## Supplementary Materials



Supplementary Figure 1. Transverse cryosectioning of mouse retinal punch illustrates the method used to isolate the outer retina. A punch of retina was flatmounted, embedded in OCT cutting medium, and flash frozen, then mounted in a cryostat and sectioned at $20 \mu \mathrm{~m}$ sections through retinal layers. Layers were then stained for PKC-alpha (a marker of bipolar cells in the inner nuclear layer) and cone arrestin (a marker of photoreceptors in the outer retina). Labeling shows successful isolation of outer retinal tissue from inner retinal cells, which was then used for amplification of libraries. RPE was peeled away prior to sectioning.


Supplementary Figure 2. Scatter plots track variants across rounds of selection. Scatterplots illustrate the behavior of individual variants over all rounds of selection for the $\sim$ Ancestral- 7 mer library and the 588-Loopswap library. Variants overrepresented in the original library are colored blue. Variants that had the greatest fold increase in representation in the final round of selection are shown in magenta. Variants that were overrepresented in the original library and increased significantly in representation over rounds of selection are colored orange. Black dots in AAV2-7mer scatter plots indicate the variant NHP\#9.

| Round | NHP ID | Age/Weight/Sex | Libraries injected | Amount Virus Injected | Notes |
| :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | V002278 | Approx. 7 years (age unknown at import) 5.98 kg / Male | Loop Swap Ancestral-7mer | $2 \times 10^{11} \mathrm{vg}$ per library; $100 \mu 1$ volume. |  |
| 2a | V002262 | Approx. 7 years (age unknown at import) 6.35 kg / Male | Recovered variants from round 1 AAV2-7mer | $\begin{gathered} 2.5 \times 10^{11} \mathrm{vg} \text { ONL } \\ 2.5 \times 10^{11} \mathrm{vg} \text { RPE } \\ 5 \times 10^{10} \mathrm{vg} \text { AAV2-7mer; } 100 \\ \mu \mathrm{l} \text { volume. } \end{gathered}$ | No variants were PCR amplified following injection from this round, no obvious immune response noted. |
| 2b | V002148 | Approx. 7 years (age unknown at import) $6.48 \mathrm{~kg} /$ Male | Recovered variants from round 1 AAV2-7mer | $\begin{gathered} 1.3 \times 10^{11} \mathrm{vg} \text { ONL } \\ 1.3 \times 10^{11} \mathrm{vg} \text { RPE } \\ 5 \times 10^{10} \mathrm{vg} \text { AAV2- } 7 \mathrm{mer} ; 100 \\ \mu \mathrm{l} \text { volume. } \end{gathered}$ | Repeat of previous round |
| 3 | V002265 | 8 years 5 months $4.92 \mathrm{~kg} /$ Male | Recovered variants from round 3 | $\begin{gathered} 4.3 \times 10^{12} \mathrm{vg} \text { ONL } \\ 3.7 \times 10^{12} \mathrm{vg} \text { RPE; } 100 \mu \mathrm{l} \\ \text { volume. } \end{gathered}$ | Error prone PCR conducted No adverse events |
| 4 | V002540 | 6 years 9 months $4.59 \mathrm{~kg} /$ Male | Recovered variants from round 4 | $\sim 1 \times 10^{12}$ vg per library; 100 $\mu$ volume. | No adverse events |
| 5 | V002861 | 6 years 6 months $6.60 \mathrm{~kg} /$ Male | Recovered variants from round 5 | $\begin{gathered} 2.4 \times 10^{12} \mathrm{vg} \text { ONL } \\ 6.3 \times 10^{12} \mathrm{vg} \text { RPE; } 100 \mu 1 \\ \text { volume. } \end{gathered}$ | No adverse events |
| $\begin{aligned} & \text { GFP- } \\ & \text { barcode } \end{aligned}$ | V002361 | 9 years 5 months $6.00 \mathrm{~kg} /$ Male | Barcoded individual variants | $\sim 1 \times 10^{10} \mathrm{vg}$ each variant; 100 $\mu \mathrm{l}$ volume. | Both eyes injected with GFP-BC library. Hyphema in left eye resolved in 12 days |
| Variant validation | 106 | 9 years 5 months $14.5 \mathrm{~kg} /$ Male | 7 m 8 and NHP9 | $\begin{gathered} \sim 1.5 \times 10^{12} \mathrm{vg} \\ 7 \mathrm{~m} 8-\mathrm{pR} 1.7-\mathrm{GFP}+ \\ 1.5 \times 10^{12} \mathrm{vg} \\ 7 \mathrm{~m} 8-\mathrm{SNCG}-\mathrm{tdTomato} ; 100 \\ \mu \mathrm{l} \text { volume. } \\ \mathrm{OR} \\ \sim 1.5 \times 10^{12} \mathrm{vg} \\ \text { NHP\#9-pR1.7-GFP }+ \\ 1.5 \times 10^{12} \mathrm{vg} \\ \text { NHP\#9-SNCG-tdTomato; } \\ 100 \mu 1 \text { volume. } \end{gathered}$ | No adverse events |
| Variant validation | 735 | 17 years <br> Male | NHP26 in one eye | $\begin{gathered} \sim 5 \times 10^{10} \mathrm{vg} \\ \text { NHP\#26-scCAG-GFP; } 100 \\ \mu \text { l volume. } \end{gathered}$ | No adverse events |

