

Gene Editing of Livestock

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Presentation Primer

- 1. Why humans work to modify the genome of food animals**
- 2. Strategies to edit genes in livestock cells**
- 3. Strategies to generate gene edited livestock**
- 4. Conscientious considerations**

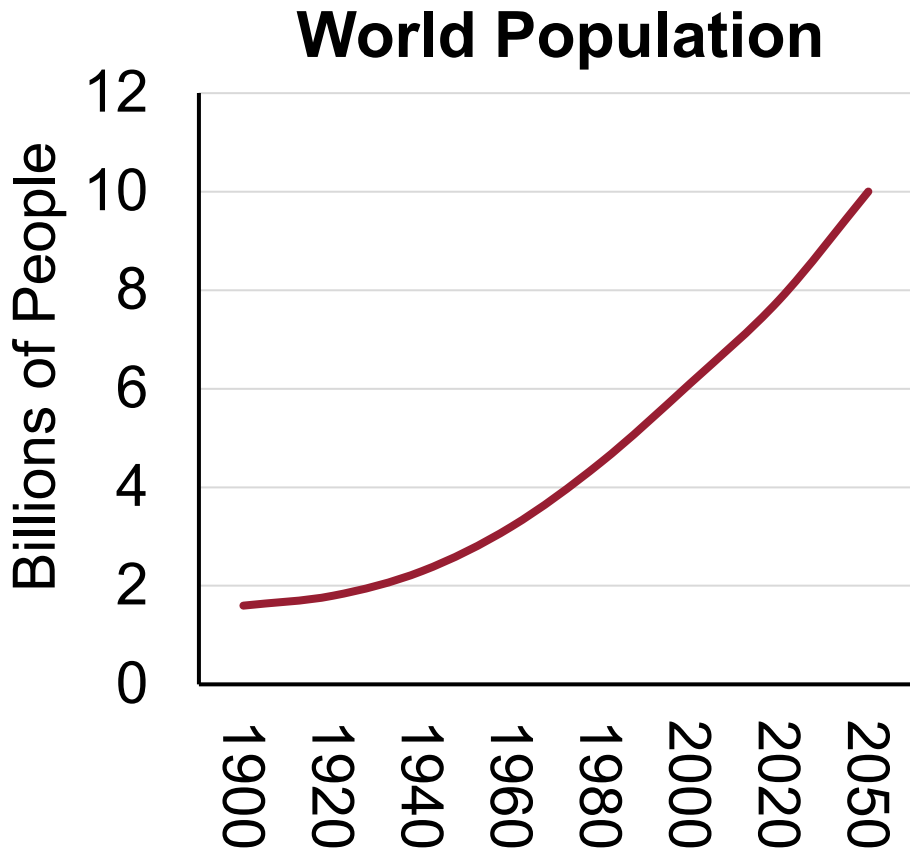
Animal Genetic Engineering

Genome Shaped via Human Intervention



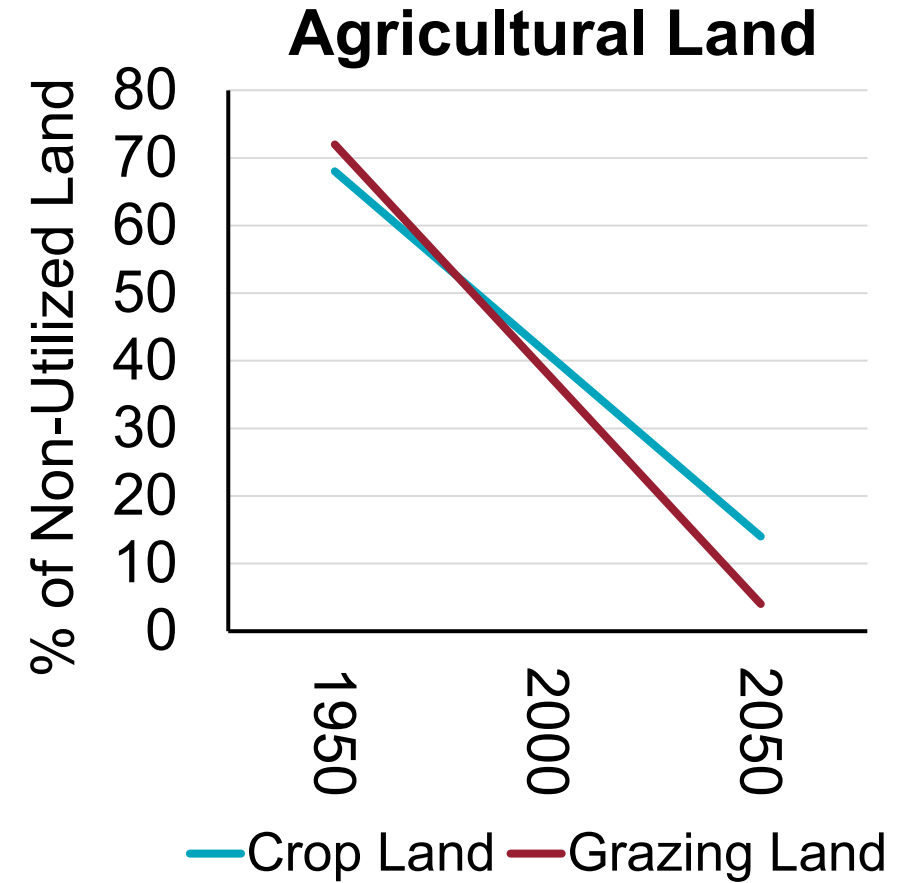
Credit: Getty Images

Food (In)Security



Source: US Census Bureau

60%
