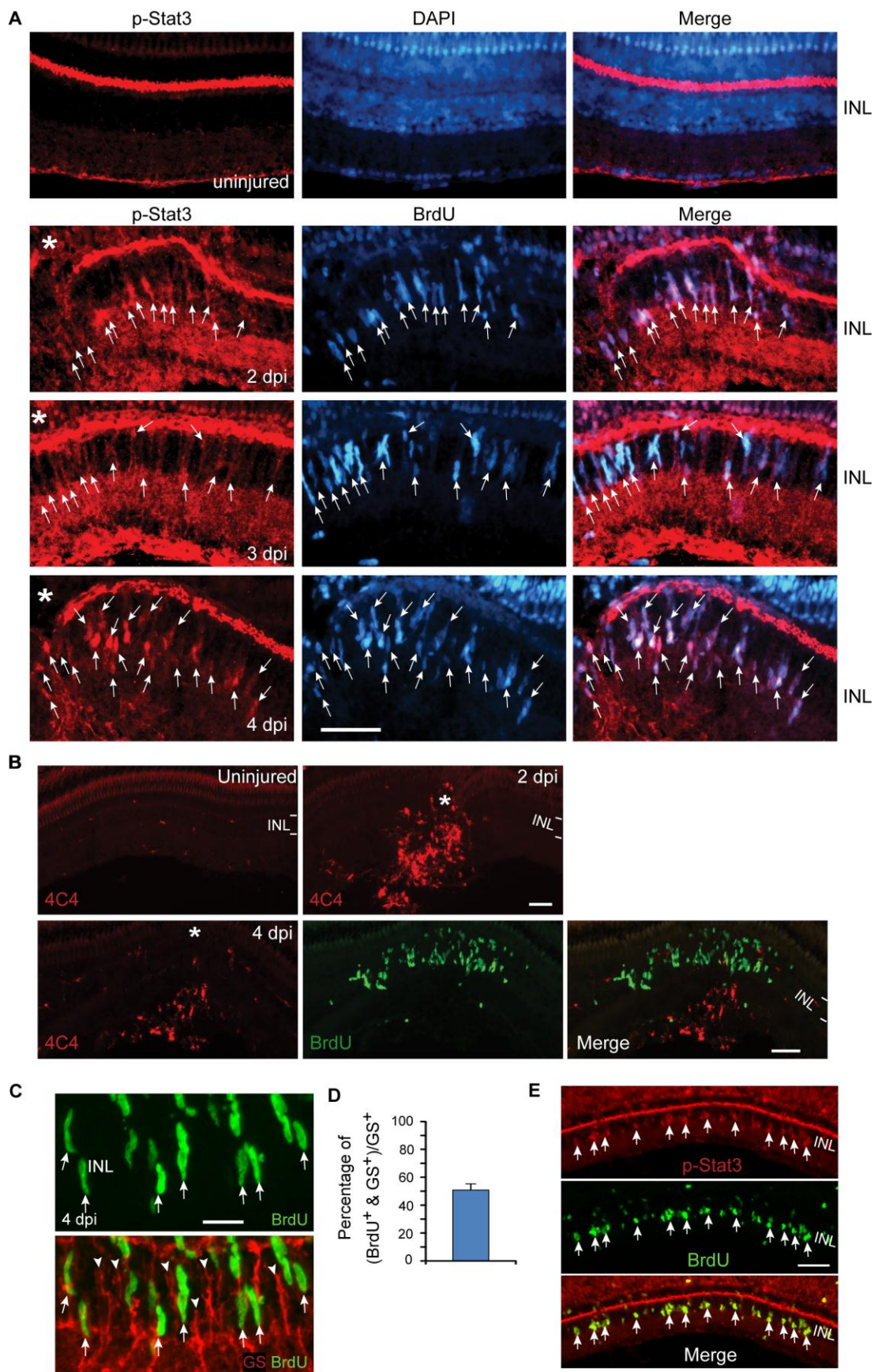


Figure S1. Injury-dependent activation of the Jak/Stat3 signaling pathway. Related to

Figure 1. (A) p-Stat3 (red) and BrdU (blue) immunofluorescence shows p-Stat3 is induced in BrdU⁺ MG-derived progenitors at the injury site at 2-4 dpi; n=3. Arrows point to p-Stat3⁺/BrdU⁺ double labelled cells. (B) 4C4 immunofluorescence (red) shows microglia, diffusely scattered throughout the uninjured retina, accumulate at the injury site, but do not proliferate (lack BrdU co-labeling, green). The BrdU⁺ cells (green) are MG-derived progenitors confined to the INL. (C, D) Immunofluorescence shows ~50% of glutamine synthetase (GS)⁺ MG (red) incorporate BrdU⁺ 4 days after a 30 min exposure to UV light; n=3. Arrows point to BrdU⁺/GS⁺ double labelled cells. Arrowheads point to BrdU/GS⁺ quiescent MG. (E) Immunofluorescence shows p-Stat3 is restricted to BrdU⁺ MG-derived progenitors 4 days after a 30 min exposure to UV light. Asterisks in (A) point to the injury site (needle poke). Scale bar, 20 μ m (C); 50 μ m (A,E). INL, inner nuclear layer.