Supplemental Table 1. Summary of the rounds of selection performed in primates. The table indicates the age and weight of the primates injected, the virus and titer injected at each round, and notes on the rounds of selection completed. ONL refers to virus libraries recovered from ONL samples. RPE refers to virus libraries recovered from RPE samples, which were processed in parallel. Round 2 b was a repeat of the $2^{\text {nd }}$ round of selection, which did not result in PCR amplification of variants.

| Primer | Sequence |
| :---: | :---: |
| SDM1 | GACCTTAATCACAATCTTTTAAAACCCCGGCATGGCGGCT |
| SDM2 | GGCTCGTGGACAAGTAAAGGGATTACCTCGGA |
| Neb Genomic_F | GTAAGGGTCTGCTCCATTGCCACTT |
| Neb Genomic_R | CTAAATCAAAAAAGAGTGAAAAGTTAGGAGG |
| IFA_F | TGGCTCGTGGACAAGGTAAGGGTCTGCTCCATTGC |
| IFA_R | CTCCGAGGTAATCCCCTAAATCAAAAAAGAGTGAAAAGTT |
| HindIII_F1 | GACGTCAGACGCGGAAGCTTC |
| NotI_R1 | GGTTTATTGATTAACAAGCGGCCG |
| Ascl_R1 | TGGCGGACTTTATAGGCGCG |
| Spel_R1 | GCCCAGTTCGAATAGCGAGT |
| LS588_Forward adapter | AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCTNNNNNGTTCTGTATCTACCAACCTCCA |
| LS588_rev_index1 | CAAGCAGAAGACGGCATACGAGATCGTGATGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGACCATGCCTGGAAGAACGCC |
| LS453 Forward adapter | AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCTNNNNNATCGACCAGTACCTGTATTACT |
| LS453_rev_index1 | CAAGCAGAAGACGGCATACGAGATCTGATCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGGTCCCGAATGTCACTCGCTC |
| Anc_Forward adapter | AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCTNNNNNCCTGCAGTCGTCTAACACCGC |
| Anc rev index 1 | CAAGCAGAAGACGGCATACGAGATCTCTACGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTAAGGCTCCCTGGCTGTTGAC |
| 5_Forward adapter | AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCTNNNNNATGGCCACCAACAACCAGAGC |
| 5_rev_index1 | CAAGCAGAAGACGGCATACGAGATATCAGTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGAGGTTGTACGTGCCGGTCGC |
| 4 Forward adapter | AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCTNNNNNACCTACCTGGCGGTGACCAGA |
| 4_rev_index1 | CAAGCAGAAGACGGCATACGAGATTAGTTGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTCCACGGTCGGCAGGTTGCTG |
| LS588_rev_index2 | CAAGCAGAAGACGGCATACGAGATACATCGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGACCATGCCTGGAAGAACGCC |
| LS588 rev index3 | CAAGCAGAAGACGGCATACGAGATGCCTAAGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGACCATGCCTGGAAGAACGCC |
| LS588_rev_index4 | CAAGCAGAAGACGGCATACGAGATTGGTCAGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGACCATGCCTGGAAGAACGCC |
| LS588 rev index5 | CAAGCAGAAGACGGCATACGAGATCACTGTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGACCATGCCTGGAAGAACGCC |
| LS588_rev_index6 | CAAGCAGAAGACGGCATACGAGATATTGGCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGACCATGCCTGGAAGAACGCC |
| LS588_rev_index 7 | CAAGCAGAAGACGGCATACGAGATGATCTGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGACCATGCCTGGAAGAACGCC |
| LS588 rev index8 | CAAGCAGAAGACGGCATACGAGATTCAAGTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGACCATGCCTGGAAGAACGCC |
| LS453_rev_index2 | CAAGCAGAAGACGGCATACGAGATAAGCTAGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGGTCCCGAATGTCACTCGCTC |
| LS453_rev_index3 | CAAGCAGAAGACGGCATACGAGATGTAGCCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGGTCCCGAATGTCACTCGCTC |
| LS453 rev index4 | CAAGCAGAAGACGGCATACGAGATTACAAGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGGTCCCGAATGTCACTCGCTC |
| LS453_rev_index5 | CAAGCAGAAGACGGCATACGAGATTTGACTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGGTCCCGAATGTCACTCGCTC |
| LS453_rev_index6 | CAAGCAGAAGACGGCATACGAGATGGAACTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGGTCCCGAATGTCACTCGCTC |