**Agricultural
Production**



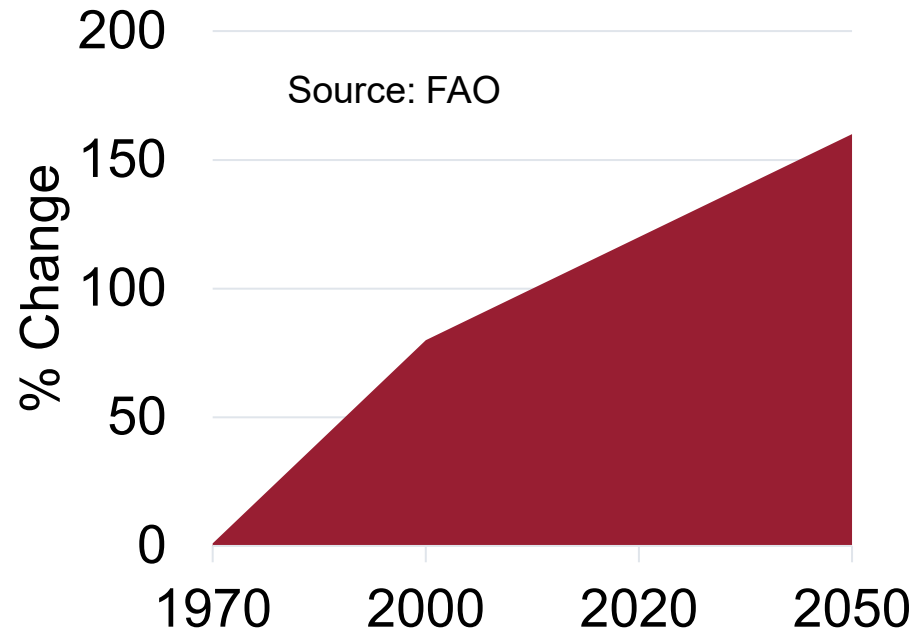
Source: Food and Agricultural Organization of the United Nations

Global Problem



↓ Agricultural Land ↑ Demand for Food

Demand for Animal Protein

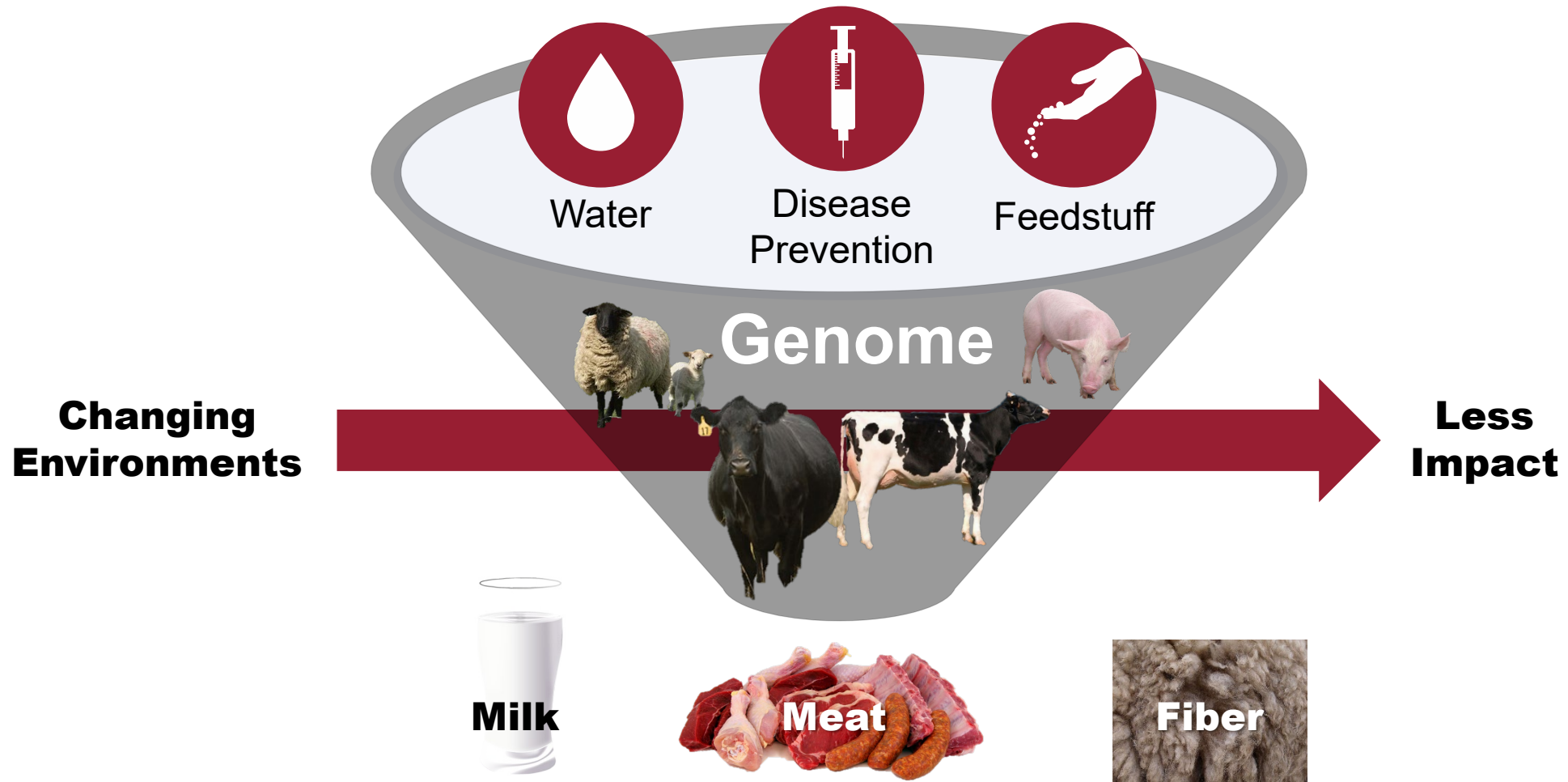


A photograph of a herd of black cattle in a field. The central focus is a close-up of a cow's face, looking directly at the camera. Its ears are large and slightly flared. In the background, other cows and a small calf are visible in a field of dry, yellowish grass under a clear sky.