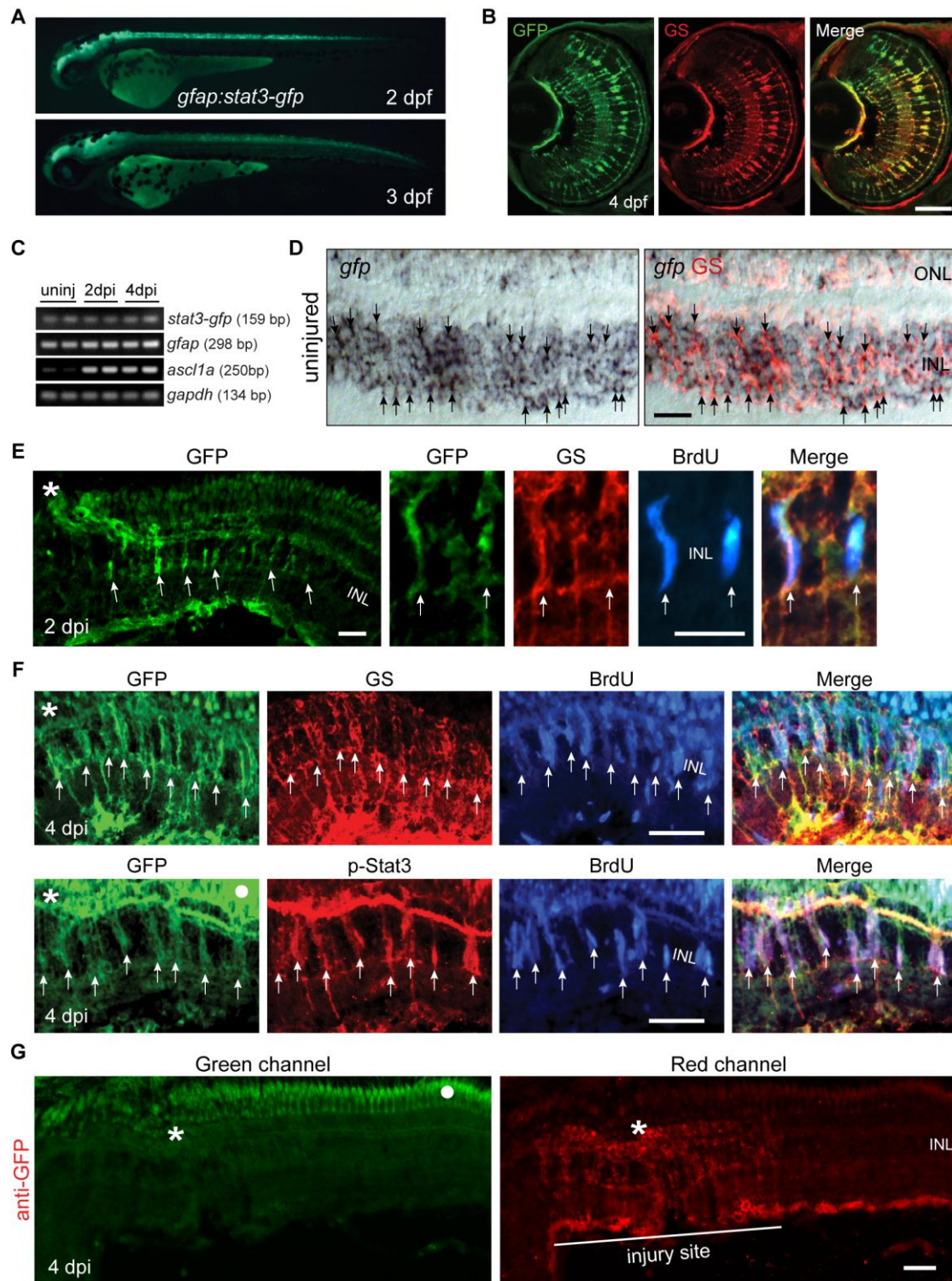


Figure S2. Stat3-GFP expression in developing and adult *gfap:stat3-gfp* fish. Related to Figure 1. (A) Stat3-GFP is expressed throughout brain and spinal cord in live *gfap:stat3-gfp* larva at 2 and 3 dpf. (B) Immunofluorescence on retinal sections shows co-localization of Stat3-GFP with glutamine synthetase (GS)⁺ MG at 4 dpf. (C) Whole retina RT-PCR showing constitutive *gfap* mRNA expression in the uninjured and injured adult retina, while *ascl1a* mRNA was induced after retinal injury. (D) *In situ* hybridization and immunofluorescence shows expression of *gfap* RNA in GS⁺ MG in adult uninjured retina

(arrows). (E) Immunofluorescence shows Stat3-GFP expression co-localizes with $GS^+/BrdU^+$ MG-derived progenitors localized to the injury site at 2 dpi in the adult retina. (F) Immunofluorescence shows co-localization of Stat3-GFP with $GS^+/p-Stat3^+/BrdU^+$ MG-derived progenitors at 4 dpi in the adult retina. White dot indicates autofluorescence unique to the green channel. Asterisks mark the injury site (needle poke) and arrows point to MG-derived progenitors. (G) Anti-GFP immunofluorescence on a retinal section prepared at 4 dpi from *gfap:stat3-gfp* fish using a secondary antibody coupled to a red fluor shows autofluorescence in the ONL when viewed in the green channel (left-hand panel, dot marks autofluorescence) that is not evident in the red channel (right-hand panel). Asterisk marks the injury site. Scale bar, 50 μm (B, D-G). INL, inner nuclear layer; dpf, days post fertilization.

