

Description of Additional Supplementary Files

Supplemental Dataset 1: snRNA/ATAC-Seq and scRNA-Seq data for LD and NMDA samples.

Supplemental Dataset 2: Genes, peaks, and motifs selectively enriched in LD vs. NMDA samples (snRNA/ATAC-Seq+scRNA-Seq).

Supplemental Dataset 3: Regulatory relationships among TFs selectively active in LD vs. NMDA samples.

Supplemental Dataset 4: Genes selectively expressed in each of major retinal cell type in snRNA-seq and scRNA-Seq injury datasets.

Supplemental Dataset 5: snRNA/ATAC-Seq and reanalyzed scRNA-Seq data for developing retinal samples.

Supplemental Dataset 6: Genes, peaks, and motifs selectively enriched in injury vs. development samples (snRNA/ATAC-Seq+scRNA-Seq).

Supplemental Dataset 7: Regulatory relationships among TFs selectively active in injury vs. development samples.