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## Efficient mouse genome engineering by CRISPR-EZ (CRISPR RNP Electroporation of Zygotes) technology

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### Abstract

CRISPR/Cas9 technology has transformed mouse genome editing with unprecedented precision, efficiency, and ease; however, the current practice of microinjecting CRISPR reagents into pronuclear-stage embryos remains rate-limiting. We thus developed CRISPR-EZ (**CRISPR RNP Electroporation of Zygotes**), an electroporation-based technology that outperforms pronuclear and cytoplasmic microinjection in efficiency, simplicity, cost, and throughput. In C57BL/6J and C57BL/6N mouse strains, CRISPR-EZ achieves 100% delivery of Cas9/sgRNA ribonucleoproteins (RNPs), facilitating indel mutations (insertions or deletions), exon deletions, point mutations, and small insertions. In a side-by-side comparison in the high-throughput KnockOut Mouse Project (KOMP) pipeline, CRISPR-EZ consistently outperformed microinjection. Here, we provide an optimized protocol covering single guide RNA (sgRNA) synthesis, embryo collection, RNP electroporation, mouse generation, and genotyping strategies. Using CRISPR-EZ, a graduate-level researcher with basic embryo manipulation skills can obtain genetically modified mice in 6 weeks. Altogether, CRISPR-EZ is a simple, economic, efficient, and high-throughput technology that is potentially applicable to other mammalian species.

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#### AUTHOR CONTRIBUTIONS

AJM conceived the original idea that leads to the development of the CRISPR-EZ technology, drafted the manuscript and methods, analyzed the raw experimental data, and produced all figures and tables. SC and AJM trained staff at MBP at UC-Davis to perform CRISPR-EZ experiments, optimized electroporation conditions and produced a video for all experimental procedures. SC conceived of the Tyrediting scheme for optimizing the CRISPR-EZ protocol, and drafted and revised the manuscript. BW contributed to sgRNA design, RNP preparation, genotype screening, data analysis, and manuscript preparation. JAW and KL contributed to study design, collaboration setup, and paper revision. LH contributed to experimental design, initiated and established the collaboration with MBP at UC-Davis, worked closely with AJM and SC to interpret the data, to draft and revise the manuscript.

**EDITORIAL SUMMARY:** CRISPR-EZ (**CRISPR RNP Electroporation of Zygotes**) achieves 100% delivery of Cas9/sgRNA ribonucleoproteins (RNPs) by zygote electroporation, enabling efficient indels, exon deletions, point mutations, and small insertions in mouse genome, and outperforming microinjection-based methods.

**TWEET:** CRISPR-EZ employs zygote electroporation to deliver RNPs for efficient mouse genome editing

#### COMPETING FINANCIAL INTERESTS

The authors declare no competing financial interests.

## Keywords

Mouse genome editing; CRISPR-Cas9; CRISPR-EZ; CRISPR; Cas9; sgRNA; ribonucleoprotein; RNP; zygote; embryo; electroporation; zygote electroporation

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## Introduction

The advent of CRISPR/Cas9 technology has greatly simplified and expedited *in vivo* genome editing in mice<sup>1–3</sup>. Standard practice involves pronuclear and/or cytoplasmic microinjection of *Cas9* mRNA/sgRNA into pronuclear-stage embryos<sup>4–8</sup>, but microinjection remains a rate-limiting step due to its slow, costly, and technically demanding nature. This procedure is often performed by experienced personnel at dedicated transgenic facilities whose technical expertise commands a premium rate. Furthermore, *Cas9* mRNA/sgRNA microinjection is prone to suboptimal editing efficiency and undesired mosaicism<sup>9–11</sup>, as the Cas9 protein must first be translated, folded, and complexed with sgRNAs prior to editing<sup>12–15</sup>.

To expedite genome editing *in vivo*, we developed CRISPR-EZ (**CRISPR RNP Electroporation of Zygotes**), an electroporation-based mouse genome engineering technology to overcome these limitations<sup>16</sup>. By delivering preassembled Cas9/sgRNA ribonucleoproteins (RNPs) into pronuclear-stage embryos, editing occurs rapidly and transiently, maximizing efficiency while minimizing mosaicism<sup>9–11</sup>. CRISPR-EZ completely bypasses microinjection by utilizing a series of electrical pulses to deliver RNPs with 100% efficiency. In both mouse strains tested, C57BL/6J and C57BL/6N, CRISPR-EZ vastly surpassed *Cas9* mRNA/sgRNA microinjection in editing efficiency, at approximately one quarter of the cost. CRISPR-EZ has now been successfully utilized by seven independent labs (personal communications), generating mice with editing schemes ranging from simple indels and exon deletions via the Non-Homologous End Joining (NHEJ) pathway, to precise point mutations and small insertions via the Homology-Directed repair (HDR) pathway (Fig. 1a). Other electroporation-based protocols have been developed by independent groups, with efficiencies exceeding that of microinjection<sup>9,17,18</sup>. These methods employ costly or proprietary equipment setups and were only tested on a limited number of editing experiments. In contrast, CRISPR-EZ uses commonly available reagents and equipment, requires only basic embryo handling skills, and has been comprehensively tested across a wide range of targets and editing schemes, making our protocol readily amenable to many investigators. Altogether, CRISPR-EZ is a powerful tool that stands to replace microinjection as the standard mouse genome editing method for most common editing schemes.

## Protocol overview

Using CRISPR-EZ, one can quickly test sgRNAs directly in cultured mouse embryos in 1–2 weeks, and subsequently generate edited mice in one month (Fig. 1b). This protocol can be broken into 7 major stages, including sgRNA design (Steps 1–2), sgRNA synthesis (Steps 3–15), superovulation and mating (Steps 16–19), embryo culture, collection and processing (Steps 20–28), RNP assembly and electroporation (Steps 29–34), embryo culture and

genotyping (Step 35A), and oviduct transfer (Step 35B). We have also included a video demonstrating all key procedures (Supplementary Video 1).

## Development of protocol

Development of CRISPR-EZ was inspired by the reported success of delivering *Cas9* mRNA/sgRNA into mouse and rat zygotes by electroporation<sup>19–22</sup>. However, the highly anionic properties of *Cas9* mRNA may explain inefficient pronuclear delivery<sup>23</sup>. In contrast, CRISPR-EZ achieves 100% delivery by electroporating the relatively less anionic *Cas9*/sgRNA RNPs<sup>23,24</sup>. More importantly, preassembled *Cas9*/sgRNA RNPs initiate editing in embryos more rapidly than *Cas9* mRNA, and their transient presence could reduce mosaicism and potential off-target effects<sup>9–11</sup>.

We developed CRISPR-EZ as a simple, high-throughput methodology that lowers the technical and financial barriers for mouse genome engineering. In a proof-of-principle editing experiment targeting the key pigment synthesis gene *Tyrosinase* (*Tyr*) (Fig. 2a, b, Supplementary Fig. 1a, b), we previously achieved 88% bi-allelic gene disruption and 42% HDR-mediated editing in C57BL/6J mice using a highly efficient sgRNA<sup>24</sup>. We also demonstrated that CRISPR-EZ could efficiently mediate a variety of editing schemes, including indels, precise mutations, exon deletions, and small insertions<sup>24</sup> (Fig. 1a). We have since further optimized CRISPR-EZ electroporation conditions to achieve better editing efficiency without impacting embryo viability (Fig. 2c-i, Supplementary Table 1), and expanded its application to the widely used C57BL/6N mouse strain (Fig. 2d, h, Supplementary Table 1). Under these optimized conditions (30V, 6 pulses, 3 ms), we achieved 100% bi-allelic *Tyr* editing in both mouse strains, as shown by the albino coat in all mice generated (Fig. 2c, d, g, h), as well as 62.5% HDR efficiency in the C57BL/6J strain (Supplementary Table 1).

We have performed side-by-side comparisons of CRISPR-EZ versus microinjection targeting the *Tyr* gene in the C57BL/6J strain<sup>24</sup>, as well as the *Sh3rf2* gene in the C57BL/6N strain (Fig. 3a, b, c, d). In both cases, CRISPR-EZ achieved markedly greater editing efficiency<sup>24</sup> (Fig. 3d). We then comprehensively compared 21 CRISPR-EZ and 27 microinjection-based knockout experiments integrated within the KnockOut Mouse Project (KOMP) pipeline, as part of their ongoing effort to knockout all protein-coding genes in mice. Remarkably, CRISPR-EZ exhibited on average ~3-fold increased editing efficiency over microinjection in C57BL/6N mice, while retaining comparable live birth rates (Fig. 3e, g, Supplementary Fig 2,3, Supplementary Table 2 and 3). Importantly, CRISPR-EZ efficiently generated edited founders for multiple genes that were refractory to editing by microinjection (Fig. 3f, Supplementary Fig. 3, Supplementary Tables 2 and 3). Based on these findings, the Mouse Biology Program at UC-Davis, a major operation center for KOMP, has adopted CRISPR-EZ as their primary method for knockout mouse generation. Altogether, CRISPR-EZ constitutes an economic, high-throughput, and highly efficient mouse genome engineering technology that stands to replace microinjection for a variety of editing strategies.

## Application of Protocol

CRISPR-EZ is broadly applicable to a wide range of mouse genome editing schemes for biomedical research. So far, CRISPR-EZ has been successfully employed in 34 editing projects across seven independent labs, generating edited mice with genomic deletions (n=28), indel mutations (n=3), and point mutations (n=3). Thus, CRISPR-EZ can replace microinjection for common editing schemes and multiple mouse strains.

CRISPR-EZ also provides a cost-effective means to test sgRNA designs *in vivo*. Despite recent advancements in sgRNA prediction algorithms, empirical testing remains necessary before *in vivo* experiments are attempted. Due to the high cost of microinjection, sgRNAs are frequently screened using immortalized or transformed mouse cell lines, whose editing efficiencies may differ from that of the embryo<sup>25</sup>. Using CRISPR-EZ, investigators can rapidly screen sgRNAs by directly genotyping electroporated embryos after culturing to the morula stage, thereby increasing confidence in obtaining the desired edited mice.

CRISPR-EZ may be a powerful tool for generating compound mouse genetic models to study complex genetic interactions or highly redundant gene families. Traditionally, mice harboring multiple edited alleles are generated through lengthy breeding of individually edited mice. With the advent of Cas9/sgRNA microinjection, it became possible to simultaneously edit multiple loci<sup>26</sup>, and CRISPR-EZ makes this even more accessible and efficient. Such approaches could be particularly useful when engineering multiple genetically linked loci, which are virtually impossible to obtain through breeding.

Given the similarities in preimplantation development among some mammalian species, CRISPR-EZ could potentially edit a much wider range of model organisms than the tested mouse strains. Notably, highly efficient porcine genome editing was recently demonstrated using a similar strategy<sup>27</sup>. The amenability of a specific mammalian zygote to CRISPR-EZ depends on its innate sensitivity to experimental manipulation, as well as the physical and biochemical properties of its zona pellucida and cellular membrane. Optimization will be necessary for each additional mouse strain or mammalian species. Given its high efficiency, CRISPR-EZ may be particularly suited for editing organisms whose embryos are scarce or difficult to obtain.

## Comparison with other methods

The current *de facto* standard for mouse genome editing is microinjection of CRISPR reagents into pronuclear-stage embryos. Edited mice are routinely generated by microinjection of *Cas9* mRNA/sgRNAs, and more recently, of Cas9/sgRNA RNPs<sup>28–30</sup>. It has been reported that Cas9/sgRNA RNPs are superior to *Cas9* mRNA/sgRNAs owing to their immediate editing activity upon delivery<sup>12,14,15,31,32</sup>. In spite of its popularity, microinjection possesses several caveats, including high cost, low throughput, and technical difficulty, all of which are addressed by CRISPR-EZ. In our studies, CRISPR-EZ outperforms microinjection of *Cas9* mRNA/sgRNAs for mouse genome engineering. Similar studies have demonstrated better mouse editing efficiency by CRISPR-EZ when compared with that of Cas9 RNP microinjection<sup>33</sup>.

While CRISPR-EZ efficiently performs a variety of simple editing schemes, its potential for more complex genome engineering, such as the insertion of fluorescent tags or conditional alleles, remains unclear, as we have only demonstrated electroporation of ssODN donors up to 162 nt<sup>24</sup>. Complex HDR-mediated editing schemes may require alternative approaches such as genetic engineering of embryonic stem cells (ESCs), microinjection of large double-stranded donors, or more recently, microinjection of synthesized long ssODNs<sup>34–37</sup>. As engineered ESCs can be clonally expanded and validated prior to mouse generation, this classic approach is still the preferred method for most sophisticated editing schemes.

In recent years, multiple electroporation-based methods have been utilized to deliver *Cas9* mRNA into mouse zygotes under both *ex vivo* and *in utero* conditions<sup>19–21</sup>. However, the reported efficiencies for these approaches were generally suboptimal (~10% for indels, for example<sup>21</sup>) compared to CRISPR-EZ, possibly due to inefficient *Cas9* mRNA delivery and/or delayed *Cas9* expression in the embryo<sup>38,39</sup>. Several other groups also reported success in mouse genome engineering using zygote electroporation of *Cas9* RNPs<sup>9,18,40</sup>. However, their specific experimental conditions were only tested on a limited number of editing experiments (no more than 1–2 targets per editing scheme were reported), and the variability across different targets, editing schemes and mouse strains, as well as high-throughput scalability were not clearly addressed<sup>9,18,40</sup>. Using CRISPR-EZ, we clearly demonstrated a 100% *Cas9* RNP delivery in the zygotes of two different mouse inbred strains. We also demonstrated the effectiveness, robustness and reliability of CRISPR-EZ in a high-throughput pipeline across 21 different knockout targets (Fig. 3e, g, Supplementary Figs. 2 and 3, Supplementary Tables 2 and 3), showing an average of 35% editing success rate among the liveborn pups. For NHEJ-mediated deletion schemes, CRISPR-EZ typically uses 60 embryos to generate ~3 edited founders in most experiments (Fig. 3e). This editing efficiency and robustness significantly exceeds that of microinjection-based technology in the same editing schemes (Figs. 2 and 3); other *Cas9* RNP electroporation protocols have not reported this level of robustness in a high throughput knockout pipeline. Finally, CRISPR-EZ is significantly more cost-effective compared to other *Cas9* mRNA or *Cas9* RNP electroporation methodologies, owing to its high editing efficiency, its technical simplicity, and its use of inexpensive, off-the-shelf equipment and consumables. Thus, CRISPR-EZ offers significant advantages over other microinjection-based and electroporation-based technologies, and can be employed as a standard technology for mouse editing experiments to generate indels, deletions, point mutations and small insertions.