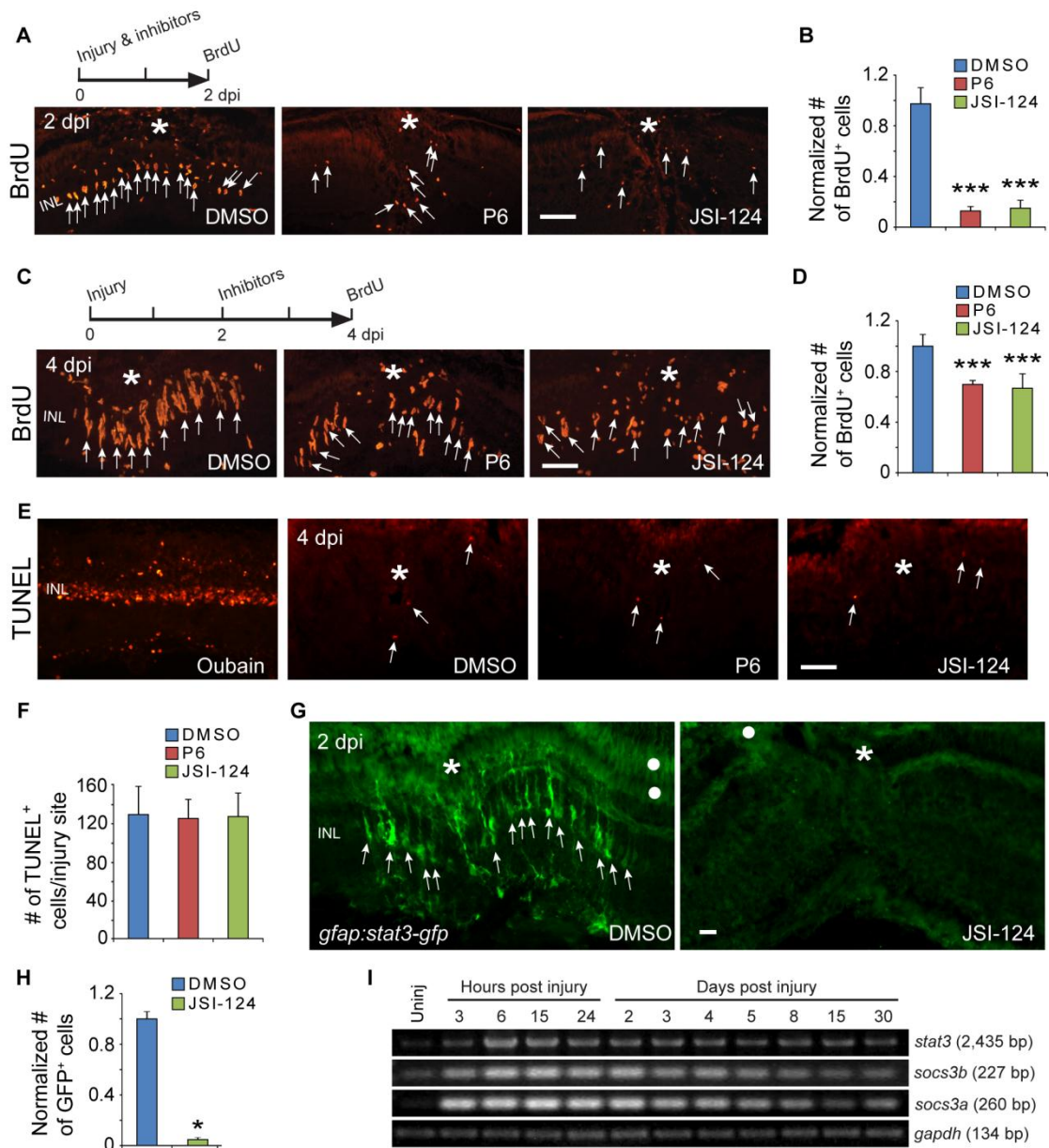


Figure S3. Jak/Stat signaling regulates proliferation of MG-derived progenitors and Stat3-GFP expression. Related to Figure 2. (A, B) BrdU immunofluorescence on retinal sections shows that exposing fish to Jak inhibitors P6 or JSI-124 from 0-2 dpi, inhibits progenitor proliferation at 2 dpi; ***P<0.001, n=4. (C, D) BrdU immunofluorescence shows that exposing fish to Jak inhibitors P6 or JSI-124 from 2-4 dpi, inhibits progenitor proliferation at 4 dpi; ***P<0.001, n=4. (E, F) TUNEL staining for apoptotic cells at 4 dpi in retinas treated with DMSO, P6 or JSI-124. Oubain was used as a positive control. In panel F, n=3. (G, H) GFP immunofluorescence shows that exposing *gfap:stat3-gfp* fish to Jak inhibitors JSI-124 from 0-2 dpi, inhibits Stat3-GFP expression; *P<0.05, n=3. (I) RT-PCR analysis of retinal *stat3* and *socs3* gene expression at various times after injury. Asterisks mark the injury sites (needle poke) and arrows point to MG-derived progenitors. White dot marks regions with autofluorescence. Error bars, s.d. Scale bar, 20 μ m (A, C, E, G). INL, inner nuclear layer.