Solution

**Enhance livestock production by
engineering resiliency & efficiency**

Livestock Production Efficiency & Resiliency



Selective Breeding Shapes the Genome

+ Genetic Gain, but...

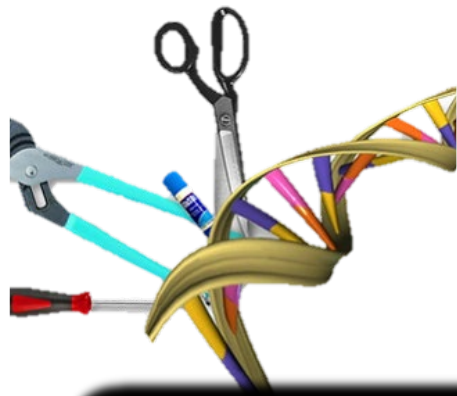


Based Primarily on Observable Phenotype

5+ Generations & 10+ Years for Incremental Changes

Genetic Drift & Inadvertent Negative Phenotypes

Promise of Gene Editing



**Molecular
Tools**

+



**DNA
Information**



**Efficient & Precise
Genetic Engineering
1-2 Generations**

Gene Editing Strategies

CRISPRs

ZFNs

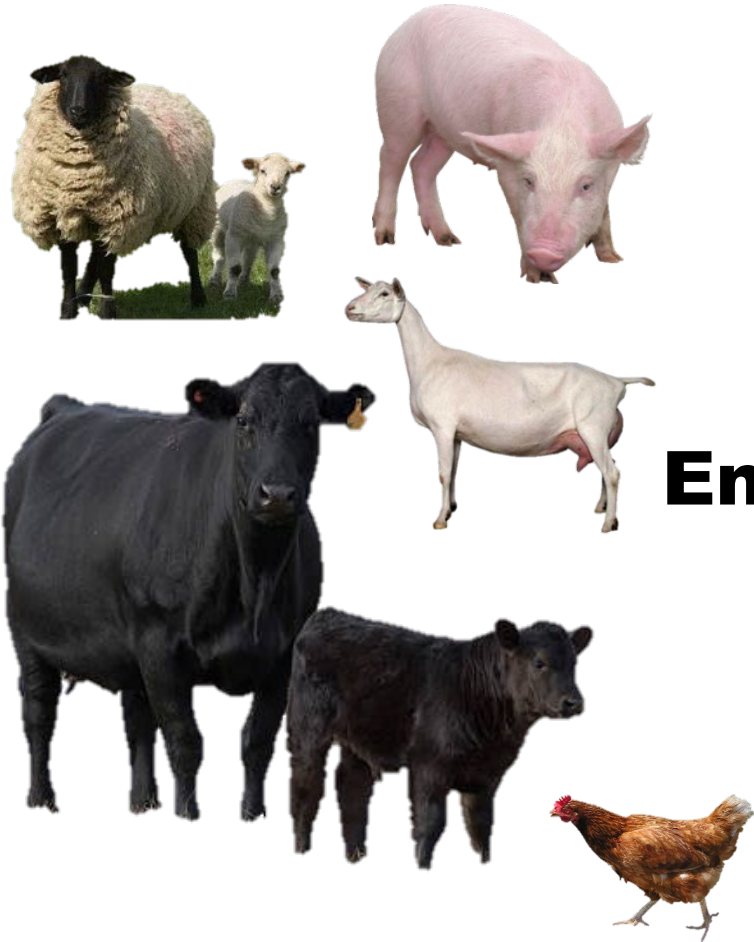


TALENs

Absence of Foreign DNA Sequence

Genomic Changes that Could Arise in Nature

The Leading Edge of Gene Editing for **Livestock Resilience**



Welfare

Disease Resistance

Environmental Adaption

Growth

Reproduction

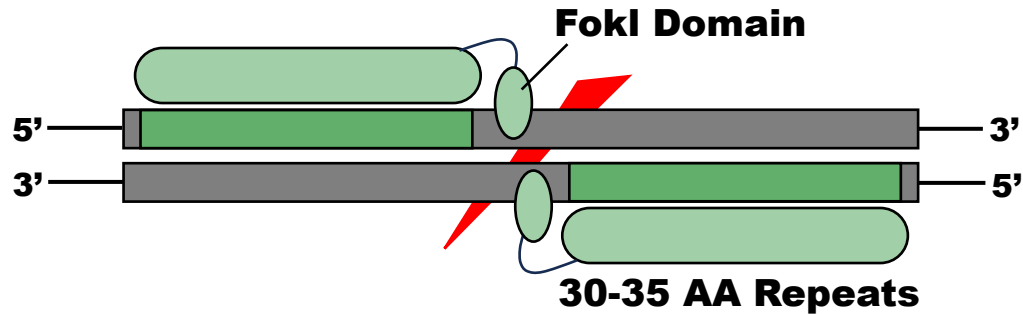
**Future of
Food Security**

Strategies to Edit Livestock Genomes

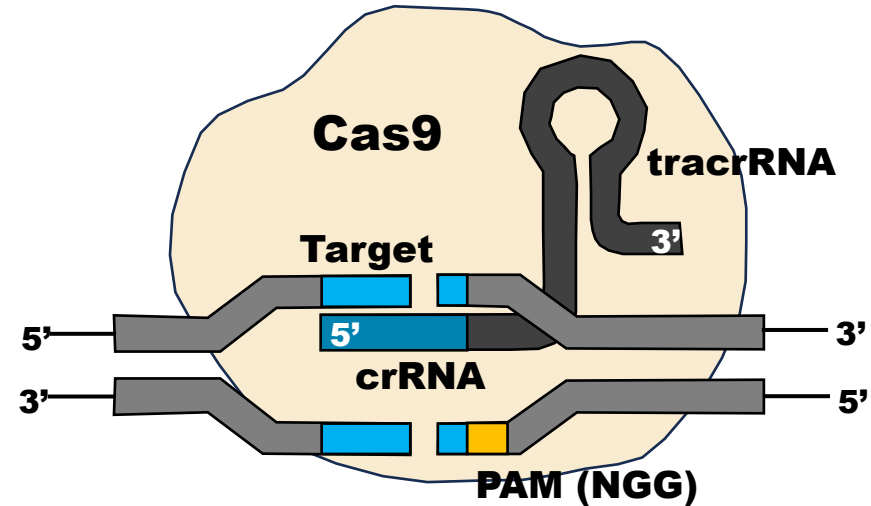
Conventional & Next Generation

Conventional Strategies are the Primary Use with Food Animals

TALEN



CRISPR-Cas9



+ Repair Template (ssDNA, long dsDNA, plasmid)



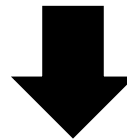
Site-Specific DNA Double Strand Break (DSB)



- **Non-Homologous End Joining (NHEJ) – Random Insertion-Deletion Mutations (INDELs)**
- **Homology Directed Repair (HDR) – Sequence-Specific Insertions or Replacements**

Cas Variants

	Cas9	Cas9 Nickase (D10A/H840A)	Cas12
Cas Family	Type II	Type II	Type V
PAM sequence	G-rich	G-Rich	T-rich
Cut type	Blunt double-strand , 3 bp upstream of PAM	Blunt single strand , 3 bp upstream of PAM	Staggered, 18–23 bp downstream of PAM
RNAs needed	crRNA + tracrRNA	crRNA + tracrRNA	crRNA
Major Application	Mammalian gene editing	Mammalian gene editing	Diagnostics