### Level of expertise required

CRISPR-EZ requires basic mouse embryo manipulation skills and molecular biology training. To successfully perform CRISPR-EZ in cultured mouse embryos, researchers will perform PCR, *in vitro* transcription, RNA purification, and RNP assembly to generate functional *Cas9*/sgRNA complexes, and then perform surgeries to collect pronuclear-stage embryos from superovulated females for RNP electroporation. To test editing strategies, the researchers will culture pronuclear-stage embryos to morula for genotyping analysis. A graduate student could master these skills after 2–3 months of training. To generate edited mice, researchers will transfer electroporated embryos into the oviducts of pseudopregnant

mothers. This technically demanding procedure requires survival surgery and fine embryo manipulation skills and can also be learned by a graduate level researcher but will require considerably more hands-on experience than routine embryo manipulation. Researchers can utilize the service of a transgenic facility or acquire training to perform oviduct transfer independently. We refer researchers to “Manipulating the Mouse Genome” by Dr. András Nagy, if more experimental details are desired for this procedure<sup>41</sup>.

## Limitations

CRISPR-EZ is well suited for generating indels and exon deletions through the NHEJ pathway (Figs. 2 and 3, Supplementary Figs. 1–3). We demonstrated that CRISPR-EZ facilitates deletions up to ~2.6 kb in length (Supplementary Fig. 2a, Supplementary Table 3), which greatly exceeds the average length of a protein-coding exon (~170 bp)<sup>42</sup>. However, deletion efficiency is inversely correlated with deletion size (Supplementary Fig. 1c), such that engineering very large deletions could present a challenge, albeit this caveat is not unique to CRISPR-EZ.

Cas9/sgRNA RNPs can be co-electroporated with ssODNs to mediate precise sequence replacement via HDR. Using CRISPR-EZ, we successfully inserted a V5 tag (42 bp) into the endogenous *sox2* locus by co-electroporating a 162 nt ssODN<sup>24</sup>. Given the difficulty of synthesizing longer ssODNs, we have not yet investigated the maximal oligo length for delivery by CRISPR-EZ. A recent study has reported complex HDR-mediated editing in mice upon microinjection of Cas9/sgRNA RNPs along with long ssODN donors (~1000 nt)<sup>37</sup>. Hence, utilizing long ssODNs in CRISPR-EZ, if possible, would provide a promising strategy for complex genome editing. Currently, ESC genome editing and microinjection remain the preferred methodologies for complex mouse genome engineering.

While CRISPR/Cas9 has helped define an era of genome editing, concerns persist regarding the extent of Cas9 off-target effects<sup>43–45</sup>. We have not yet performed whole genome sequencing of edited mice generated by CRISPR-EZ or microinjection, and it is unclear if they differ in the extent of off-target effects. Engineered Cas9 variants may improve target specificity<sup>46,47</sup>, which have yet to be tested in the CRISPR-EZ system. Ultimately, one should exercise caution when interpreting the phenotype of CRISPR edited mice. To unambiguously demonstrate a causal relationship between genome modifications and the observed mouse phenotype, and to mitigate the concern of off-target artifacts, researchers may generate multiple edited mouse lines using independent sgRNA designs, and perform sufficient backcrossing to generate congenic mouse strains. When engineering mutant mice using classic ESC approaches, mutant phenotype is often validated using multiple ESC lines. Similar criteria are applied to transgenic mouse lines, where multiple founders are analyzed to definitively conclude the phenotype caused by the transgene. CRISPR editing in mice, either generated by CRISPR-EZ or by microinjection, may need to be subjected to the same level of scrutiny.

## Experimental Design

### Animal use authorization.

Live animals are used throughout this protocol. As a matter of caution and compliance, all appropriate authorizations must be acquired from institutional and/or federal regulatory bodies prior to performing this protocol. All mouse use, including but not limited to housing, breeding, production, sample collection for genotyping, and euthanasia, is in accordance with the Animal Welfare Act, the AVMA Guidelines on Euthanasia and are in compliance with the ILAR *Guide for Care and Use of Laboratory Animals*, and the UC Davis institutional animal care and use committee (IACUC) guidelines and policies. Our animal care and use protocol has been reviewed and approved by our IACUC for this project.

### sgRNA design.

Designing optimal sgRNAs is crucial for successful editing, as efficiency varies depending on Cas9 sequence preference, target specificity, genome topology, and cell type<sup>48–50</sup>. Several publicly-available design programs can be used to determine initial sgRNA candidates. Some examples include but are not limited to: Sequence scan for CRISPR (<http://crispr.dfc.harvard.edu/SSC/>), Gene Perturbation Platform (<http://portals.broadinstitute.org/gpp/public/analysis-tools/sgrna-design>), Chop-Chop (<https://chopchop.rc.fas.harvard.edu/>), CRISPR Design (<http://crispr.mit.edu/>). We recommend screening 2–3 sgRNA designs per target site directly in mouse embryos to ensure faithful *in vivo* editing (Fig. 1b). For exon deletions, we screen 2–3 sgRNAs for each cutting site and identify the paired sgRNA combination(s) with the best efficiency. Although we have had some success testing sgRNAs *in vitro* using Cas9-overexpressing mouse cancer cell lines, we have encountered cases where cell line results differ from those obtained from embryos (data not shown).

### sgRNA synthesis.

We recommend a rapid and cost-effective cloning-free method to generate large quantities of high quality *in vitro* transcribed sgRNAs<sup>51</sup> (Fig. 4). In this strategy, a unique oligo containing a T7 promoter and the guide sequence is annealed to a universal oligo containing the tracrRNA sequence, forming a PCR template that is amplified using two additional primers. The product of this reaction serves as the template for T7 *in vitro* transcription, which typically yields >100 µg of sgRNA.

### Superovulation and mating.

Mouse genome engineering requires large numbers of pronuclear-stage embryos. Consecutive injection of pregnant mare serum gonadotropin (PMSG) and human chorionic gonadotropin (hCG) is a well-established method to induce superovulation in rodents and generate large numbers of embryos (Fig. 5a-h). Both PMSG and hCG are functional analogs of gonadotropins normally produced in the anterior pituitary glands for natural sexual development and reproductive function<sup>52</sup>. Superovulated females are immediately housed with stud males to produce pronuclear-stage embryos, with a typical yield of 10–20 embryos per mouse. The number of females required for each CRISPR experiment depends on the



specific experimental design. For NHEJ-mediated editing schemes, we usually electroporate ~30 embryos to test sgRNA efficiency and ~60 embryos to generate edited mice, but more complex editing schemes may necessitate additional embryos.

### **Embryo collection and processing.**

CRISPR-EZ is best performed on pronuclear-stage embryos, where the maternal and paternal pronuclei have yet to reach S-Phase or fuse (Supplementary Fig. 4). In practice, embryos collected from superovulated females contain both pronuclear-stage embryos and unfertilized oocytes. Researchers can use differential interference contrast (DIC) microscopy to select healthy pronuclear-stage embryos for CRISPR-EZ. Alternatively, overnight culture of electroporated embryos allows healthy embryos to develop into 2-cell embryos, which can be selected for oviduct transfer.

Pronuclear-stage embryos are surrounded by the zona pellucida, a glycoprotein layer that plays an essential role for spermatozoa binding, acrosome reaction, and fertilization<sup>53</sup> (Fig. 6a). Partial zona erosion by transient exposure to Acid Tyrode's (AT) solution dissociates the zona protein constituents and enhances RNP delivery (Fig. 6b). Zona dissolution is a critical step of the CRISPR-EZ protocol, which greatly facilitates the Cas9 RNP delivery under electroporation conditions with minimal detrimental effects on embryo viability. We empirically determined ~30% zona erosion as suitable for allowing efficient RNP delivery without affecting embryo viability (Fig. 6b), as prolonged Acid Tyrode's treatment may impair preimplantation development and blastocyst hatching<sup>54</sup>.

### **RNP assembly and electroporation.**

Processed pronuclear-stage embryos are mixed with the *in vitro* assembled Cas9/sgRNA RNPs and subjected to a series of electrical pulses that transiently permeabilize the zona and cell membrane. For mouse embryos, constant, low-voltage, repeated electrical pulses with a long duration (Square Wave Form) offers the best electroporation conditions to achieve efficient RNP delivery with minimal impact on embryo viability<sup>55</sup>. In short, the first electrical pulse opens pores in the zona and embryo cell membrane, while subsequent pulses maintain pore integrity and can presumably electrophorese the RNP molecules through the cytoplasm<sup>56</sup>. Our optimal electroporation condition is empirically determined as six 3-ms pulses at 30 volts for a pool of 35–100 embryos, which consistently achieves 100% Cas9 RNP delivery efficiency<sup>24</sup> (Fig. 2c,d).

### **Embryo culture and genotyping.**

One of the key benefits of CRISPR-EZ is its capacity to rapidly test sgRNAs *ex vivo* in mouse embryos prior to generating live animals by first culturing electroporated embryos until the morula stage of preimplantation development (Fig. 1b). Depending on the experimental design, editing events can be validated in individual mouse morula by various methods, including restriction fragment length polymorphism (RFLP)<sup>57</sup>, T7 endonuclease digestion<sup>58</sup>, and PCR-based analysis. The exact sequence modification is then validated by sequencing. The same genotyping strategies can also be applied to mice using tail DNA.

RFLP is a powerful genotyping strategy when an endogenous restriction enzyme recognition site is predicted to be altered by editing, such that unedited, partially edited (heterozygous or mosaic editing), and bi-allelic edited embryos can be distinguished and quantified. We present an example RFLP assay for our *Tyr* editing scheme (Fig. 2a. b).

When an appropriate restriction enzyme site is absent from the predicted target site, a T7 endonuclease assay can be employed to estimate the sgRNA efficiency. T7 endonuclease selectively cleaves DNA heteroduplexes that form after edited and unedited DNA is heat denatured and reannealed. However, as a single embryo lacks sufficient DNA heterogeneity to form heteroduplexes, multiple embryos (we recommend >8 embryos) must be pooled for one T7 reaction. While more versatile than RLFP, the T7 endonuclease assay provides less precise information.

A PCR-based genotyping strategy is often employed for CRISPR-mediated exon deletions. We typically design a PCR strategy to amplify both the deleted and unedited sequence. The relative abundance of PCR product from the predicted deletion allele versus the unedited allele provides an estimation of the extent of editing.

### Oviduct transfer.

Live animals can be generated from electroporated embryos by performing oviduct transfer into pseudopregnant CD1 females (Fig. 7a-j). Electroporated embryos are typically implanted into the ampulla of pseudopregnant females either immediately following electroporation, or at the 2-cell stage 24 hours later. If pronuclear-stage embryos can be pre-screened by DIC microscopy, we recommend same-day transfer, as this minimizes *in vitro* culture time and increases overall viability. Oviduct transfer is technically demanding but achievable with sufficient training. At first, researchers will likely require the expertise of a highly trained technician from a transgenic facility.

## MATERIALS

### REAGENTS

#### sgRNA synthesis:

- Phusion high fidelity DNA polymerase (New England Biolabs, cat. no. M0530)
- HiScribe T7 high yield RNA synthesis kit (New England Biolabs, cat. no. E2040)
- DNase I, RNase-free (New England Biolabs, cat. no. M0303)
- SeraMag Speedbeads (GE Healthcare, cat. no. 65152105050250)
- 100% Ethanol (EtOH, Koptec, cat. no. V1016)

**CAUTION** Flammable: store in appropriate storage container. Eye irritant: wear safety glasses.

- Nuclease free water (Ambion, cat. no. AM9937, molecular biology grade)
- 1 M Tris-HCl pH 7.4 (Sigma, cat. no. T2663-1L, molecular biology grade)

- 0.5 M EDTA (Sigma, cat. no. 03690–100ML, molecular biology Grade)
- Oligos for sgRNA synthesis, Donor Oligo and PCR Primers for genotyping (Table 1, Integrated DNA Technologies, custom DNA oligonucleotides)

### Superovulation and mating

- 3–8-month-old male mice and 3–5-week-old female mice (C57BL/6J (Jax 000664) or C57B/6N (Jax 005304)  
  
CAUTION Experiments involving live mice must conform to appropriate institutional regulations. All animal experiments in this protocol were performed in accordance with the UC Davis institutional animal care and use committee (IACUC) guidelines and policies.
- Pregnant mare serum gonadotropin (PMSG; ProspecBio, cat. no. HOR-272, lyophilized)
- Human chorion gonadotropin (hCG; Millipore, cat. no. 230734, lyophilized)
- Dulbecco's phosphate buffered saline (DPBS, calcium and magnesium free; Gibco, cat. no. 14190–144)

### Embryo collection and processing:

- M2 medium (Zenith Biotech, cat. no. ZFM2–050)
- Bovine serum albumin (BSA; Sigma, cat. no. A3311, embryo culture grade)
- Hyaluronidase/M2 (Millipore, cat. no. MR-051-F)
- Acid Tyrode's solution (Sigma, cat. no. T1788, embryo culture grade)

### RNP assembly and electroporation:

- HEPES (Sigma, cat. no. H4034, cell culture grade)
- Potassium chloride (KCl; Sigma, cat. no. P9333, molecular biology grade)
- Magnesium chloride (MgCl<sub>2</sub>; Sigma, cat. no. M8266, anhydrous)
- 100% glycerol (Fisher, cat. no. BP229, molecular biology grade)
- Tris (2-carboxyethyl) phosphine hydrochloride (TCEP; Sigma, cat. no. C4706)
- OptiMEM reduced serum media (Thermo, cat. no. 31985062)
- Cas9 protein (Alt-R™ S.p. Cas9 Nuclease 3NLS (IDT, cat. no. 1074181, 61 μM)