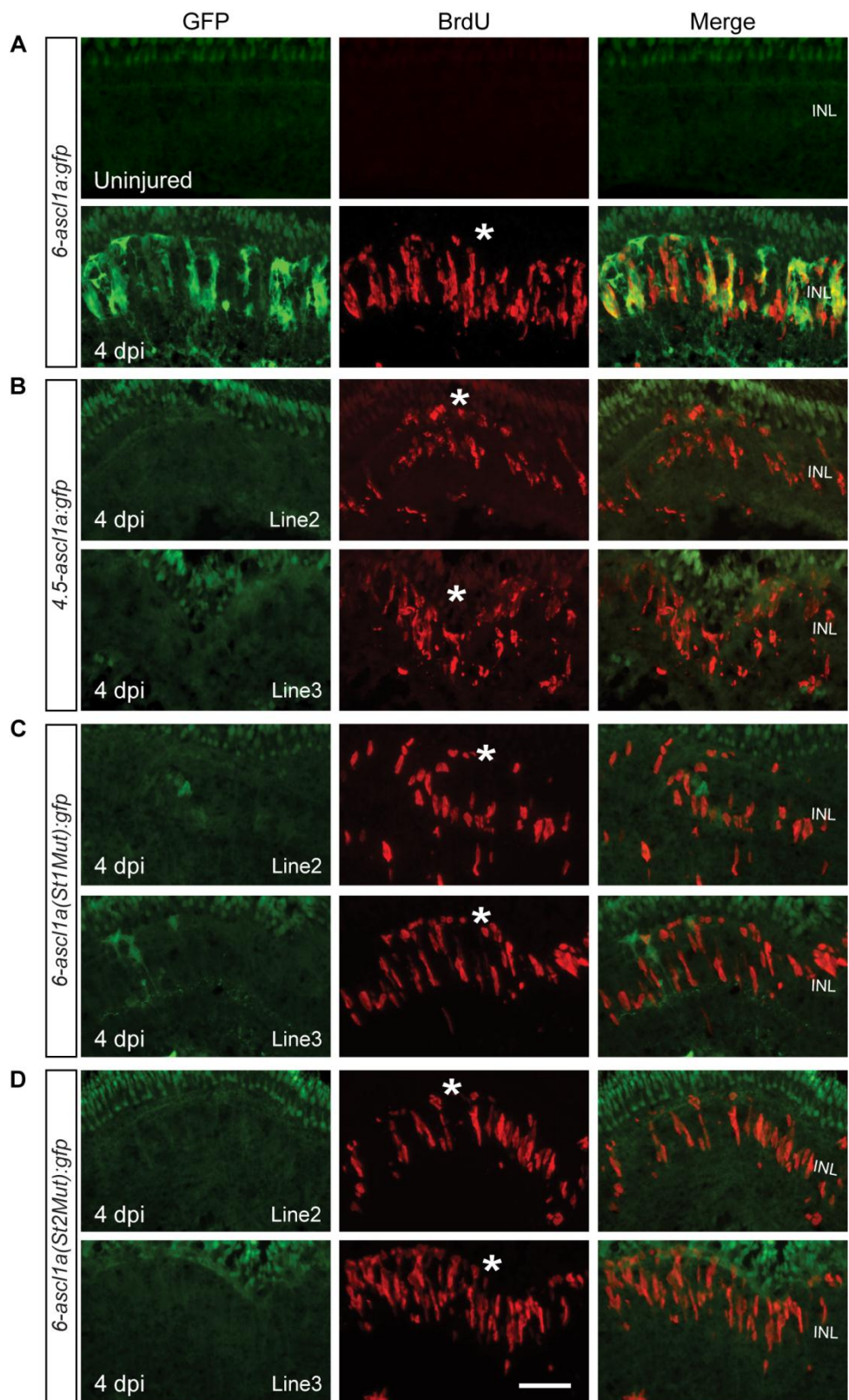


Figure S4. Jak/Stat3 signaling mediates injury-dependent activation of the *ascl1a* promoter. Related to Figure 3. (A) GFP and BrdU immunofluorescence co-localize beneath the injury site in *6-ascl1a:gfp* fish at 4 dpi. (B) A distal 1.5 kb *ascl1a* promoter fragment is necessary for injury-dependent transgene induction. (C, D) Consensus Stat3 sites located within this distal 1.5 kb region are necessary for injury-dependent transgene expression. BrdU⁺ immunofluorescence identifies the injury site and a normal regenerative response. The asterisks mark the injury sites. See Figure 3 for promoter schematic. Scale bar, 50 μ m. INL, inner nuclear layer.