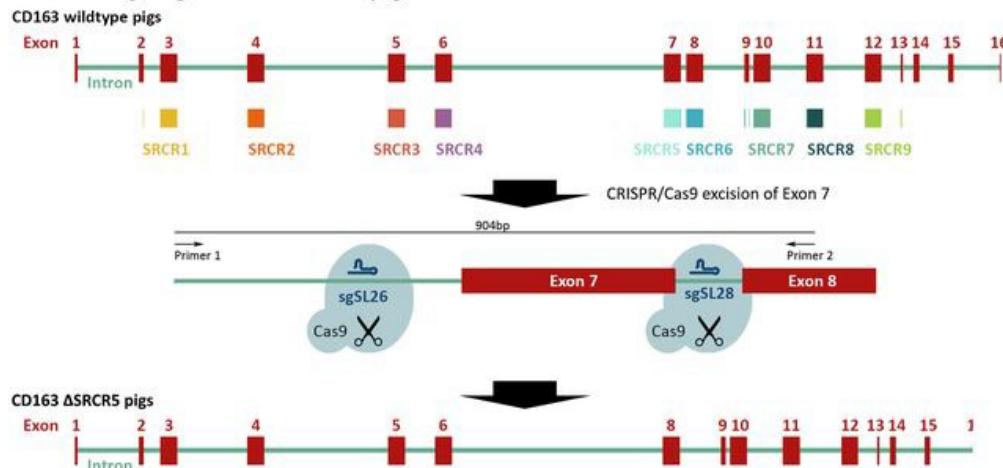


Generate gDNA breaks for INDEL or HDR

Livestock w/ Engineered INDEL Alleles

PRRS Resistant Pigs – Editing CD163 *University of Missouri & Roslin Institute*

A Genome editing to generate Δ SrcR5 pigs

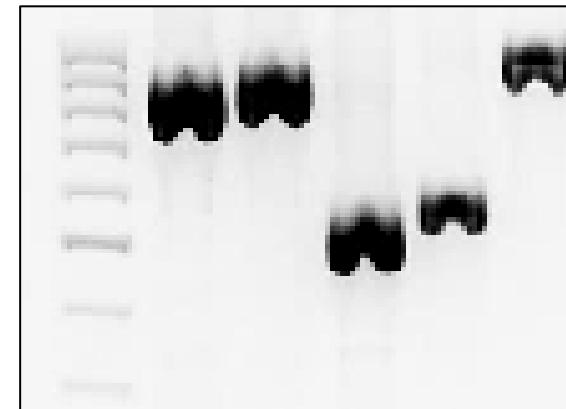


Burkard et al., 2018, J Virol

Surrogate Sires – Editing *NANOS2* *Washington State University*



Knockout via Large Exon Deletion



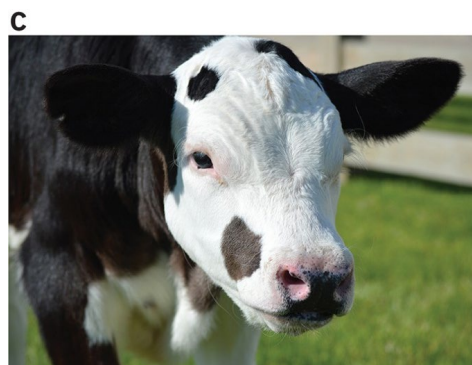
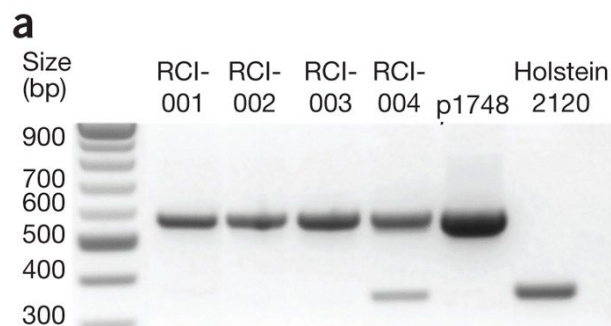
160 bp Δ / 510 bp Δ

Miao et al., 2019, Biol Repro; Ciccarelli et al., 2020, PNAS

Livestock w/ Engineered HDR Alleles

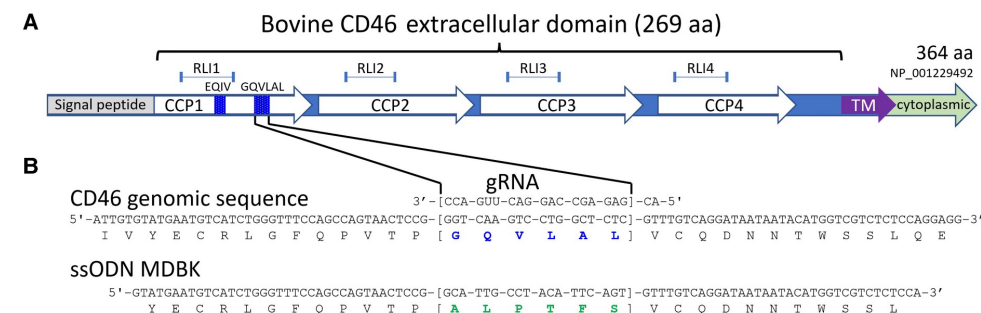
Hornless Dairy Cattle – Polled Allele *Recombinetics*

212 bp Introgression to Replicate Celtic Polled Allele

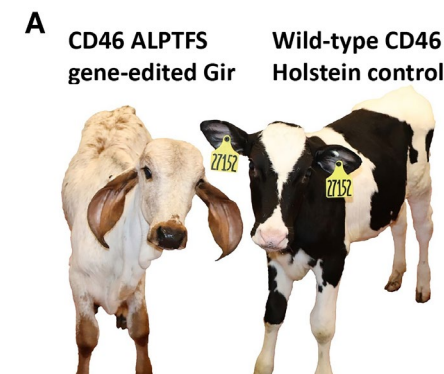


Carlson et al., 2016, Nat Biotech