### Embryo culture and genotyping

- KSOM+AA medium (Potassium-supplemented simplex optimized medium plus amino acids; Zenith Biotech, cat. no. ZEKs-050)
- Mineral oil (Millipore, cat. no. ES-005C, embryo grade)
- Potassium chloride (KCl, Fisher, cat. no. P217–3, certified ACS grade)

- 1M Tris-HCl, pH 8.5 Solution. (Teknova, cat. no. T1085, molecular biology grade)
- Ethylenediaminetetraacetic acid (EDTA; Sigma, cat. no. EDS-100G, anhydrous)
- Magnesium chloride (MgCl<sub>2</sub>; Fisher, cat. no. M33–500 certified ACS grade)
- Gelatin type B (Fisher, cat. no. G7–500, laboratory grade)
- Nonidet P-40 substitute (NP-40; Sigma, cat. no. 74385)
- Tween-20 (Sigma, cat. no. P7949–500, molecular biology grade)
- Proteinase K (Fisher, cat. no. BP1700–100, molecular biology grade)
- GoTaq (Promega, cat. no. M712)
- HinfI (NEB, cat. no. R0155S, 10,000 units/ml)
- EcoRI (NEB, cat. no. R3101S, 20,000 units/ml)
- Additional restriction enzymes for RFLP (Optional, NEB)
- LE agarose (BioExpress, cat. no. E-3120–500, analytical grade)
- Tris Base (Fisher cat. no. 77–68-1, molecular biology grade)
- Acetic acid (Fisher cat. no. 64–19-7, Glacial Certified ACS)
- 1 kb Plus DNA ladder (Thermo, cat. no.10787018)
- EtBr (Ethidium Bromide 10 mg/ml; Thermo, cat. no. 15585011, Ultrapure)  
CAUTION EtBr is considered to be a potent mutagen. Avoid skin, eye, mouth and upper respiratory exposure by wearing appropriate personal protective equipment. Non-Toxic alternatives can be used, such as Sybr-Safe (Thermo, cat. no. S33102)
- Gel loading dye, 6x (Thermo, cat. no. R0611, 6X)

### Oviduct transfer

- 7–9-week-old CD1 recipient female mice and 2–8-month-old vasectomized males (Charles River strain code 022, vasectomy code VASEX)  
CAUTION Experiments involving live mice must conform to appropriate institutional regulations. All animal experiments in this protocol were performed in accordance with the UC Davis institutional animal care and use committee (IACUC) guidelines and policies.
- Artificial tears ointment (Akorn, cat. no.17478–062-35)
- Betadine (Purdue products, cat. no. 67618–150-01)
- Buprenorphine hydrochloride injection CIII (Pfizer, cat. no. 00409–2012-32)  
CAUTION Partial opioid agonist and Schedule III controlled substance. Obtain appropriate authorizations prior to use.

- Ketamine (VetOne, cat. no. 501072, 100 mg/ml)
- Xylazine (Akorn, cat. no. 59399011020, 100mg/ml)

CAUTION Seek institutional approval prior to acquiring and using anesthetics.

## EQUIPMENT

### sgRNA synthesis:

- RNase-free 1.5 ml microcentrifuge tubes (VWR, cat. no. 20170–333)
- RNase-free 8-well PCR strip tube. (VWR, cat. no. 82006–606)
- Thermalcycler (Biorad S1000 or equivalent)
- Gel electrophoresis apparatus (Biorad 1704489EDU or equivalent)
- Gel doc imager XR+ system (Biorad 1708195 or equivalent)
- Magnetic stand (Invitrogen, cat. no. 12321D)
- Nanodrop ND-2000-US (Thermo, or equivalent spectrophotometer)
- Agilent Bioanalyzer 2100 (Optional, typically a facility owned equipment)

### Superovulation and mating:

- 26 G x ½” needle (BD, cat. no. 305111)  
CAUTION Sharps hazard. Please dispose of needles in sharps container.
- 1 ml syringe (BD, cat. no. 309659, without needle, latex free)

### Embryo collection and processing:

- Stereomicroscope (Nikon SMZ-U or equivalent)
- 0.22 µm PVDF syringe filter (Millipore, cat. no. SLGV033RB)
- 0.45 µm PVDF syringe filter (Millipore, cat. no. SLHV033RB)
- 15-inch rubber aspirator tube assembly (Sigma, cat. no. A5177)
- Capillary tubes (Sigma, cat. no. P0674)
- CAUTION Sharps hazard. Dispose of used needles in sharps container.
- 60 mm tissue culture dish (Greiner Bio-One, cat. no. 628–160)

### RNP assembly and electroporation:

- 0.1 cm gap electroporation cuvette (Biorad, cat. no. 1652089)
- Electroporator (Biorad Gene Pulser XCell, cat. no. 1652660, including CE and PC modules)

### Embryo culture and genotyping

- 35 mm tissue culture dish (Greiner Bio-One, cat. no. 627–160)

- Cell culture incubator (Thermo, Napco Series 8000 DH or equivalent)

#### **Oviduct transfer:**

- Warm stage/platform (C & A Scientific XH-2001 or equivalent)
- Weighing scale (OHAUS, cat. no. LS200)
- Fine dissection forceps (Roboz, cat. no. RS-4976)
- Dumont straight forceps (Fine Science Tools, cat. no. 11252–23)  
CAUTION Sharps hazard. Dispose of needles in sharps container.
- Dissection scissors (Roboz, cat. no. RS-5960)
- Serrefine forceps (Fine Science Tools, cat. no. 18050–28)
- Bead Sterilizer (Fine Science Tools, cat no. 18000–45)  
CAUTION Sterilizer could reach extremely high temperatures.
- 30 G x ½” needle (BD, cat. no. 305106)  
CAUTION Sharps hazard. Dispose of needles in sharps container.
- Hair trimmer (Andis: model no. D-4)
- Cotton tipped applicator (Medline, cat. no. MDS202000Z)
- Wound clip (Braintree Scientific, cat. no. EZC CS)
- Wound clip applier (Braintree Scientific, cat. no. EZC APL)
- Absorbable Suture (CP Medical, cat. no. 421A)  
CAUTION Sharps hazard. Dispose of needles in sharps container.

## **REAGENT SETUP**

### **Superovulation hormones:**

Reconstitute lyophilized hormones by adding pre-chilled DPBS directly to the glass bottles containing PMSG or hCG powders. Mix by gentle inversion and pipetting to ensure full reconstitution. Prepare 50 IU/ml stock solutions for PMSG and hCG. Dispense 600 µl aliquots into pre-chilled 1.5 ml centrifuge tubes, and immediately flash freeze by submerging in a liquid nitrogen bath or a dry ice/ethanol bath. PMSG and hCG stocks can be stored at –80°C for up to 1 year.

**CRITICAL**—All reagents and vessels are pre-chilled on ice, as hormones should be kept cold throughout preparation with minimal exposure to ambient temperature.

### **Mice for superovulation:**

Injections for superovulation should be timed such that female mice are ~4 weeks old at the time of mating. Females up to 5 weeks can be used but females older than 5 weeks do not respond as well to superovulation. In our experience, reliable male studs are 3–8-month-old.

Replace males if they fail to produce copulatory plugs after 3 consecutive mating attempts. After each mating, males should be rested for at least 4 or more days to recover sperm count.

**CRITICAL**—In order to ensure fertilization success, we test the plugging rate of stud males with reproductively mature females (8+ weeks). We consider males to have proven fertility if they leave copulatory plugs after three consecutive matings. Optimal superovulation conditions vary by mouse strains, and may need to be empirically determined. The described procedures are tested for C57BL/6J and C57BL/6N mice.

**Cas9 protein:**

After first thaw, make 4  $\mu$ l or 8  $\mu$ l Cas9 protein aliquots for single or double use, respectively. We recommend testing new lots of protein against previous lots in embryos to ensure efficacy. Avoid freeze thaw and store in  $-80^{\circ}\text{C}$  freezer for up to 10 weeks.

**Embryo culture media (KSOM+AA+BSA):**

Supplement KSOM+AA medium with BSA to a final concentration of 1 mg/ml, and sterilize through a 0.22  $\mu$ m PVDF syringe filter. Make 1 ml aliquots and store at 4  $^{\circ}\text{C}$  for up to 1 week once BSA is added.

**Embryo manipulation media (M2+BSA):**

Supplement M2 media with BSA to a final concentration of 4 mg/ml, and sterilize through a 0.22  $\mu$ m PVDF syringe filter. Make 1 ml aliquots and store at 4  $^{\circ}\text{C}$  for up to 1 week. M2 without BSA can be stored for up to 4 months at 4  $^{\circ}\text{C}$ .

**Ethanol, 80% (vol/vol):**

Combine 8 ml of 100% ethanol with 2 ml nuclease free water. Store at room temperature (20–25 $^{\circ}\text{C}$ ), indefinitely.

**Ethanol, 70% (vol/vol):**

Combine 7 ml of 100% ethanol with 3 ml nuclease free water. Store at room temperature, indefinitely.

**TE Buffer (10 mM Tris-HCl, 1 mM EDTA, pH 7.4):**

Prepare TE buffer by combining 10 mL of 1 M Tris-HCL (pH 7.4) with 2 mL 0.5 M EDTA and 988 mL of nuclease free water. TE Buffer can be stored at room temperature, indefinitely.

**RNP Buffer for Cas9 RNP assembly:**

Prepare RNP buffer stock by combining the reagents listed in the table below. Prepare 500  $\mu$ l aliquots and store at  $-20^{\circ}\text{C}$  for up to 6 months. Immediately prior to RNP assembly, dilute 100 mM TCEP 1:20 in RNP buffer to make complete RNP buffer (by adding 1  $\mu$ l of 100 mM TCEP to 19  $\mu$ l of RNP buffer). This complete reagent is single use and should not be stored.

**CRITICAL**—TCEP acts as a reducing agent and prevents Cas9 aggregation. TCEP is unstable in phosphate buffers at neutral pH and is therefore added just prior to complex formation. 100mM TCEP should be stored as single-use, 5  $\mu$ l aliquots at  $-20^{\circ}\text{C}$  indefinitely.

#### **Embryo Lysis Buffer for genotyping:**

Prepare Embryo Lysis Buffer by combining the reagents listed in the table below, and store as 500  $\mu$ l aliquots at  $-20^{\circ}\text{C}$  for up to 6 months. Prior to embryo lysis, dilute 20 mg/ml Proteinase K 1:100 in Embryo Lysis Buffer to a final concentration of 0.2 mg/ml (by adding 5  $\mu$ l of 20 mg/ml Proteinase K to 495  $\mu$ l Embryo Lysis Buffer). This complete reagent is single use and should not be stored.

#### **Pseudopregnant female mice for oviduct transfer:**

7–9-week-old female CD1 mice are mated to 2–8-month-old vasectomized males. Pseudopregnant CD1 females are identified by the presence of copulatory plugs the next morning.

**CRITICAL STEP**—Vasectomized males should be tested for infertility at least once by pairing with a reproductively mature female (8+ weeks), checking presence of copulatory plug, and confirming absence of sired litter. Replace males if they fail to produce copulatory plugs after 3 consecutive mating attempts.

#### **Anesthetic cocktail preparation:**

Dilute Ketamine (stock 100 mg/ml) and Xylazine (stock 100 mg/ml) by aseptically combining 1 ml stock Ketamine with 0.1 ml stock and 8.9 ml of sterile water. Store at  $4^{\circ}\text{C}$  until the expiration of either component. Check manufacturer information for expiration dates.

## **EQUIPMENT SETUP**

#### **Mouth pipet for embryo manipulation:**

For multiple CRISPR-EZ procedures, mouse embryos are manipulated using a mouth pipet loaded with a glass transfer pipet (Supplementary Fig. 5). Glass transfer pipets are produced by pulling glass capillary tubes over an open flame. An ideal glass transfer pipet possesses a 4–5 cm tip of near uniform diameter, with an opening of approximately 5x the diameter of an oocyte. To make a mouth pipet, first cut the 15-inch rubber aspirator tube in half and connect both new ends to a 0.45  $\mu$ m PVDF syringe filter to prevent contamination from the researcher. Next, insert a glass transfer pipet into the clear pipet holder.

#### **Embryo culture plates:**

For prolonged embryo culture, prepare culture plates by even distributing 5–6 20  $\mu$ l droplets of KSOM+AA+BSA media into a 35-mm cell culture plate (Supplementary Fig. 6). Carefully overlay with 2–3 ml of mineral oil using a serological pipet, and equilibrate by pre-incubating for at least 4 hours in a water jacketed, 5%  $\text{CO}_2$  incubator at  $37^{\circ}\text{C}$  and 95% humidity. **CRITICAL** As KSOM is a sodium bicarbonate buffered solution, pH is extremely



sensitive to air exposure. Mineral oil serves to buffer against temperature and pH fluctuations outside the incubator<sup>59</sup>.

Embryo manipulation plates: For embryo manipulation, collection, treatment, and washes, prepare M2+BSA medium plates immediately prior to use by evenly distributing droplets of 40  $\mu$ l M2 + 4 mg/ml BSA on 60 mm plates. As M2 is a HEPES buffered solution, it is refractory to pH fluctuations and therefore does not require a mineral oil overlay (Supplementary Fig. 7).

### Embryo workstation:

For optimal expediency, we recommend establishing an embryo workstation containing the following equipment, tools and supplies within easy reach: stereomicroscope, electroporator, CO<sub>2</sub> incubator, embryo culture/manipulation plates, hand pipets, mouth pipets, and dissection tools. Arrange these items such that dissected mouse oviducts can be immediately processed under the stereomicroscope for embryo collection. Pronuclear-stage embryos are rapidly isolated, pulsed with the electroporator, and placed into the incubator, with minimal exposure of embryos to ambient temperature.

## PROCEDURE:

### sgRNA design ● TIMING 1 day; 1–2 hours hands on

1. Candidate sgRNAs can be selected from several widely used online algorithms. To engineer indels, point mutations or small insertions, we recommend selecting 2–3 candidate sgRNAs for the target cut site based on algorithm score criteria. To engineer genomic deletions, design a pair of sgRNAs to flank the region to be excised, using one of the commonly used algorithms for sgRNA design. Some publicly-available design programs include but are not limited to: Sequence scan for CRISPR (<http://crispr.dfc.harvard.edu/SSC/>), Gene Perturbation Platform (<http://portals.broadinstitute.org/gpp/public/analysis-tools/sgrna-design>), Chop-Chop (<https://chopchop.rc.fas.harvard.edu/>), CRISPR Design (<http://crispr.mit.edu/>).