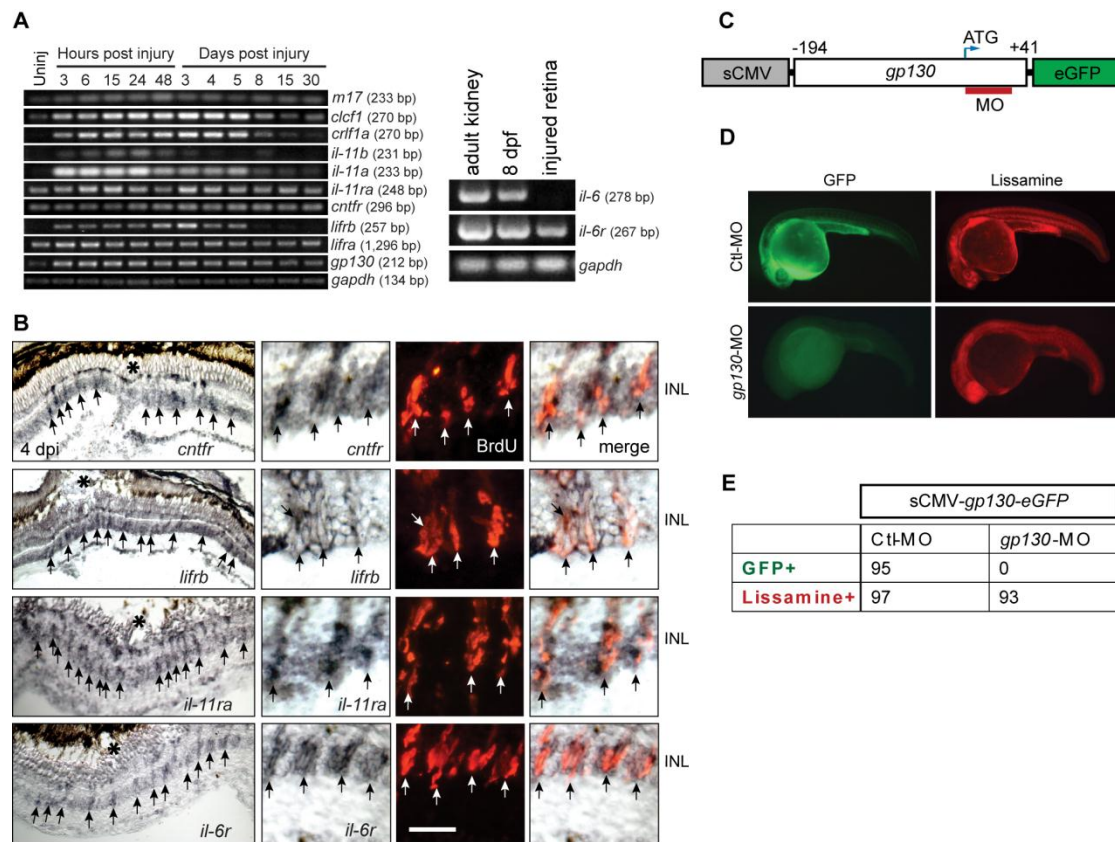


Figure S5. Genes encoding IL-6 family cytokines are expressed in MG-derived progenitors upon retinal injury. Related to Figure 4. (A) RT-PCR analysis of retinal gene expression at various times after injury. *il-6* is expressed in adult kidney and 8 dpf larva, but undetectable in the injured retina. (B) *In situ* hybridization shows induction of *il-6* family member genes in BrdU⁺ MG-derived progenitors at the injury site. The asterisks mark the injury sites and the arrows point to MG-derived progenitors. Scale bars, 20 μ m. (C) Diagram of sCMV:*gp130-egfp* reporter in which a fragment of *gp130* DNA, containing the MO target sequence, is in-frame with the *egfp* coding sequence and under control of the sCMV promoter. (D) Injection of the sCMV:*gp130-egfp* reporter together with either lissamine-tagged control (Ctl) MO or *gp130*-targeting MO into one cell stage of zebrafish embryos and examined by fluorescence 24 hours later. Red fluorescence shows embryos received lissamine-tagged MOs. (E) Quantification of GFP-expressing embryos at 24 hpf. No GFP was detected in the *gp130*-targeting MO injected group, while GFP was readily detected in the control group. Similar results were obtained in 3 independent experiments. dpf, days post fertilization ; INL, inner nuclear layer.