BVD Resistant Cattle – Editing *CD46* *USDA-MARC & Recombinetics*



6AA Substitution – Viral Binding Domain



Workman et al., 2023, PNAS Nexus

Array of Editing Outcomes

Peer-Reviewed Scientific Literature + Washington State University (Unpublished Data)
Mice, Cattle, Pigs, Goats, and Sheep

Allelic Inactivation/Alteration via INDELs

- ❖ **Range of modifications (1 bp to >1 kb)**
- ❖ **Insertions from other chromosomes**
- ❖ **Retrotransposon insertions (e.g. LINE1)**
- ❖ **Mono-allelic or bi-allelic editing**
- ❖ **Alleles with same edit or each allele edited differently**

Site-Specific Insertion/Replacement via HDR

- ❖ **Designed edit with no unexpected modification**
- ❖ **Designed edit with unexpected INDEL**
- ❖ **Rearranged insert**
- ❖ **Concatemer integration**

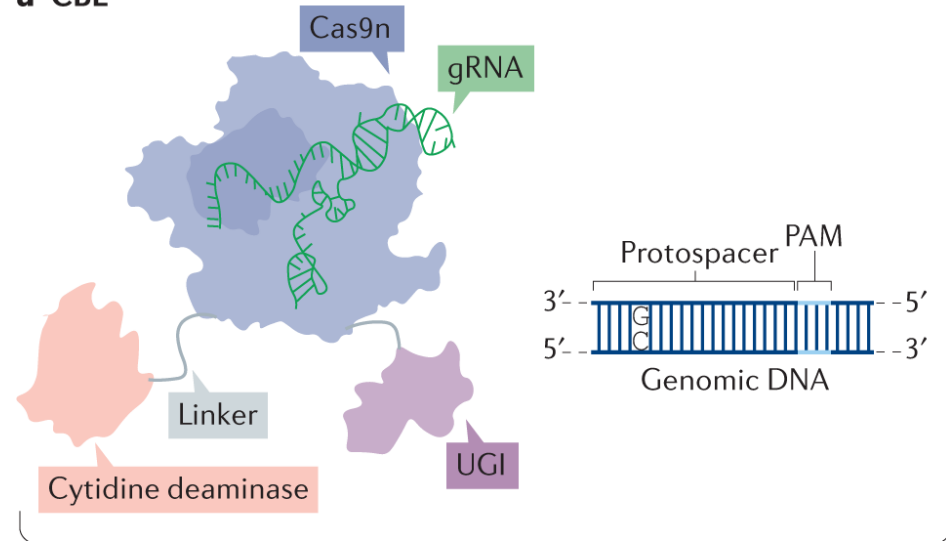
What transmits through the Germline to the Next Generation is Key