**CRITICAL STEP**—Although it remains unclear how extensive Cas9 off-target activity is in edited mice, it is important to avoid sgRNA designs with highly possible off-target sites. Not all algorithms provide off-target analysis on predicted sgRNAs. It is necessary, in some cases, to perform BLAT analyses on the selected sgRNA sequences to manually identify potential off-target sites. In our experience, the most successful sgRNAs are often highly ranked across multiple algorithms.

2. Replace the variable region with the 19–20 nt sgRNA sequence selected from Step 1, (Fig. 4, Table 1), order the sgRNA-specific oligos as custom DNA oligos from vendor (such as IDT) with standard desalting. PAGE purification is not required.

**sgRNA synthesis ● TIMING 2 days; 3 hours hands on**

3. Generate the DNA template for sgRNA *in vitro* transcription through PCR by preparing the following reaction mixture in sterile RNase-free 8-well PCR strip tube. Clean up is not required after the PCR reaction.

Place the PCR reaction in a thermal cycler using the following cycling conditions:

4. Confirm the presence of template by running 5 µl of PCR product combined with 1 µl of 6x DNA loading dye on a 2% (wt/vol) agarose gel. Expected product size will be 127bp (Fig. 4).

**? TROUBLESHOOTING**

**PAUSE POINT:** The completed PCR reaction can be stored at –20°C until needed.

5. For T7 *in vitro* transcription, follow manufacturer's instructions. Briefly, prepare the following reaction mixture in an RNase-free 8-well PCR strip tube, and mix the reagents by gentle pipetting. All components except DNA template (generated in Step 3) are provided by the NEB HiScribe T7 high yield RNA synthesis kit.

**CRITICAL STEP:** The *in vitro* transcription reaction is highly sensitive to RNase contamination. It is essential to ensure an RNase-free work environment by use of RNase-free reagents and supplies.

6. Incubate the *in vitro* transcription reaction mixture for >18 hours at 37°C in a thermal cycler.
7. Add 1 µl of RNase-free DNase I (2 units/reaction), and incubate for 20 minutes at room temperature (20–25°C) to remove DNA template.
8. Add 129 µl 100% ethanol to the reaction to facilitate subsequent small RNA binding to the magnetic SeraMag Speedbeads for sgRNA purification.
9. Thoroughly vortex SeraMag Speedbeads for 10 seconds to resuspend. Add 100 µl bead mixture to the 150 µl *in vitro* transcription reaction. Gently pipet the reaction 10 times to mix, and incubate for 5 minutes at room temperature.
10. Place the RNA+bead reaction on the magnetic stand, and wait for 5 min to allow beads to pellet towards the magnet.
11. Discard supernatant using a pipet. Add 200 µl of 80% (vol/vol) EtOH and wash the beads by gently pipetting for 5 times. Incubate for 5 minutes on the magnetic stand to allow beads to re-pellet.
12. Repeat the wash step (Step 10), then air dry for 5–10 minutes.
13. Elute RNA with 20 µl of nuclease-free water, pipet to mix 10 times, and incubate for 2 minutes at room temperature.

14. Place the reaction tube on the magnetic stand for 5 minutes at room temperature. Using a pipet, carefully transfer the RNA-containing supernatant to an RNase-free 1.5 ml microcentrifuge tube.
15. Determine the quality and quantity of sgRNA preparation using a bioanalyser and spectrophotometer, respectively. Alternatively, run 2  $\mu$ l of sgRNAs on an RNase-free 2% (wt/vol) agarose gel to check quality.

### ?TROUBLESHOOTING

**PAUSE POINT:** Purified sgRNAs can be stored for up to a year at  $-80^{\circ}\text{C}$ . To minimize freeze-thaw cycles, store sgRNAs as 5  $\mu$ l aliquots.

### Superovulation and mating ● TIMING 3 days; 2 hours hands on

16. Determine the number of females required for superovulation prior to the experiment, based on the assumption of obtaining 10–20 embryos per female. For most editing strategies, we aim to electroporate ~30 embryos per condition for testing sgRNA design in embryos, and ~60 embryos per condition for generating edited mice. Hence, 2–3 superovulated females are sufficient for testing sgRNA designs in embryos, and 4–5 recipient females are sufficient for one mouse editing experiment. It is prudent to include some extra embryos to allow for procedural embryo loss, embryo lethality and unfertilized oocytes.
17. Follow the hormone injection schedule for superovulation (Fig. 5a) by administering 5 IU PMSG (100  $\mu$ l) to ~4-week-old female mice through intraperitoneal (IP) injection at 2:00–3:00 PM on day 1. To perform IP injection, restrain the mouse by grasping behind its neck with thumb and forefinger, while securing the tail between pinky and ring finger. Invert the mouse to expose the lower abdomen, and with a 26-gauge syringe, inject at a  $45^{\circ}$  angle between two visible nipples on either side of the abdomen (Fig. 5b).
18. Approximately 46–48 hours after PMSG injection (1:00–3:00 PM on day 3), administer 5 IU hCG (100  $\mu$ l) to the same female mice by IP injection to induce ovulation.

**CRITICAL STEP**—Perform IP injections within 30 minutes of thawing either PMSG or hCG, as hormones rapidly lose activity at room temperature.

### ?TROUBLESHOOTING

19. After hCG injection, pair each hormone stimulated female with one 3–8 month old stud male with proven fertility in a mating cage.

### Embryo collection and processing ● TIMING 2 hours

20. The morning after hCG IP injection (66–68 hours after PMSG, 7:00–9:00 AM on day 4), collect females with copulation plugs for embryo collection. Typically, 60% of superovulated females display copulation plugs, an indication of a successful mating (Fig. 5c).

**CRITICAL STEP**—We provide a detailed video for all experimental procedures described following this Step (Supplementary Video 1).

### ?TROUBLESHOOTING

21. Euthanize females by CO<sub>2</sub> asphyxiation and/or additional measures according to institutional guidelines. Lay the animals on their backs and spray the abdominal area thoroughly with 70% ethanol. Using dissection scissors, open the abdominal cavity and locate the ovaries on both sides (Fig. 5d). Surgically isolate the ovaries with attached oviducts and place each ovary/oviduct in a 50 µl droplet of M2+BSA media in an embryo manipulation plate (Fig. 5e, f). We typically array 9 droplets of M2+BSA media for oviduct dissection around the perimeter of a 60 mm dish, leaving one droplet in the center for embryo collection (Supplementary Fig.7).
22. Under a stereomicroscope, nick the ampulla of the oviduct using dissection forceps (Fig. 5f), releasing the cumulus oocyte complex containing oocytes or pronuclear stage embryos surrounded by supporting cumulus cells (Fig. 5g, h, Supplementary Video 1).
23. Transfer all dissected cumulus oocyte complexes to a single 50 µl droplet of M2+BSA media using a hand pipet (Fig. 5g, h).
24. Prepare a 60-mm plate with a 100 µL droplet of Hyaluronidase/M2, and transfer the cumulus oocyte complexes to the Hyaluronidase/M2 droplet using a pipet with as little carryover as manageable. Gently pipet the cumulus oocyte complexes up and down until the majority of the embryos are dissociated from the cumulus cells. In our experience, this typically takes ~1 minute, but the length of treatment depends on the number of cumulus oocyte complexes collected. An additional 50 µL of Hyaluronidase/M2 can be added to this droplet if cumulus cell dissociation does not occur within 2 minutes.

**CRITICAL STEP**—Embryos should not be left in Hyaluronidase/M2 for longer than 5 minutes.

25. Hyaluronidase treatment releases embryos from the cumulus oocyte complex. Transfer embryos to a new 50 µl M2+BSA droplet using a mouth pipet to dilute active Hyaluronidase and wash away surrounding cumulus cells.
26. Pass the embryos by mouth pipet through 3–6 additional 50 µl M2+BSA droplets until the cumulus cells are completely removed. If cumulus cells remain, additional washes are necessary, which have minimal impact on embryo viability. At this point, embryo morphology and quality can be assessed by visual inspection under stereomicroscope (Fig. 6a). The status of fertilization can be assessed by checking for the presence of both the male and female pronuclei using a DIC capable microscope.

**PAUSE POINT:** Embryos can be temporarily maintained for no more than 30 minutes in a 50 µl M2+BSA droplet in a water jacketed, 5% CO<sub>2</sub> incubator at 37°C and 95% humidity.

**CRITICAL STEP**—Embryos are extremely sensitive to culture conditions, including temperature, CO<sub>2</sub> level, and media conditions. Be sure to test incubator conditions and check media expiration dates once a month to ensure successful *ex vivo* embryo culture.

### ?TROUBLESHOOTING

27. Mildly erode the zona pellucida by brief embryo exposure to Acid Tyrode's solution. Using a mouth pipet, transfer the embryos to a droplet of Acid Tyrode's solution in a 60 mm plate. Observe the embryos under a stereomicroscope, until the zona pellucida is ~30% eroded. This usually occurs at approximately 30–40 seconds (Fig. 6b, Supplementary Video 1).

**CRITICAL STEP:** The timing of Acid Tyrode's treatment is crucial, as prolonged exposure can completely dissolve the zona pellucida and compromise embryo viability. Expect some heterogeneity in the extent of zona erosion among the embryos.

### ?TROUBLESHOOTING

28. Using a mouth pipet, swiftly pass the embryos sequentially through at least four 50 µl droplets of M2+BSA to dilute out the Acid Tyrode's solution. Obtain a final embryo count to confirm sufficient numbers for experimental conditions. It is not uncommon to lose up to 5% of embryos during these manipulations steps.

**PAUSE POINT:** Embryos can be temporarily stored for no more than 30 minutes in a 50 µl M2+BSA droplet in a water jacketed, 5% CO<sub>2</sub> incubator at 37°C with 95% humidity.

**CRITICAL STEP:** As the exact timing of fertilization is unknown, and embryos from multiple females are combined, it is important to realize that the embryos will be within a few hours of each other on a development timeline and this heterogeneity should not be misinterpreted as a developmental delay or consequence of manipulation. Some embryos will fail to be fertilized (oocytes) or develop and do not progress beyond the 1–2 Cell stages.

## RNP assembly and electroporation ● TIMING 1 hour

29. Immediately following embryo processing, prepare Cas9 RNP complex reaction mixtures. To assemble Cas9/sgRNA RNP complex for NHEJ mediated indel, use option A; to engineer HDR mediated editing, use option B; to make defined deletions, use option C.

### A. For NHEJ-mediated indels

- i) Combine the following reagents at room temperature in a sterile, RNAse free 8-well PCR strip.
- ii) Incubate the reaction mixture in a thermal cycler at 37°C for 10 minutes to allow complex formation.

**B. For HDR mediated editing**

- i) Combine the following reagents at room temperature in a sterile, RNase free 8-well PCR strip.
- ii) Incubate the reaction mixture in a thermal cycler at 37°C for 10 minutes to allow complex formation.

**C. For engineering deletions using paired sgRNAs**

- i) Combine the following reagents at room temperature in a sterile, RNase free 8-well PCR strip.
- ii) Incubate the reaction mixture in a thermal cycler at 37°C for 10 minutes to allow complex formation.

**PAUSE POINT:** Assembled RNP complex made using option A,B or C can be stored at room temperature for up to 2 hours.

30. Transfer the processed embryos from Step 28 through two 50 µl droplets of OptiMEM to dilute out the M2+BSA in preparation for electroporation. **CRITICAL STEP:** It is important to remove BSA, as it provides undesired resistance during embryo electroporation.
31. Transfer 30 (for *in vitro* testing) or 60 (for animal derivation) embryos to a 10 µl droplet of OptiMEM. Add 10 µl pre-assembled Cas9/sgRNA RNP complex (from Step 29) to this droplet and mix thoroughly by pipetting up and down 10 times, until the embryo/RNP mixture is homogenous by visual inspection under stereomicroscope.

**CRITICAL STEP:** The Cas9/sgRNA RNP mixture has a thicker viscosity than OptiMEM due to the presence of glycerol. The embryo/RNP mixture must be pipetted gently to reach homogeneity.

32. Transfer the entire 20 µl embryo/RNP mixture to a 0.1 cm gap electroporation cuvette, avoid making bubbles, and electroporate embryos in a Biorad Gene Pulser XCell apparatus with a square wave protocol and the following parameters:

**?TROUBLESHOOTING**

33. Recover electroporated embryos by flushing the cuvette at least three times with 50 µl KSOM+AA+BSA media using a pipet, transferring and combining each wash into a 60 mm plate. Avoid introducing air bubbles during this procedure. It is optional to repeat this procedure to maximize the recovery of electroporated embryos. We typically recover >95% of the originally loaded embryos.
34. Using a mouth pipet, transfer the recovered embryos to one droplet of KSOM +AA+BSA media in an equilibrated embryo culture plate. This serves to dilute the remaining RNP/OptiMEM mixture. Then transfer the embryos to a second

droplet in the same plate for subsequent embryo culture (Step 35A) or in preparation for oviduct transfer (Step 35B).

**PAUSE POINT:** Embryos can be cultured for up to four days in the equilibrated embryo culture plate while in a water jacketed, 5% CO<sub>2</sub> incubator at 37°C with 95% humidity.

**CRITICAL STEP:** Prior to use, the embryo culture plate containing KSOM+AA+BSA media overlaid with mineral oil that and is equilibrated by pre-incubation in a cell incubator for at least 4 hours is prepared according to Supplementary Fig. 6.

35. At this point, embryos can either be cultured to the morula stage for embryo genotyping (option A) or transferred to a recipient female oviduct for the generation of CRISPR edited mice (option B). When attempting this protocol for the first time, subject the embryos to *ex vivo* culture conditions according to option A to test sgRNA editing efficiency. Once the results from the *ex vivo* testing conditions are adequate, one can transfer electroporated embryos to generate mice using option B.