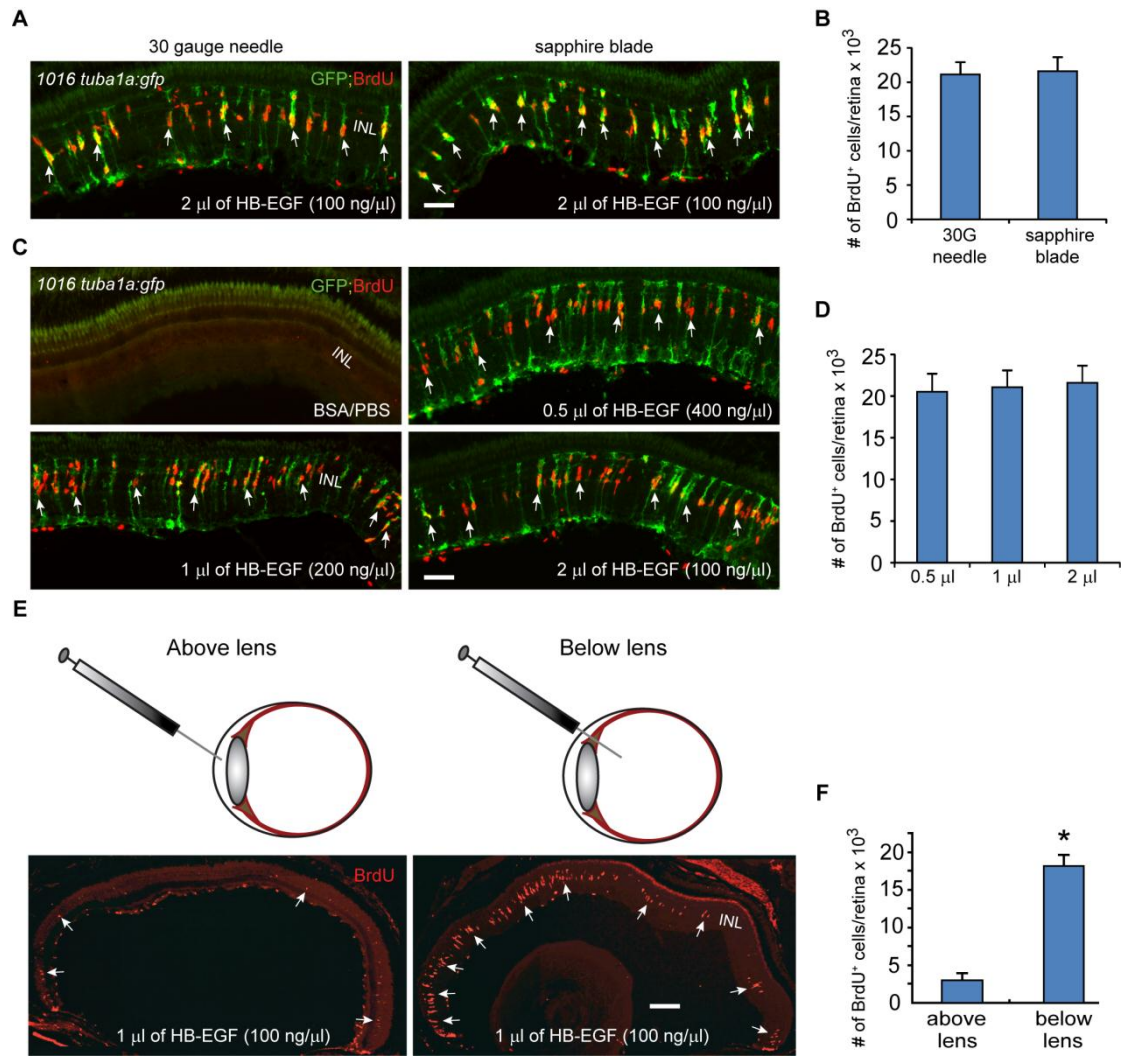


Figure S6. HB-EGF stimulates MG reprogramming and proliferation in the uninjured retina. Related to Figure 4. (A) A corneal puncture with a 30 gauge needle or a corneal incision made with a sapphire blade was followed by intravitreal injection of recombinant HB-EGF into the uninjured eye of *1016tuba1a:gfp* fish. HB-EGF stimulated GFP expression and BrdU incorporation in MG throughout the retina's inner nuclear layer. No GFP expression or BrdU incorporation is observed in PBS/BSA-injected eyes. (B) Quantification of BrdU⁺ cells in (A) shows similar effect of intravitreally delivered HB-EGF on MG proliferation when the cornea was either punctured with a needle or cut with a sapphire blade; n=3 per group. (C) A sapphire blade was used to make a small incision in the cornea and 200 ng of HB-EGF in 0.5 to 2 µl volumes was intravitreally injected into uninjured eyes of *1016tuba1a:gfp* fish. (D) Quantification of BrdU⁺ cells following intravitreal delivery of different HB-EGF volumes reveals a similar effect on MG proliferation regardless of injection volume; n=3 per group. (E, F) HB-EGF must be delivered below the lens to stimulate MG proliferation; *P<0.05, n=3. Arrows point to MG-derived progenitor. Error bars, s.d. Scale bar, 50 µm (A, C); 150 µm (E). INL, inner nuclear layer.