Next Generation Editing

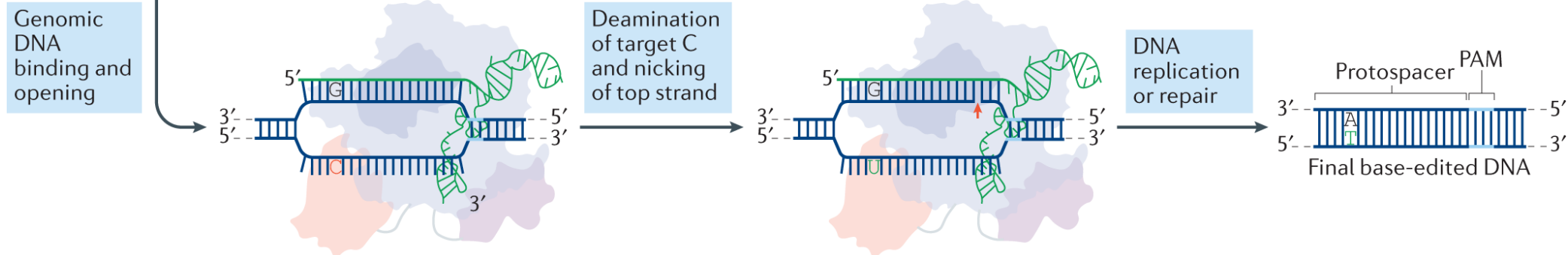
***Expand the Toolbox for
Specificity & Precision***

Base Editing

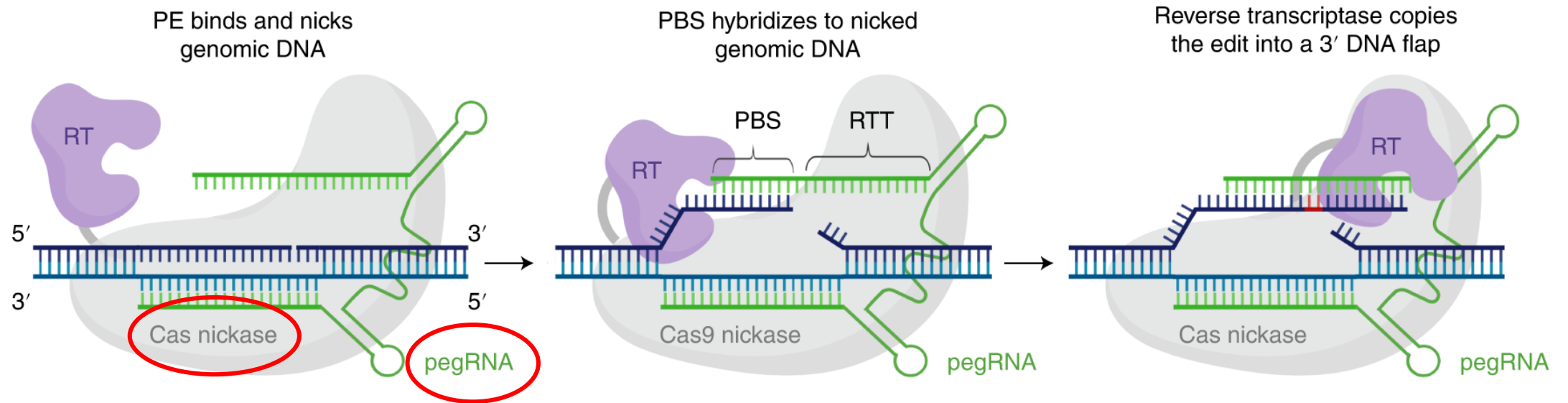
a CBE



+ Potential for engineering precise point mutations
- High rates of off-target editing
- Yet to be tested with animal generation