| LS453 rev index 7 | CAAGCAGAAGACGGCATACGAGATTGACATGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGGTCCCGAATGTCACTCGCTC |
| LS453_rev_index8 | CAAGCAGAAGACGGCATACGAGATGGACGGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGGTCCCGAATGTCACTCGCTC |
| Anc_rev_index 2 | CAAGCAGAAGACGGCATACGAGATGCGGACGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTAAGGCTCCCTGGCTGTTGAC |
| Anc rev index 3 | CAAGCAGAAGACGGCATACGAGATTTTCACGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTAAGGCTCCCTGGCTGTTGAC |
| Anc_rev_index 4 | CAAGCAGAAGACGGCATACGAGATGGCCACGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTAAGGCTCCCTGGCTGTTGAC |
| Anc_rev_index 5 | CAAGCAGAAGACGGCATACGAGATCGAAACGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTAAGGCTCCCTGGCTGTTGAC |
| Anc_rev index6 | CAAGCAGAAGACGGCATACGAGATCGTACGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTAAGGCTCCCTGGCTGTTGAC |
| Anc_rev_index 7 | CAAGCAGAAGACGGCATACGAGATCCACTCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTAAGGCTCCCTGGCTGTTGAC |
| Anc_rev_index8 | CAAGCAGAAGACGGCATACGAGATGCTACCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTAAGGCTCCCTGGCTGTTGAC |
| 5 rev index2 | CAAGCAGAAGACGGCATACGAGATGCTCATGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGAGGTTGTACGTGCCGGTCGC |
| 5_rev_index 3 | CAAGCAGAAGACGGCATACGAGATAGGAATGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGAGGTTGTACGTGCCGGTCGC |
| 5_rev_index 4 | CAAGCAGAAGACGGCATACGAGATCTTTTGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGAGGTTGTACGTGCCGGTCGC |
| 4 rev index 2 | CAAGCAGAAGACGGCATACGAGATCCGGTGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTCCACGGTCGGCAGGTTGCTG |
| 4_rev_index 3 | CAAGCAGAAGACGGCATACGAGATATCGTGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNTCCACGGTCGGCAGGTTGCTG |
| 4 rev index 4 | CAAGCAGAAGACGGCATACGAGATTGAGTGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNTCCACGGTCGGCAGGTTGCTG |
| 2-7mer Forward adapter | AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCTNNNNNTCTACCAACCTCCAGAGAGG |
| rev_index1 | CAAGCAGAAGACGGCATACGAGATCGTGATGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev index2 | CAAGCAGAAGACGGCATACGAGATACATCGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev_index3 | CAAGCAGAAGACGGCATACGAGATGCCTAAGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev_index4 | CAAGCAGAAGACGGCATACGAGATTGGTCAGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev index5 | CAAGCAGAAGACGGCATACGAGATCACTGTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev_index6 | CAAGCAGAAGACGGCATACGAGATATTGGCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev_index 7 | CAAGCAGAAGACGGCATACGAGATGATCTGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev index8 | CAAGCAGAAGACGGCATACGAGATTCAAGTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev_index9 | CAAGCAGAAGACGGCATACGAGATCTGATCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev_index 10 | CAAGCAGAAGACGGCATACGAGATAAGCTAGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev index11 | CAAGCAGAAGACGGCATACGAGATGTAGCCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| rev_index 12 | CAAGCAGAAGACGGCATACGAGATTACAAGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNGTTGACATCTGCGGTAGCTG |
| F_adapter_GFPBC | AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCTNNNNNGGCCATCAAGCTTATCGATACC |
| R adapter GFPBC | CAAGCAGAAGACGGCATACGAGATCGTGATGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCTNNNNNCTGATCAGCGAGCTCTAGTCG |
| NHP GAPD F | TGCACCACCAACTGCTTAGC |
| NHP GAPD R | GGCATGGACTGTGGTCATGAG |
| K9 GAPDH F | TGTCCCCACCCCCAATGTATC |
| K9 GAPDH R | CTCCGATGCCTGCTTCACTACCTT |

Supplemental Table 2. Primers used in the study.