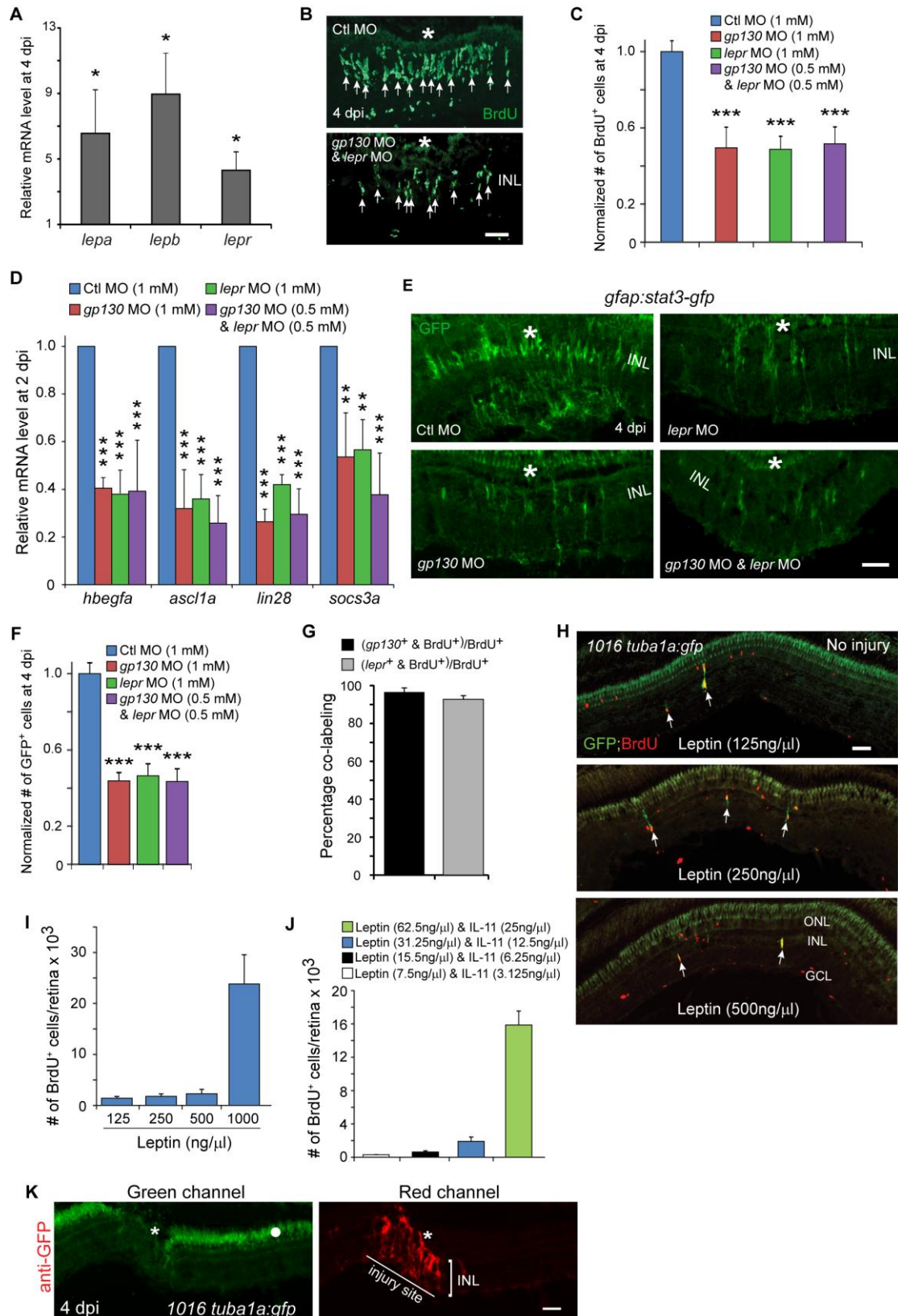


Figure S7. Leptin signaling regulates the generation of MG-derived progenitors. Related to Figure 5. (A) qPCR quantifying induction of gene expression in FACS purified GFP⁺ MG-derived progenitors from the retina of *1016tuba1a:gfp* fish at 4 dpi in comparison

to that in FACS purified GFP⁺ MG from the uninjured retina of *gfap:gfp* fish; *P<0.05, n=3. (B) BrdU immunofluorescence shows that Leptin receptor and Gp130 knockdown inhibits the generation of BrdU⁺ MG-derived progenitors at 4 dpi. Asterisks mark the injury sites and the arrows point to MG-derived progenitors. (C) Quantification of the effects Gp130 and Leptin receptor knockdown have on progenitor formation (Figure 4C; Figure 5C); ***P<0.001, n=4. (D) qPCR showing Gp130 and Leptin receptor knockdown, individually or in combination, inhibits injury-dependent induction of reprogramming genes at 2 dpi; **P<0.01, ***P<0.001, n=4. (E) Knockdown of Gp130 inhibits the expression of Stat3-GFP at 4 dpi. (F) Quantification of the number of GFP⁺ cells in the retina at 4 dpi following Gp130 or Leptin knockdown; ***P<0.001, n=4. Error bars, s.d. (G) Quantification of the number of BrdU⁺ cells expressing *gp130* or *lepr* RNA in the retina at 4 dpi; n=3. Error bars, s.d. (H) Effects of different Leptin concentrations on the generation of BrdU⁺ MG-derived progenitors and transgene expression in the uninjured retina of *1016tuba1a:gfp* fish. The arrows point to MG-derived progenitors. (I, J) Quantification of the effect different Leptin (I) or Leptin/IL-11 (J) concentrations have on progenitor formation in the uninjured retina; n=3. Error bars, s.d. (K) GFP immunofluorescence using a secondary antibody coupled to a red fluor shows autofluorescence in the green channel (left-hand panel, dot) that is localized to the ONL and pigment epithelium, while the red channel shows GFP expression restricted to the injury site (right-hand panel, asterisk). Scale bars, 20 μm (B); 50 μm (E, H, K). ONL, outer nuclear layer; INL, inner nuclear layer; GCL, ganglion cell layer.

Table S1. Primer sequences for PCR. Related to Experimental Procedures.