Prime Editing



❖ **Low efficiency in mammalian embryos**

❖ **Limited reports of success in livestock**

- Zhou et al., 2023, BMC Genomics: ~1% efficiency with sheep embryos

❖ **Recent report of increased efficiency in mice**

- Kim-Yip et al., 2024, Nat Biotech: ~60% efficiency with mouse embryos

Strategies for Generating Food Animals Possessing DNA Edits

Goal

Germline Transmission = Heritable

Current: Edited Embryos

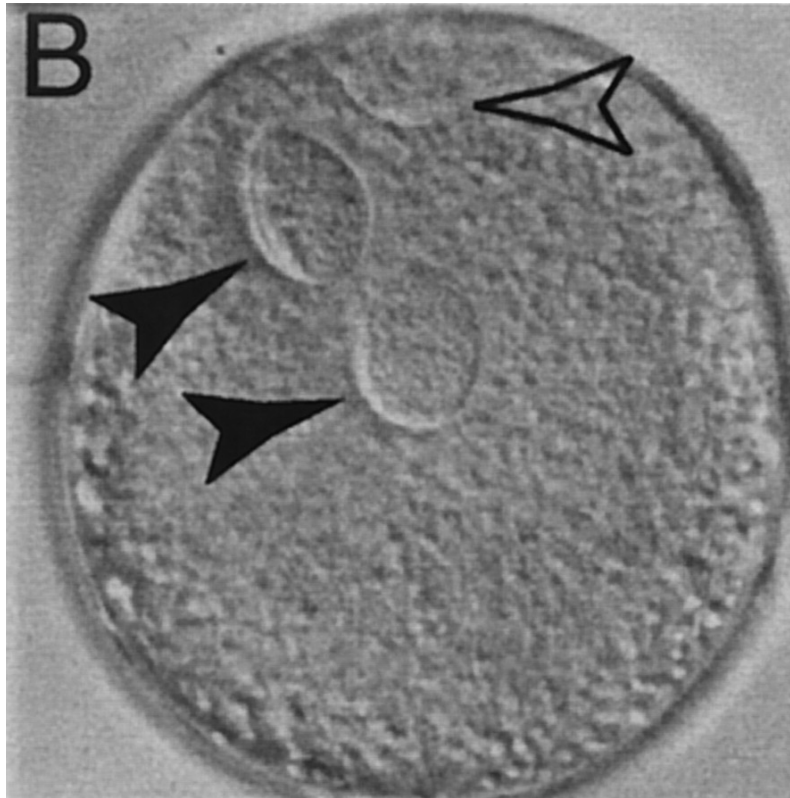


➡ **Pregnancy** ➡ **Offspring
(Founder)**

Zygote Manipulation

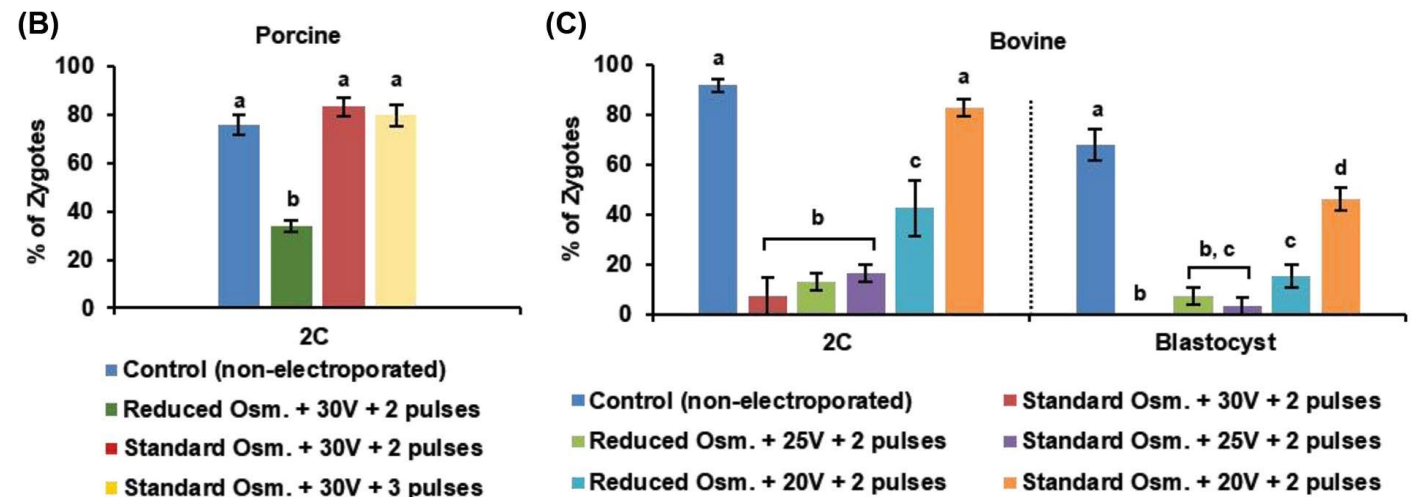
Introduction of Editing Components

Bovine Zygote



Chan et al., 1998, PNAS

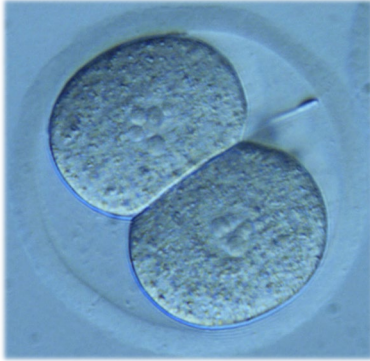
- **Viral Vectors**
 - ❖ Limited utility for livestock
- **Microinjection**
 - ❖ Cytoplasmic deposit but technical skill needed
- **Electroporation**
 - ❖ Simple but not standard across species



Miao et al., 2019, Biol Reprod

2C Microinjection or Electroporation

Introduction of Editing Components



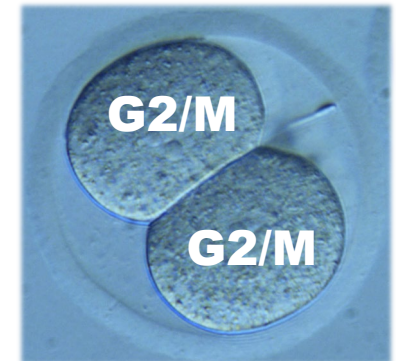
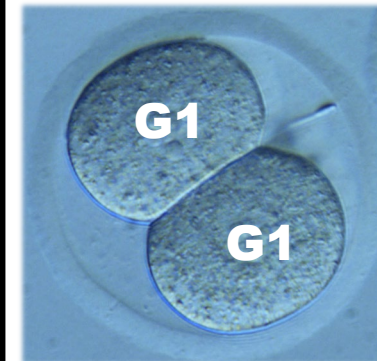
**Higher Rate of HDR
vs
Zygote Manipulation**