**(A) Embryo culture and genotyping ● TIMING 4–5 days; 4–6 hours hands on**

- i. The following day, under a stereoscope, transfer healthy 2-cell embryos to a fresh droplet of KSOM+AA+BSA, and culture to morula/blastocyst embryos in an equilibrated embryo culture plate while in a water jacketed, 5% CO<sub>2</sub> incubator at 37°C with 95% humidity. Typically, pronuclear stage embryos develop to morula after ~72 hours of culture, and to blastocysts after ~84 hours of culture.
- ii. Wash morula/blastocyst embryos by sequentially passing them through two droplets of DPBS using a mouth pipet.
- iii. Using a pipet set to 1 µl, transfer each individual embryo into one well of an 8-well PCR strip containing 10 µl of Embryo Lysis Buffer + Proteinase K (0.2 mg/ml).
- iv. Lyse the embryos to extract DNA by heating the samples in a thermal cycler at 55°C for 4 hours, followed by a 10-minute incubation at 95°C to inactivate Proteinase K.

**PAUSE POINT:** Embryo lysates can be stored at 4°C for 2–3 days or –20°C for up to 2 weeks before genotyping analysis. Avoid multiple freeze-thaws.

**CRITICAL STEP:** Embryo genotyping assays vary depending on the exact editing scheme. Provided are procedures to assess NHEJ-or HDR-mediated editing for the *Tyr* editing scheme using RFLP. As this workflow is highly optimized and reproducible, we strongly recommend performing the *Tyr* editing experiment as a positive control in CRISPR-EZ experiments, particularly for the first-time users.

- v. Perform PCR #1 of a nested PCR to amplify DNA fragments flanking the targeted site from small amounts of DNA from crude single-embryo lysates (from Step 35Aiv). Prepare the first PCR reaction mix as described:

Place the PCR reaction in a thermal cycler using the following cycling conditions:

**PAUSE POINT:** PCR product can be stored at 4°C for 2–3 days or –20°C for up to 2 weeks before genotyping analysis. Avoid multiple freeze-thaws.

- vi. Perform PCR #2 of a nested PCR reaction to generate a highly specific PCR product using 2 µl of 1:10 diluted first PCR reaction as the template. Dilution of the first PCR product serves to reduce primer carryover in the nested PCR. Prepare the nested PCR reaction mix as described:

Place the PCR reaction in a thermal cycler using the following cycling conditions:

**CRITICAL STEP:** Annealing temperature depends on primer sequences. This particular reaction is showing sgTyr conditions. We typically achieve >90% reliable PCR reactions when amplifying products up to 500 bp from individual embryos.

#### ?TROUBLESHOOTING

- vii. To genotype NHEJ-mediated *Tyr* indel mutations, digest 10 µl of the nested PCR products from Step 35Aiv with 10 units of *HinfI* in a 20 µl reaction and incubate for 4 hours at 37°C. Run digestion products on a 2% (wt/vol) agarose gel, using non-digested PCR product as a loading control for 30 minutes at 135 volts. There is sufficient loading dye carry over from the previous PCR reaction.
- viii. To genotype HDR-mediated *Tyr* small sequence substitution, digest 10 µl of the nested PCR products from Step 35Aiv with 20 units of *EcoRI* in a 20 µl reaction and incubate for 4 hours at 37°C. Run the digestion products on a 2% (wt/vol) agarose gel, using non-digested PCR product as a loading control for 30 minutes at 135 volts. There is sufficient loading dye carry over from the previous PCR reaction.

**CRITICAL STEP:** In any HDR-mediated editing, both HDR and NHEJ editing events can occur, which are not mutually exclusive within the same embryo. We recommend to perform both NHEJ and HDR RFLP analyses to determine the genotype of each embryo in a HDR editing experiment.

#### (B) Oviduct Transfer ● TIMING 19 Days; 2–3 hours hands on

- i. On the day before the embryo transfer, prepare 5 pseudopregnant recipient females by combining 5 females with 5 vasectomized males, so that each mating cage has a single male and female mouse.

**CRITICAL STEP:** Typically, 60 embryos are electroporated, from which 40+ viable embryos are expected to be recovered, which can be transferred to 2x CD1 pseudopregnant



mothers. If more embryos are available for transfer, be sure to prepare additional mating cages for pseudopregnant females.

- ii. On the morning of the embryo transfer, prepare for survival surgery by sterilizing all surgical tools with 70% ethanol and placing into a bead sterilizer until use. Prepare an empty mouse cage by placing it on a warm stage/platform set at 37 °C.
- iii. On the same morning, identify 0.5 days post coitum (dpc) females with visible copulatory plugs. Weigh each pseudo-pregnant female and IP inject 0.1 ml of anesthetic cocktail for every 10g of body weight (dosage of Ketamine at 100 mg/kg and Xylazine at 10mg/kg) using a 1 ml syringe attached to a 26G x ½” needle.

**CRITICAL STEP:** Once administered, mice typically succumb to anesthetics within 1–2 minutes which lasts for 60–120 minutes. Mice should be monitored after experiment according to institutional guidelines. Ketamine over dosage may cause death. Repeat injections should be avoided if at all possible.

- iv. After one minute, check if the mouse is fully anesthetized by pinching its toes with forceps. Once the mouse is unresponsive, place the anesthetized mouse under the stereomicroscope and apply artificial tears ointment to its eyes to prevent drying during surgery (Fig 7a). Using a hair trimmer, shave the hair over the surgical area, along the midline of the back near the last rib (Fig. 7b).
- v. To disinfect the surgical area, apply betadine by cotton tipped applicator, scrubbing from the center of the surgical site and radiating outwards to disinfect approximately a 2-inch square area (Fig. 7c). Subsequently wipe off with 70% (vol/vol) ethanol. Repeat this procedure twice to fully sterilize the surgical area (Supplementary Video 1).
- vi. Make a small longitudinal incision ( 1cm) parallel to the midline at the level of the last rib and slide the skin to expose the body wall. Pick up the body wall with forceps over the site of the ovary and make a small incision through the body wall (Fig. 7d).

**CRITICAL STEP:** Take precaution to avoid cutting blood vessels, as bleeding may obstruct the oviduct transfer and increase risk of surgical complications.

- vii. Identify the fat pad attached to the ovary, and using blunt fine forceps, pull it out along with the ovary and upper part of the uterus from the incision site (Fig. 7e). Clip the fat pad with serrefine forceps and lay down the ovary over the middle of the back (Fig. 7f).
- viii. Load the glass transfer pipet by drawing up M2+BSA medium about 1 cm up the pipet, then one small air bubble, then 10 embryos from Step 34 within a minimal amount of media, and finally another small air bubble (Fig. 7j).

**CRITICAL STEP:** Up to 20 embryos can be transferred into the oviducts of each pseudopregnant female (10 embryos per oviduct). Typically, 60 embryos are electroporated where 40+ viable embryos are expected to be recovered and transferred to 2 CD1 pseudopregnant mothers.

- ix. Insert the transfer pipette with loaded embryos into the infundibulum, and gently blow the embryos into the infundibulum of the oviduct until the two bubbles are seen in the ampulla to ensure successful transfer (Fig. 7g, j). Alternatively, if the infundibulum is too difficult to locate, use a 30 G x ½” needle to puncture a small hole upstream of the ampulla and insert the transfer needle into this newly made opening (Fig. 7h, k).
- x. Remove the transfer pipette and unclip the serrefine forceps. Using blunt forceps, gently place the fat pad, ovary, and uterus back into the peritoneal cavity, and sew up the body wall with absorbable sutures (Fig. 7i, Supplementary Video 1).
- xi. Transfer embryos into the other oviduct of the same mouse by sliding the skin over to expose the opposite side of the back, and repeat Steps 35Bvi-x.
- xii. Hold the skin together and close the opening with wound clips.
- xiii. For pain relief, deliver 0.1 ml of Buprenorphine to the mouse by subcutaneous injection into the lower abdominal region using a 26 G x ½” needle.
- xiv. Place the mouse into the pre-warmed empty cage and monitor for vital signs (e.g. respiratory rate) until awake. House mice in an IACUC facility until pups are born. For C57BL6/J mice, gestation period is typically 18–19 days.

**CRITICAL STEP:** Animals should be monitored daily for at least 10 days following surgery and additional analgesia can be given at the recommendation of the veterinarian.

After the gestational period, collect tail samples from the pups for genomic DNA extraction<sup>60</sup>, in accordance with institutional guidelines. Perform genotyping analysis to determine editing efficiency by repeating Steps 35Av-viii xv.

● **TIMING:** Steps 1–2, sgRNA design, 1 day; 1–2 hours hands on.

Steps 3–15, sgRNA synthesis, 2 days; 3 hours hands on.

Steps 16–19, Superovulation and mating, 3 days; 2 hours hands on.

Steps 20–28, Embryo collection and processing, 2 hours.

Steps 29–34, Complex formation and electroporation, 1 hour hands on.

Step 55A, Embryo culture and genotyping, 4–5 days; 4–6 hours hands on

Step 35B, Oviduct Transfer, 19 days; 2–3 hours hands on

? **TROUBLESHOOTING:** Troubleshooting advice can be found in Table 2.

**ANTICIPATED RESULTS:** Once the desired editing strategy is designed, our protocol typically yields >100 µg of sgRNA at >6000 ng/µl concentration for efficient Cas9/sgRNA RNP assembly. In both C57BL/6J and C57BL/6N mouse strains, one can achieve 100% embryo delivery of Cas9/sgRNA RNPs, with or without a ssODN (tested up to 162 nt in length). For NHEJ-mediated editing experiments, our superovulation protocol typically recovers 10–20 viable embryos per plugged female, ~80% of which are fertilized and viable, and ideal for electroporation. The editing efficiency ranges from 3% to 100% when engineering genomic deletions (21 independent experiments, 1275 zygotes electroporated, 249 pups born; Fig. 3e, g, Supplementary Fig 2,3, Supplementary Table 2, 3), from 50% to 100% when engineering indel mutations (3 independent experiments, 405 embryos electroporated, 97 pups born; Fig. 2c-i, Supplementary Table 1; data not shown), and from 14% to 63% when engineering precise small sequence replacements (3 independent experiments, 439 zygotes electroporated, 125 pups born; Supplementary Table 1; personal communications, Polina Lishko and Ellen Robey). Taken together, CRISPR-EZ outperforms microinjection in a variety of CRISPR genome engineering applications in mice.

### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

### ACKNOWLEDGMENTS

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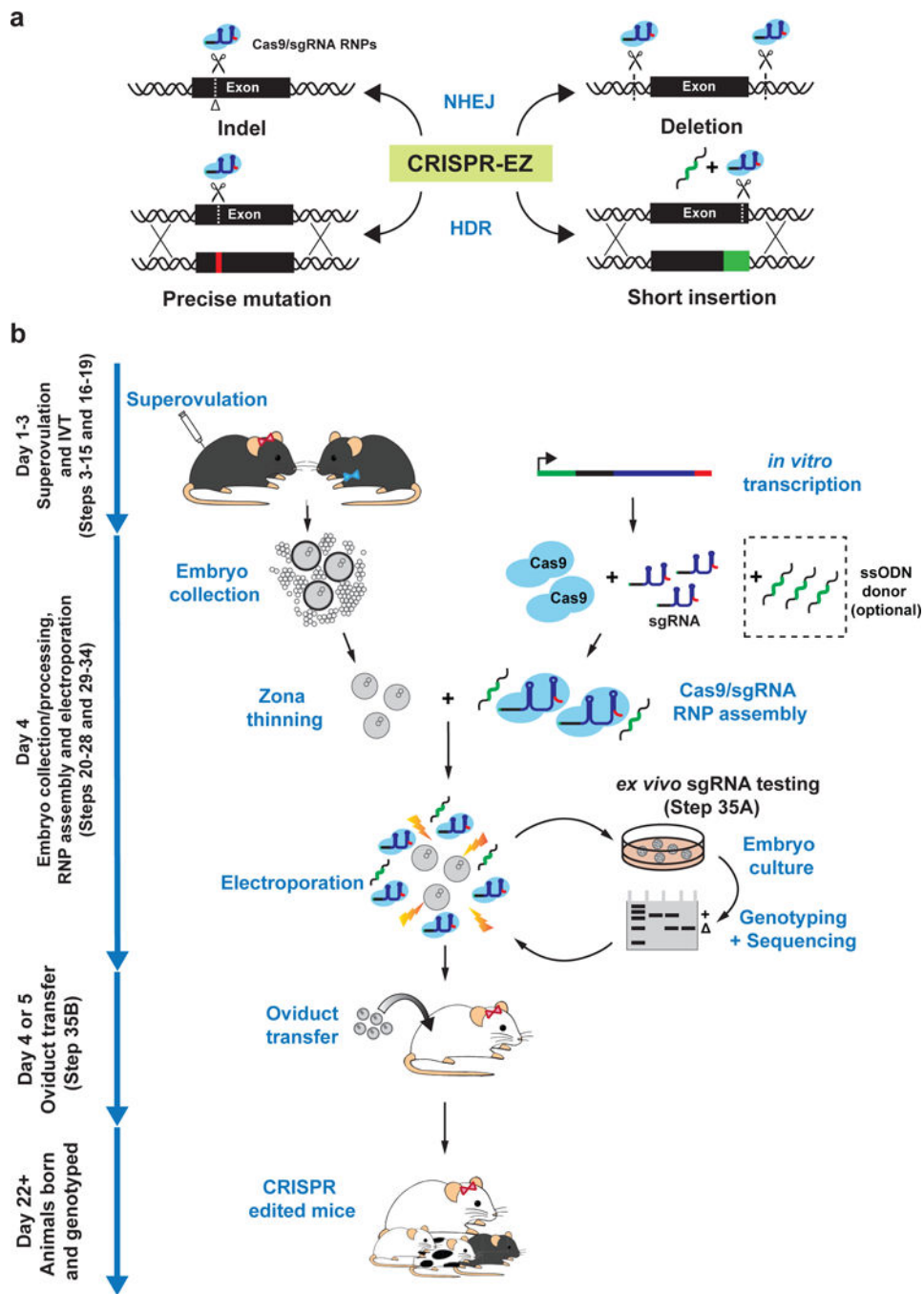
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**Figure 1. Overview of CRISPR-EZ technology and workflow.**

(a) An illustration of the most successful CRISPR-EZ editing strategies. A single sgRNA can be used to create a small indel via NHEJ repair or in conjunction with a ssODNs to create a precision mutation or a small insertion by HDR. Multiple sgRNAs can be used to engineer a genomic deletion by NHEJ repair. Design sgRNAs, HDR donor oligos, and editing validation assays prior to CRISPR-EZ experiments. (b) A graphic overview of CRISPR-EZ workflow. Day 1–3: ~4 week-old females are superovulated, first by PMSG injection, 46–48 hours later by hCG injection, before being housed with stud males for

breeding. In parallel, sgRNAs are *in vitro* transcribed and purified. Day 4: Pronuclear stage embryos are collected and processed for electroporation, while Cas9/sgRNA complexes are assembled *in vitro*. Embryos (harvested at 0.5 dpc), Cas9/sgRNA RNPs, and optional ssODNs are combined in an electroporation cuvette and subjected to a series of electrical pulses. We recommend *ex vivo* validation of sgRNA editing efficiency in cultured morulae or blastocysts before generating edited mice. With a validated sgRNA design, electroporated embryos can be transferred to the oviduct of 0.5 dpc, pseudopregnant mothers to generate genetically engineered mice, which are then genotyped to confirm editing efficiency.