Gene	Sequence 5' - 3'
<i>lepa</i>	F, TTTCCAGCTCTCCGCTCAACC R, CGGCGTATCTGGTCAACATGC
<i>lepb</i>	F, CATTGCTCGAACCACCATCAGC R, TCTTTATGCACCGGGGTCTCG
<i>lepr</i>	F, CAGTACGAGCTGCAATTCAAGG R, TAAAATGCGCCAGAAGTCTGG
<i>m17</i>	F, CTTGATTGCCGTTCAAGTTAGTGC R, TTAAGATGCGCTCCGATTCAAGT
<i>clcf1</i>	F, GAAAGTTGGTCAGGTTGCTGTGC R, CATAAGTCCACACGTGTTGCTGC
<i>crlf1a</i>	F, GGGATTCTGGGATCTAGGAAAGC R, TCCTTGAAGAACCTGGTTGCG
<i>il-11a</i>	F, CTCCTCATCGCTGCTTCTCTCG R, TTGCGAAGTCACTGGCTCTGC
<i>il-11b</i>	F, GCTAACAGTGTGCGCTGACTCC R, CTGTAGTTCAGTGAGGGCAGGG
<i>il-11ra</i>	F, GTTGGACTGTTGGTTTTGTTGG R, TGGATTGTGGGTAATGAAGGC
<i>cntfr</i>	F, ACACCATCACCGACGCCTATGC R, GAAGCTCACACATCACATGATGG
<i>lifra</i>	F, AAGCCGTCTCCACACAGTCTGG R, TTCCCCCATTCTGCTTTCC
<i>lifrb</i>	F, TCACAGTTGACCAGATGCTTGC R, ATGTGGGTTTTCTGAGGTGGG
<i>gp130</i>	F, AATGAAGTTCGCCGATGGAGAGG R, CGTCTTCCTTGGGCATTTCCG
<i>il-6</i>	F, GCTATTCCTGTCTGCTACACTGG R, TGAGGAGAGGAGTGCTGATCC
<i>il-6r</i>	F, TCAGCTCCTGAGACAACACTGC R, AAACGGCATAGTCTGTTTCCC
<i>stat3</i>	F, ACACATTAAGGGGTCAACATGGCCCAG R, GCATTTCCGGCAGGTGTCCATATCCGAG
<i>socs3a</i>	F, CACTAACTTCTCTAAAGCAGGG R, GGTCTTGAAGTGGTAAAACG
<i>socs3b</i>	F, GAAAACCTCCAAGATTGAGTCG R, TACTATGCGTTACCATGGCG
<i>hbegfa</i>	F, ATGTCTGACCATCATTTGGCCTCC R, ACCATTCAGCTTGCTGTGCC
<i>ascl1a</i>	F, ATCCGCGCGCTGCAGCAGCTTCTGGACG

	R, CGAGTGCTGATATTTTTAAGTTTCCTTTTAC
<i>lin28</i>	F, TAACGTGCGGATGGGCTTCGGATTTCTGTC R, ATTGGGTCCTCCACAGTTGAAGCATCGATC
<i>gfp</i>	F, CAGCAGAACACCCCATC R, CTCGTCCATGCCGAGAGT
<i>gfap</i>	F, GGATGAGATCCAGATGCTGAAGG R, CAGATCCTTCCTCTCCGTAGTGG
<i>gapdh</i>	F, ATGACCCCTCCAGCATGA R, GGCGGTGTAGGCATGAAC

Table S2. Kd, ED₅₀ and intravitreal concentration of recombinant mammalian cytokines. Related to Experimental Procedures.

Cytokine	Kd and ED₅₀	Estimated concentration after intravitreal injection*
Recombinant human Leptin	Kd ~0.06-3nM (Verkerke et al., PLoS One 9:e94843, 2014; Uotani et al., Diabetes 48:279, 1999). ED ₅₀ ~0.4-2 ng/ml in a cell proliferation assay (http://www.rndsystems.com/Products/398-LP).	62.5 ng = 0.195 μM 1 μg = 3.125 μM
Recombinant rat CNTF	Two classes of receptor binding sites: high affinity site, Kd ~1pM and low affinity site Kd ~1nM (Wong et al., J Biol Chem 270:313, 1995). ED ₅₀ ~1 ng/ml in a cell proliferation assay (Gearing et al, Proc Natl Acad Sci USA 91:1119, 1994).	25 ng = 54.8 nM 100 ng = 219 nM
Recombinant human IL-6	Two classes of receptor binding sites: high affinity site, Kd ~10pM and low affinity site, Kd ~1nM (Yamsaki et al., Science 241:825, 1988). ED ₅₀ ~10 ng/ml in a cell proliferation assay (Hibi et al., 63:1149, 1990).	100 ng = 211 nM
Recombinant mouse IL-11	Two classes of receptor binding sites: high affinity site, Kd ~0.3-0.8nM and low affinity site, Kd ~10nM; ED ₅₀ ~5 ng/ml in a cell proliferation assay (Hilton et al., EMBO J. 13:4765, 1994).	25 ng = 65.3 nM 100 ng = 2.6 μM

*Cytokine concentrations in the vitreous of injected eyes was estimated as previously described using a 20 μl vitreal volume (Fimbel et al., J Neurosci 27:1712, 2007; Ritchey et al., Exp Eye Res 99:1, 2012; Raymond et al., J Neurobiol 19:431, 1988).