Supplementary Information

Lipid nanoparticles with PEG-variant surface modifications mediate genome editing in the mouse retina

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Supplementary Figure 1. Intracellular uptake of Cy-5 tagged LNP variants. (A-E) Representative confocal images of 661W cone cells treated with Cy-5 tagged LNP variants at 24 hours treatment. All the cells were treated with Cy5-tagged LNP dose equivalent to 100 ng/well of mRNA cargo (scale bar: 25 μ m). (F) Relative cellular uptake of Cy-5 tagged LNP variants obtained by quantitative estimation of confocal images against unmodified LNP. (G) Quantification of cellular uptake using flow cytometry. (H) Gating strategy used during cellular uptake study for untreated cells and same strategy was used for other LNP variant treated cells. All data are presented as Mean ± SD. An ordinary one-way ANOVA, with Tukey's correction for multiple comparisons test was used for comparisons. Unmodified LNP values were normalized, and fold changes were compared with LNP variants. ns-not significant, **p<0.01. All representative figures and graphs have n=3 replicates. Source data are provided as a Source Data file.



Supplementary Figure 2. *In-vitro* cell viability assay. 661w cone cells were treated with LNP variants at a 100 and 200 ng fluc mRNA dose. (A) 24 hours post-treatment and (B) 48 hours post-treatment. All graphs have n=5 replicates. Source data are provided as a Source Data file.



Supplementary Figure 3. Endosomal escape of LNP variants. Representative images of the Gal8-GFP reporter cells, 24 hours after *cre* mRNA-loaded LNP variant treatments. All representative figures from n = 3 replicates.



Supplementary Figure 4. Photoreceptor colocalization study in Ai9 mice. (A-D)

Representative confocal images showing tdTomato expression in the RPE (LNP) and photoreceptors (LNPa and LNPz), which are colocalized with recoverin. Arrow indicates tdTomato expression (pink: RPE, yellow: cone photoreceptor nuclei, green: rod photoreceptor nuclei). RPE: retinal pigment epithelium, ONL: outer nuclear layer, INL: inner nuclear layer. All representative figures from n = 6 single eyes per group.



Supplementary Figure 5. Zeta potential value from mCherry encapsulated LNP variants measured by Malvern zeta sizer. n=3 LNPs per group. All Data are presented as Mean ± SD. An ordinary one-way ANOVA, with Tukey's correction was used for comparison with unmodified LNP. ns-not significant. Source data are provided as a Source Data file.

A) A*A*G*UAAAACCUCUACAAAUGGUUUUAGAGCUAGAAAUAGCAAGUUAA AAUAAGGCUAGUCCGUUAUCAACUUGAAAAAGUGGCACCGAGUCGGUG CU*U*U*U



Supplementary Figure 6. CRISPR-Cas9 delivery and sgRNA modification in Ai9 mice. (A) sgAi9 guide with modifications highlighted. In the sgRNA structure, the asterisks (blue) and red font indicate the phosphorothioate bond and 2'-O-methyl ribonucleotides, respectively. **(B)** Representative confocal image showing outer nuclear thinning and disruption 7-days post-subretinal injection with MC3-Cas9-sgAi9-LNPx (n=6 single eyes). **(C-G)** Representative confocal images showing editing events post-LP01-Cas9-sgAi9-LNPx injection. **(C)** Retinal cross section highlighting RPE editing (n=6 single eyes). **(D)** Neural retina flatmount showing tdTomato fluorescence signal within the treated area of the retina (n=9 single eyes). **(E)** Retinal cross sections highlighting editing in the Müller glia and **(F-G)** photoreceptors (n=6 single eyes). O/S: outer segment, PR: photoreceptor, RPE: retinal pigment epithelium.

Α



Supplementary Figure 7. Illumina quality control data from NGS. (A) QScore distribution indicating high quality sequencing data (over 96% Q30 and above). (B) Percentage of bases with \geq Q30 plotted by cycle showing high accuracy of base call across the whole sequencing run (n=9).

Supplementary Table 1: Antibodies used for staining

S.N.	Primary Antibody	Host	Dilution
1.	Anti-recoverin (Cat. # AB5585)	Anti-rabbit	1:500
2.	Anti-mCherry (Cat. # NBP2-25157)	Anti-rabbit	1:250
3.	Visual Arrestin (E-3) (SC-166388)	Anti-mouse	1:100

S.N.	Secondary Antibody	Host	Dilution
1.	Alexa Fluor 700-IgG (Cat. # A21038)	Goat anti-rabbit	1:1000
2.	Alexa Fluor 594-IgG (H+L) (Cat. # A-21207)	Donkey anti-rabbit	1:500
3.	Alexa Fluor Plus 488-IgG (H+L) (Cat. # A21038)	Goat anti-mouse	1:200

Supplementary Table 2: Guide RNA, primers, and cycling conditions used for editing studies

S.N.	sgRNA Name	Sequence
1.	sgAi9	AAGUAAAACCUCUACAAAUG
2.	sgGFP	CUCGUGACCACCCUGACCUA

S.N.	Primer Name	Sequence (5' -> 3')
1.	Ai9_NGS_F1_F	ACACTCTTTCCCTACACGACGCTCTTCCGATCTATACGAAGTTATTCGCGATG
2.	Ai9_NGS_F2_F	ACACTCTTTCCCTACACGACGCTCTTCCGATCTGTTGTGGTTTGTCCAAAC
3.	Ai9_NGS_R	GACTGGAGTTCAGACGTGTGCTCTTCCGATCTTGTTTCAGGTTCAGGGGGAG

Supplementary Table 3: Thermocycling methods for 1st PCR for NGS library preparation

Step	Temperature	Time
Initial Denaturation	95.0 °C	3 min
30 Cycles: Denaturation Annealing Extension	98.0 °C 60.0 °C 72.0 °C	20 sec 20 sec 30 sec
Final Extension	72.0 °C	30 sec

Supplementary Table 4: Thermocycling methods for 2nd PCR for NGS library preparation

Step	Temperature	Time
Initial Denaturation	98.0 °C	2 min
12 Cycles: Denaturation Annealing Extension	98.0 °C 62.0 °C 72.0 °C	10 sec 20 sec 30 sec
Final Extension	72.0 °C	2 min