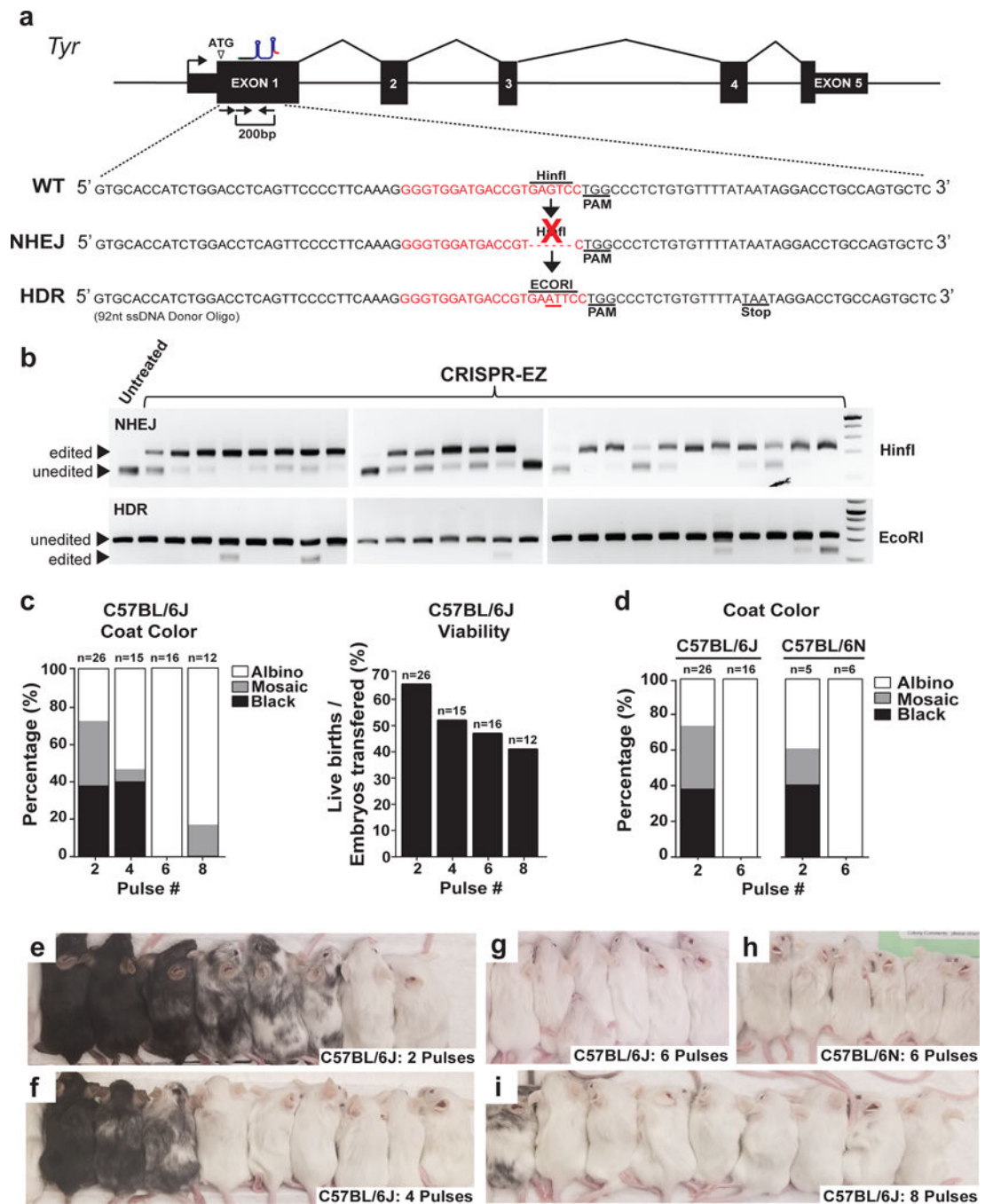
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**Figure 2. Optimization of CRISPR-EZ conditions for editing efficiency and embryo viability.** (a) A diagram illustrates the NHEJ and HDR editing strategies for exon 1 of the *Tyr* gene. A successful NHEJ editing ablates a *HinfI* site and disrupts *Tyr* gene function. A successful HDR editing replaces the *HinfI* site with an *EcoRI* site, introducing a frameshift mutation that abolishes *Tyr* gene function. (b) Representative RFLP results of *Tyr* edited mice indicate successful NHEJ editing (top) and HDR editing (bottom). (c) Since bi-allelic *Tyr* deficiency causes albinism in edited mice, the extent of albinism correlates the extent of *Tyr* editing that disrupts the genes function. Coat color (left) and viability (right) of C57B/6J edited mice

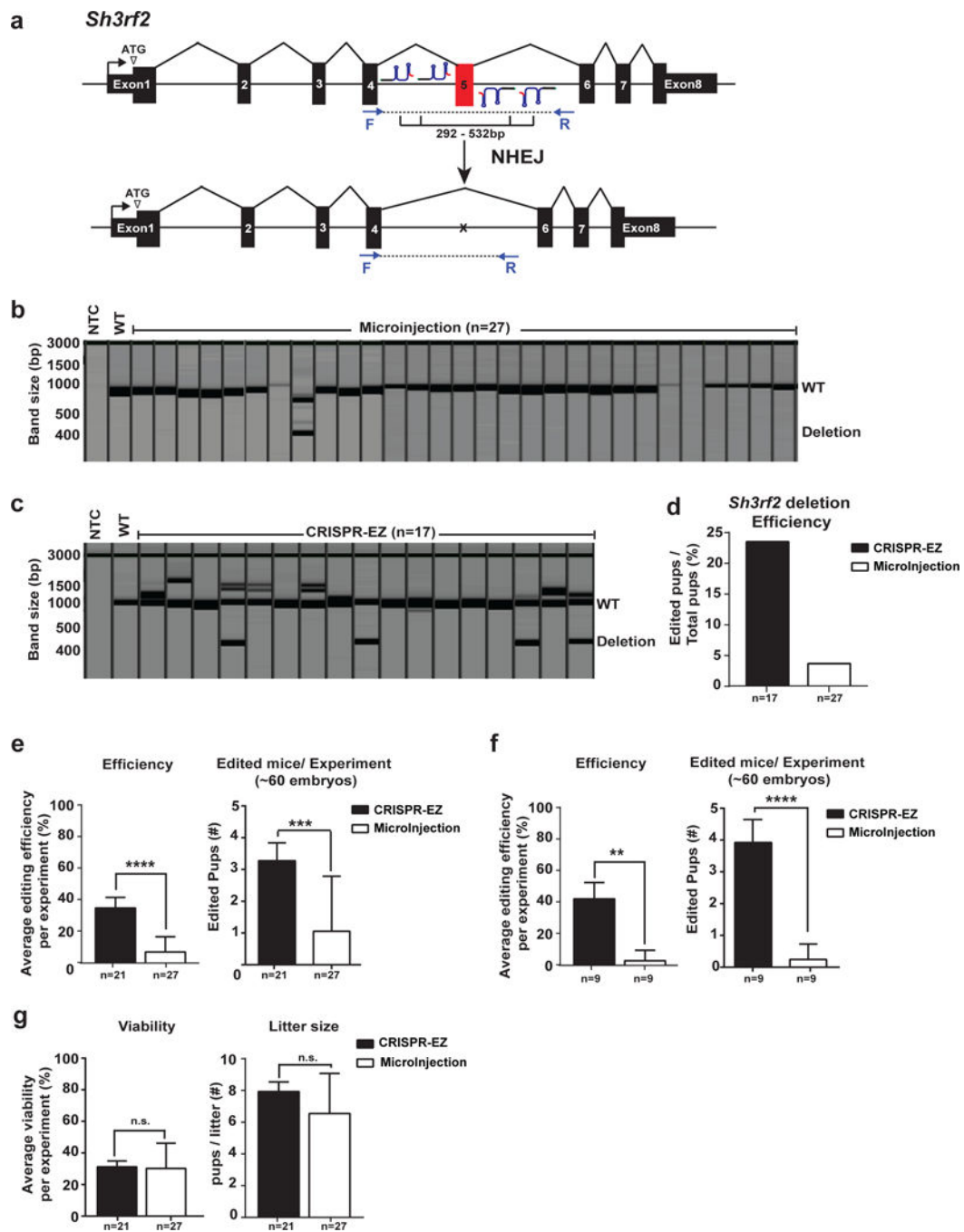
generated from 2, 4, 6 or 8 pulse CRISPR-EZ conditions. Viability is defined as the percentage of live animals born out of total embryos transferred. The 6-pulse condition maximizes editing efficiency while minimally impacting pup viability. **(d)** Comparison of editing efficiency between C57B/6J and C57B/6N mouse strain using 2 or 6-pulse electroporation conditions. The 6-pulse CRISPR-EZ condition is equally effective in both strains. **(e-i)** Representative images are shown for the coat color of edited mice from experiments shown in **(b-d)**. All animal procedures were approved by the Institutional Animal Care and Use Committee of UC Davis.

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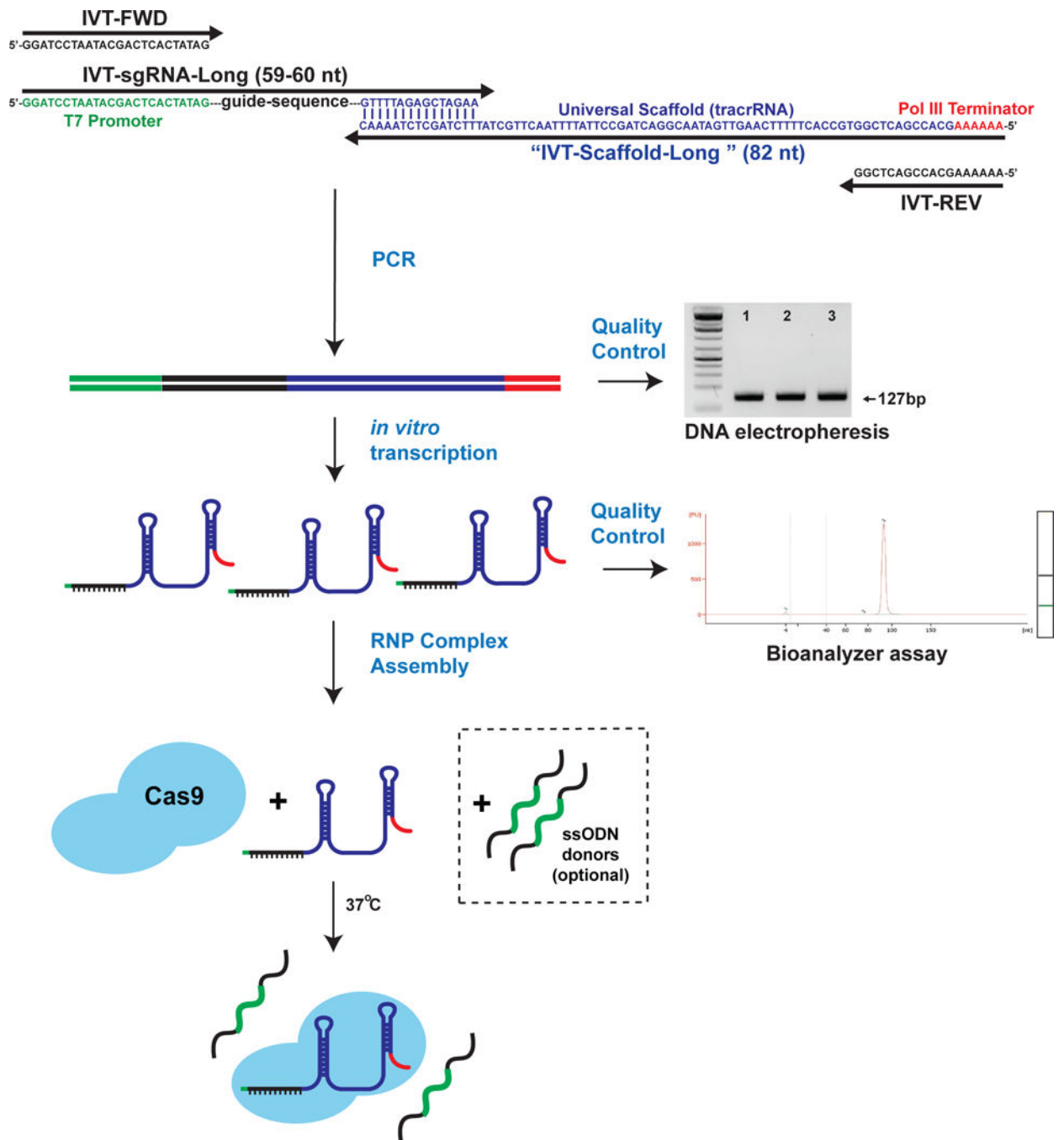
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**Figure 3. Comparing editing efficiency and viability of CRISPR-EZ and microinjection.** (a) A schematic diagram illustrates the strategy to employ two pairs of sgRNAs to mediate the deletion of *Sh3rf2* exon 5. (b,c) QIAxcel fragment analysis images show PCR genotyping results of all mice generated through microinjection (b) and CRISPR-EZ (c) in C57BL/6N mice, demonstrating a greater editing efficiency in CRISPR-EZ experiments. (NTC= non-template control, WT=wildtype) (d) Deletion efficiency of *Sh3rf2* exon 5 is shown for a side-by-side comparison between CRISPR-EZ and microinjection. (e) In a high throughput pipeline, up to two pairs of sgRNAs (4 sgRNAs) were designed to mediate exon

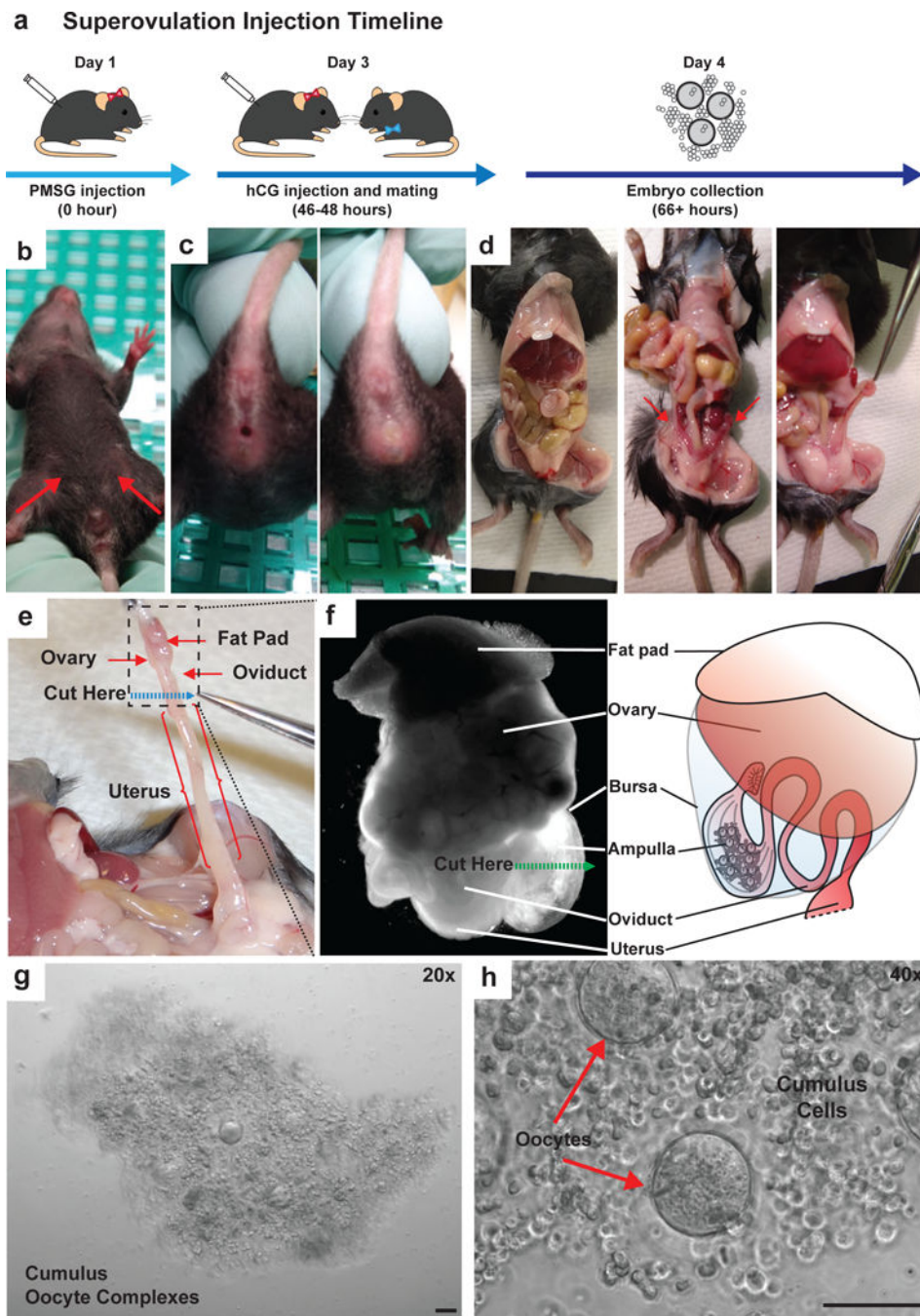
deletion in C57BL/6N mice. Average editing efficiency is calculated across 21 CRISPR-EZ experiments or 27 microinjection experiments. CRISPR-EZ outperformed microinjection in editing efficiency (left), generating an average of 3 founder animals when starting with ~60 embryos for electroporation (right). **(f)** In a direct comparison, gene knockout experiments were carried out in C57BL/6N mice using either CRISPR-EZ (n=9) or microinjection (n=9), using identical sgRNA designs. CRISPR-EZ outperformed microinjection in editing efficiency (left), generating an average of 4 founder animals when starting with ~60 embryos for electroporation. **(g)** No obvious differences are observed for animal viability (left) or litter size (right) between CRISPR-EZ and microinjection. Data are means  $\pm$  SD across all genes. \*\*  $P < 0.01$ ; \*\*\*  $P < 0.001$ ; \*\*\*\*  $P < 0.0001$ ; n.s., not significant (All  $P$ -values were calculated on a basis of an unpaired one-tailed Student's  $t$  test). All animal procedures were approved by the Institutional Animal Care and Use Committee of UC Davis.



**Figure 4. Diagram illustrating a cloning-free strategy for sgRNA synthesis.**

The sequences and purpose of each synthesized oligo are diagrammed to show how they function in this cloning free strategy. The components include a pair of PCR primers (black: IVT-FWD and IVT-REV), a common reverse template oligo (blue/red: IVT-Scaffold-Long), and an oligo containing a 5' T7 promoter and a unique sgRNA sequence (green/black/blue: IVT-VAR-sgRNA). The DNA template for sgRNA synthesis is generated by a PCR reaction. The product of this reaction is a single 127 bp amplicon which should be confirmed by gel electrophoresis prior to continuing. Shown on the right is a representative DNA

electrophoresis image of the PCR reaction. Subsequently, sgRNAs are synthesized by T7 *in vitro* transcription (IVT) and purified. Before moving forward, the quality and quantity of the newly synthesized sgRNA can be determined by submitting the sample for BioAnalyzer testing. A representative bioanalyzer trace of the IVT products is shown to the right. Recombinant Cas9 protein, purified sgRNA(s), and ssODN (optional) are assembled into RNPs *in vitro* by combining the components with a stabilizing buffer at 37°C for 10 minutes. The active RNP Complex is now ready for electroporation.

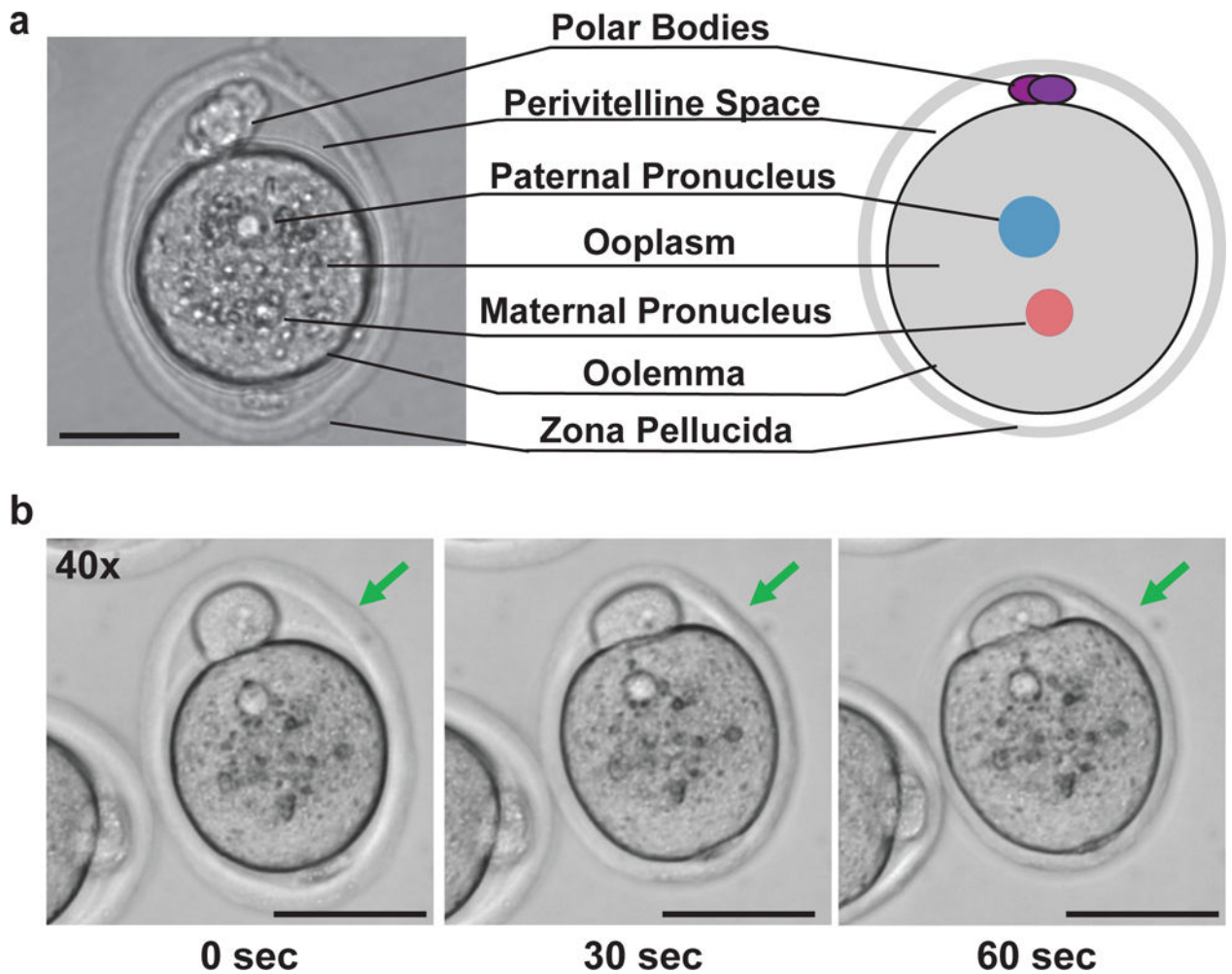


**Figure 5. Overview of key superovulation and zygote collection procedures.**

(a) A diagram illustrating the hormone injection, mating and embryo collection timeline for superovulation. (b) IP injection sites of PMSG (day 1) and hCG (day 3) are indicated in the abdominal region of a female mouse (red arrows). (c) After hCG injection at day 3, superovulated females are paired 1:1 with stud males for breeding, and copulatory plugs are evident in successfully mated females in the following morning (day 4). An example of non-plugged female (c, left) and a plugged female (c, right) is shown. (d) An image shows the internal anatomy of a female dissected abdominal cavity (d, left). Intestines are placed to the

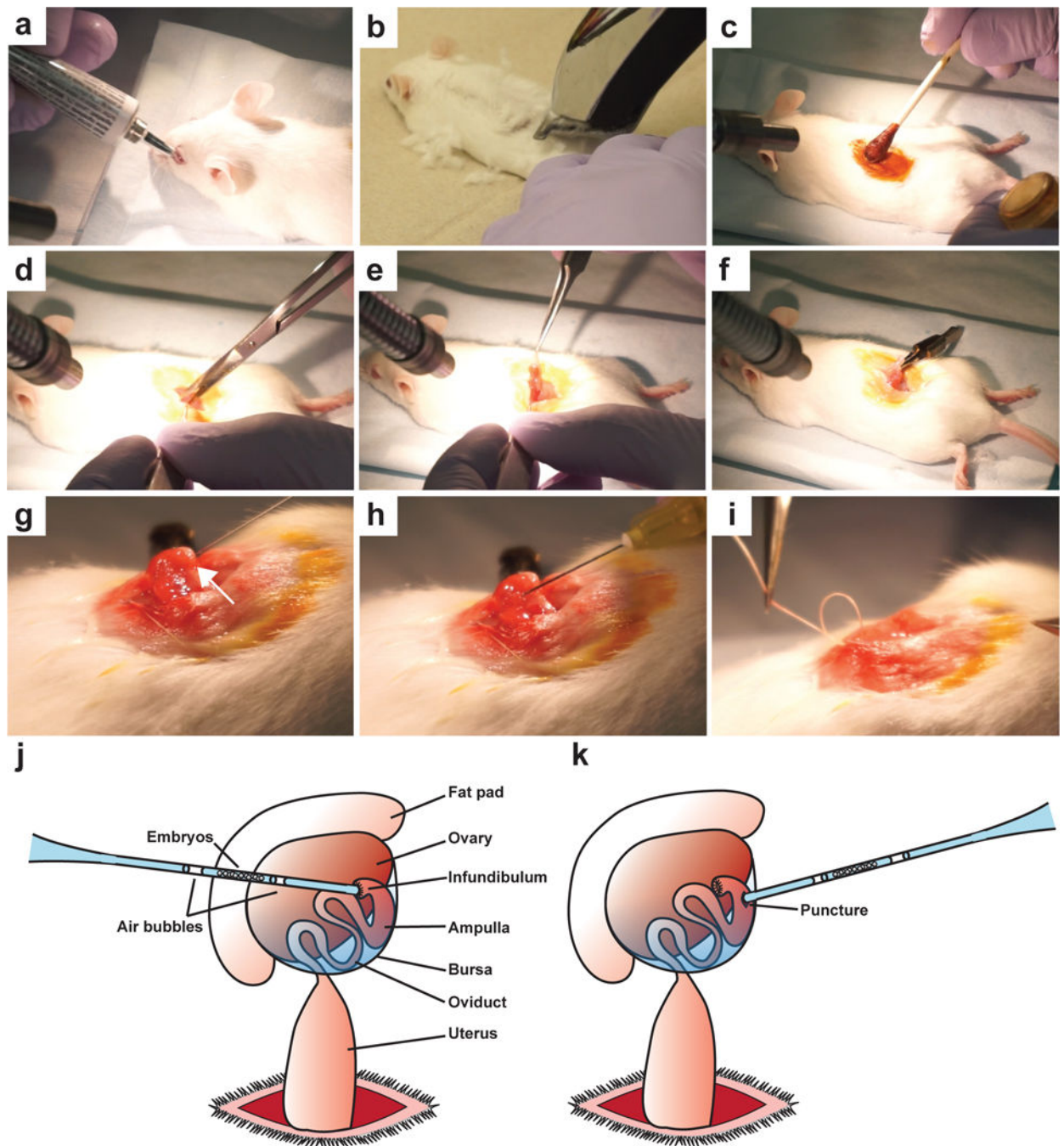
side for easier visualization and manipulation. Ovaries are located behind the intestines, each sharing a fat pad with kidney (**d**, middle, red arrows). Fat pads are used as manipulation points to avoid damaging the ovaries or oviducts during extraction. (**d**, right). Fat pads are gently detached from the kidneys by trimming along the mesometrium. (**e**) The ovary and associated oviduct is dissected by cutting the oviduct/uterine junction (dashed blue line). The tissue in the dotted black square is isolated. (**f**) A bright field image of isolated ovary and oviduct under a stereoscope (left), next to a cartoon representation of the same structure (right). Cumulus oocyte complexes can be seen through the expanded ampulla membrane, and are released by nicking the ampulla. The location and direction of oviduct nicking procedure is indicated by a green dashed line. (**g**) A representative image of cumulus oocyte complexes released as a single intact mass from an individual ampulla. (**h**) A magnified segment of panel G shows the composition of the cumulus oocyte complex, with oocytes shown with red arrows and cumulus cells between oocytes. All scale bars: 100  $\mu\text{m}$ .





**Figure 6. An overview of key zygote processing procedures.**

**(a)** Cartoon diagram illustrating various cellular structures of a pronuclear stage embryo (right), with a corresponding DIC image (left). **(b)** Representative images are shown for pronuclear stage embryos at various degrees of zona thinning upon treatment with Acid Tyrode's solution: fully intact zona (left), properly thinned zona (middle), overly thinned zona showing collapse of the perivitelline space (right). Prolonged embryo to acidic conditions impairs embryo viability. Scale Bars: 50  $\mu$ m.



**Figure 7. An overview of key procedures for oviduct transfer.**

(a) Following anesthetic cocktail administration for anesthesia, artificial tears ointment is applied to the eyes of pseudopregnant mice to prevent drying as mice are unable to close their eyes under anesthetics. (b) Shave the hair over the surgical area of the mice, along the midline of the back near the last rib. (c) Apply betadine by cotton tipped applicator to the surgical area followed by two 70% EtOH treatments using a cotton tipped applicator. (d) Using a pair of small scissors, make a small longitudinal incision along the midline at the level of the last rib and carefully position the skin to expose the body wall. Make another

small incision through the peritoneal cavity above the ovary. **(e)** Using blunt fine forceps, grasp the fat pad and gently pull out the ovary and upper part of the uterus from the incision site. **(f)** Clip the fat pad with serrefine forceps and lay down the ovary over the middle of the back. **(g)** Using a 30G x ½” needle, puncture a small hole near the ampulla, proximal to the ovary (white arrow). **(h)** Insert the glass transfer pipet with loaded embryos into the hole, and gently blow the embryos into the oviduct. **(i)** After gently returning the organs with blunt forceps, sew the body cavity closed using sutures. **(j)** Cartoon diagram illustrating the anatomy of the fat pad, ovary, oviduct, and uterus, as well as the proper location of the embryo transfer into the ampulla. The glass transfer needle is preloaded as follows: M2+BSA media, an air bubble, M2+BSA media with embryos, an air bubble, M2+BSA media. Successful transfer is indicated by the appearance of both air bubbles within the oviduct. **(k)** Cartoon diagram illustrating an alternative method to introduce embryo into the oviduct. A small opening is made between the infundibulum and ampulla using a 30G needle, and transferred embryos will settle in the ampulla.

**Table 1.**

Oligo sequences for sgRNA synthesis, donor oligo and PCR primers for genotyping

Name	Sequence	Concentration
IVT-SGRNA-LONG	5'-GGATCCTAATACGACTCACTATAG---20nt guide sequence--- GTTTTAGAGCTAGAA	1 $\mu$ M
IVT-SCAFFOLD-LONG	5'-AAAAAAGCACCGACTCGGTGCCACTTTTT CAAGTTGATAACGGACTAGCCTTATTTAAC TTGCTATTTCTAGCTCTAAAAC	1 $\mu$ M
IVT-FWD	5'-GGATCCTAATACGACTCACTATAG	100 $\mu$ M
IVT-REV	5'-AAAAAAGCACCGACTCGG	100 $\mu$ M
IVT-TYR-LONG	5'-GGATCCTAATACGACTCACTATAGGGGTGGATGACCGTGAGTCCG TTTTAGAGCTAGAA	1 $\mu$ M
TYR-ECORI-DONOR	5'- GTGCACCATCTGGACCTCAGTTCCTTCAAAGGGGTGGAT GACCGTGAATTCTGGCCCTCTGTGTTTTATAATAGGACCTG CCAGTGCTC	200 $\mu$ M
sgTyr F1	5'-TCTTTTCGGAGACTCAAATCA	100 $\mu$ M
sgTyr F2	5'-TCTGTACAATTGGGCCCCC	100 $\mu$ M
sgTyr R1	5'-GCTTTCAGGCAGAGGTTTCCT	100 $\mu$ M

Table 2.

## Troubleshooting Advice

Step	Problem	Possible Reason	Solution
3	Failed PCR amplification of DNA template for <i>in vitro</i> transcription	Failed PCR reaction, DNase contamination.	We recommend running 5 $\mu$ l of the 50 $\mu$ l reaction on a 2% (wt/vol) agarose gel to confirm PCR product size and PCR efficiency/specificity. Faint or smeared bands indicate failed PCR reactions. One can attempt the PCR genotyping again with optimized primer design and PCR reagents.
15	Failed sgRNA synthesis.	Failed <i>in vitro</i> transcription, RNase contamination.	We recommend evaluating sgRNA quality using bioanalyzer analysis. Denaturing the sample by pre-heating at 70°C for 5 minutes before running on the bioanalyzer removes additional bands by denaturing any secondary structure (Supplementary Fig. 8). If sgRNA synthesis fails, repeat IVT with RNase-free reagents.
18	Failed superovulation	Hormone leaks during IP injection, expiration of hormones.	If hormone leaks during IP injection, re-inject a half dose of PMSG or hCG (~3 IU). When using a new batch of hormone, we recommend to perform a positive control by IP injection of validated hormones that works well in the past.
20	Suboptimal embryo yield	Lack of plugged females.	In our experience, roughly 50% of non-plugged females have fertilized embryos. We suggest collecting from the non-plugged females, with the anticipation to recover fewer fertilized embryos.
26	Embryos fail to develop after electroporation.	Inappropriate embryo culture.	Embryos are sensitive to perturbation of culture conditions, such as temperature, oxygen, and humidity. Calibrate temperature and CO <sub>2</sub> level of your incubator on a monthly basis. KSOM+AA+BSA is sensitive to pH fluctuations when exposed to air. (Supplementary Fig. 6). Do not leave embryos in M2+BSA wash droplets at room temperature for longer than 5–10 minutes at a time (Supplementary Fig. 7).
26	Embryos fail to develop after electroporation.	Embryos are lysed/inviable.	Low embryo yield or poor-quality embryos can be caused by compromised hormones, poor injection technique, or improper hormone dosage. The vasculature of the ovary can provide an indication of proper superovulation—a red, highly vascularized ovary has undergone superovulation, while a pale ovary suggests unsuccessful superovulation. Embryos that do not develop or appear lysed could be due pipetting steps that are done too vigorously (Supplementary Fig. 9). If problem persists, replace hormones and ensure injections are administered in the appropriate time window (Fig 5b).
27	Embryos show low editing efficiency	Insufficient Acid Tyrode's treatment	To determine optimal Acid Tyrode's treatment duration, transfer 2–3 embryos to the droplet while viewing under a stereomicroscope. Determine how many seconds are required to completely dissolve the zona pellucida, then test again using half this amount of time. The optimal treatment duration is typically close to half the time required to completely dissolve the zona pellucida. On average, it takes about 1–2 minutes. If

Step	Problem	Possible Reason	Solution
			the Zona is still present after 2 minutes, replace aliquot and test on 3–4 more zygotes.
32	Low viability/low editing.	Pulse number optimization.	A certain balance can be struck when performing the electroporation. Fewer pulses have less impact on viability, but lower editing efficiency. More pulses tend to have higher editing but with a slight loss in viability. For complex genome editing schemes, we recommend trying different pulse numbers to enhance editing efficiency.
32	Low viability/low editing	Electroporator Functionality	If there are doubts whether the machine delivers the correct voltage at the correct intervals, an electrical engineer or the manufacturer technical services could be consulted to confirm the instrument in proper working order.
35Avi	Failed genotyping PCR	Inefficient lysis or lost embryo.	Attempt PCR with robust primers to confirm template is present. It is also possible the embryo was lost during transfer to the microcentrifuge tube. Check for the presence of an embryo immediately prior to lysis.
35Avi	All samples are edited	Premature lysis.	Since embryos are used for this protocol, essentially only a few dozen molecules of template DNA are used per reaction. It is important to use appropriate non-template and water only controls to ensure the measured editing efficiency is accurate.

<b>Component</b>	<b>Amount (ml)</b>	<b>Final concentration</b>
1 M HEPES, pH=7.5	1.0	100 mM
3 M KCl	2.5	750 mM
1 M MgCl <sub>2</sub>	0.05	5 mM
100% Glycerol	5.0	50% (vol/vol)
Nuclease-free H <sub>2</sub> O	0.95	

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<b>Component</b>	<b>Amount (ml)</b>	<b>Final concentration</b>
1 M KCl	0.5	50 mM
1 M Tris-HCl, pH=8.5	0.1	10 mM pH=8.5
1 M MgCl <sub>2</sub>	0.025	2.5 mM
Gelatin (powder)	1 mg	0.1 mg/ml
Nonidet P-40	0.045	0.45% (vol/vol)
20% Tween 20	0.225	0.45% (vol/vol)
Nuclease-free H <sub>2</sub> O	To 10 ml	-

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<b>Component</b>	<b>Amount (<math>\mu</math>l)</b>	<b>Final Concentration</b>
H <sub>2</sub> O	35.5	-
5x Phusion HF Buffer	10	1X
10 mM dNTPs	1	0.2 mM
IVT-FWD primer (1 $\mu$ M)	1	0.02 $\mu$ M
IVT-REV primer (1 $\mu$ M)	1	0.02 $\mu$ M
IVT-SGRNA-LONG oligo (100 $\mu$ M)	0.5	1 $\mu$ M
IVT-SCAFFOLD-LONG oligo (100 $\mu$ M)	0.5	1 $\mu$ M
Phusion HF (2 Units/ $\mu$ l)	0.5	0.02 Units
Total	50	-

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Cycle Number	Denature	Anneal	Extend
1	95°C, 2 min		
2–31	95°C, 10 s	72°C, 10 s	72°C, 10 s
32			72°C, 2 min

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<b>Component</b>	<b>Amount (<math>\mu</math>l)</b>	<b>Final Concentration</b>
10x HiScribe Reaction Buffer	2	1X
ATP (100 mM)	2	10 mM
GTP (100 mM)	2	10 mM
CTP (100 mM)	2	10 mM
UTP (100 mM)	2	10 mM
DNA template	8	(Step 3)
T7 RNA polymerase mix	2	-
Total	20	-

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<b>Component</b>	<b>Amount (<math>\mu</math>l)</b>	<b>Working Concentration</b>
5x RNP Buffer + TCEP	2	1x
61 $\mu$ M Cas9 Protein	1.31	8 $\mu$ M
2 $\mu$ g/ $\mu$ l sgRNA (from Step 15)	2	4 $\mu$ g
TE Buffer pH 7.4	4.69	-
Total	10	-

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<b>Component</b>	<b>Amount (<math>\mu</math>l)</b>	<b>Working Concentration</b>
5x RNP Buffer + TCEP	2	1x
61 $\mu$ M Cas9 Protein	1.31	8 $\mu$ M
2 $\mu$ g/ $\mu$ l sgRNA (from Step 15)	2	4 $\mu$ g
TE Buffer pH 7.4	3.69	-
200 $\mu$ M ssDNA oligo (optional)	1	20 $\mu$ M
Total	10	-

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Component	Amount ( $\mu$ l)	Working Concentration
5x RNP Buffer + TCEP	2	1x
61 $\mu$ M Cas9 Protein	2.62	16 $\mu$ M
2 $\mu$ g/ $\mu$ l sgRNA 1 (from Step 15)	2	4 $\mu$ g
2 $\mu$ g/ $\mu$ l sgRNA 2 (from Step 15)	2	4 $\mu$ g
TE Buffer pH 7.4	1.38	-
Total	10 $\mu$ l	-

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<b>Condition</b>	<b>Value</b>
Voltage	30 V
Pulses	4–6
Pulse Length	3 ms
Pulse Interval	100 ms

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<b>Component</b>	<b>Amount (<math>\mu</math>l)</b>	<b>Final Concentration</b>
2x GoTaq	5	1x
Single Embryo Lysate from Step 35Aiv	3	-
10 $\mu$ M sgTyr F1 (see Table 1)	1	1 $\mu$ M
10 $\mu$ M sgTyr R1 (see Table 1)	1	1 $\mu$ M
Total	10	-

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Cycle Number	Denature	Anneal	Extend
1	95°C, 2 min		
2–31	95°C, 10 s	60°C, 30 s	72°C, 10 s
32			72°C, 10 min

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<b>Component</b>	<b>Amount (<math>\mu</math>l)</b>	<b>Final Concentration</b>
Nuclease-free water	6	-
2x GoTaq	10	1x
1:10 Diluted 1 <sup>st</sup> PCR Product (from Step 35Av)	2	-
sgTyr F2 10 $\mu$ M (see Table 1)	1	1 $\mu$ M
sgTyr R1 10 $\mu$ M (see Table 1)	1	1 $\mu$ M
Total	20	-

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Cycle Number	Denature	Anneal	Extend
1	95°C, 2 min		
2–31	95°C, 10 s	60°C, 30 s	72°C, 10 s
32			72°C, 10 min

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