**Caveat/Limitation
Increased Mosaicism**

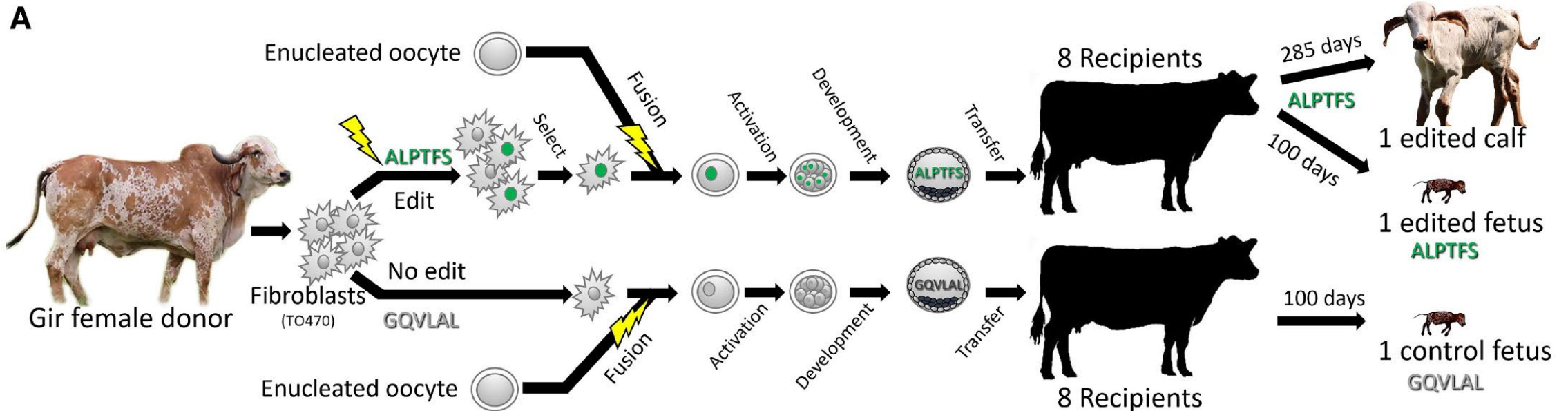
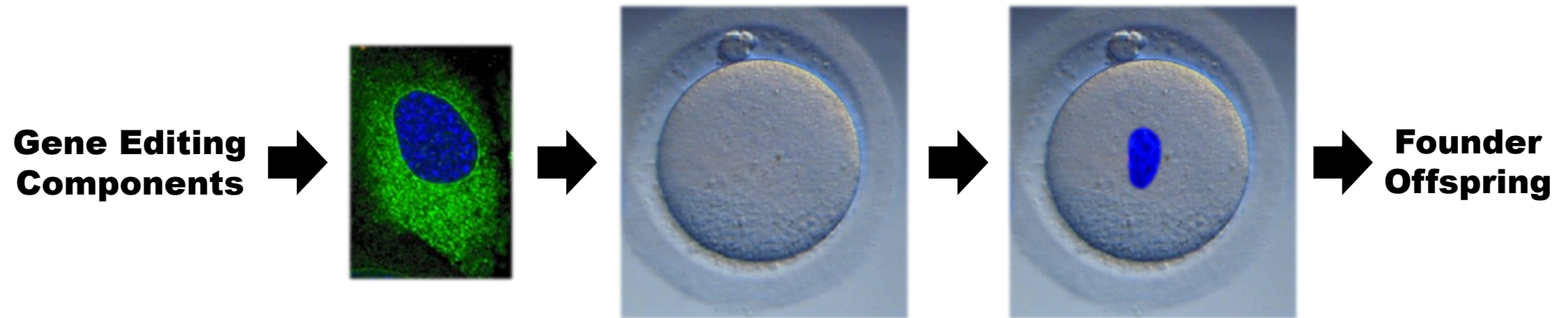
Knock-in	Repair Method	Embryos Transferred	Offspring	+ Founders
<i>Sox2-mCherry</i>	Plasmid	150	32	3 (9%)
<i>Nanog-mCherry</i>	ssDNA	75	25	2 (8%)
<i>Gata6-Halo</i>	ssDNA	60	3	3 (100%)
<i>R26-CAG-H2B-miRFP703</i>	Plasmid	75	22	2 (9%)
<i>Arrdc5-3X FLAG</i>	ssDNA	50	8	4 (50%)
<i>Arrdc5-eGfp</i>	ssDNA	50	12	2 (17%)

2N = 4 Copies

4N = 8 Copies



Editing Somatic Cell DNA + Cloning



Workman et al., 2023, PNAS Nexus

**Regardless of strategy...
intended output is founders
&
Key is Edited Germline**

Conscientious Considerations

Form of Gene Editing Components Introduced to Embryos & Cells

Editors

- ❖ **Plasmid Expression**
 -  **Risk of foreign DNA integration**
- ❖ **Synthesized RNA**
 - ❖ **sgRNA**
 - ❖ **Cas9 mRNA**
- ❖ **Ribonucleotide Protein (RNP)**
 - ❖ **sgRNA + Cas9 Protein**
 - ❖ **Standard for many applications**

Repair Template

- ❖ **Plasmid Based**
 -  **Risk of foreign DNA integration**
- ❖ **Synthesized DNA**
 - ❖ **ssDNA (ssODN) – size limits**
 - ❖ **Long dsDNA**
- ❖ **Adeno Associated Virus (AAV)**
 - ❖ **Efficient for HDR but...**
 -  **Potential for foreign sequence integration (Luqman et al., bioRxiv)**

Mosaicism

- ❖ Genotyping detection of multiple different alleles in Founders
- ❖ Soma vs Germline
- ❖ If germline mosaic, what will transmit to offspring?
- ❖ Only one edited allele will be inherited, but...which one?

CRISPR-Cas9 Editing *NANOS2* in Pigs

c

Boar #	Ear DNA Sequencing	Sperm DNA Sequencing	Genotype
136	5bpΔ/3bpΔ/3bp (insertion)	5bpΔ/3bpΔ/3bp (insertion)	Mosaic
137	1bp (insertion)/1bpΔ	1bpΔ/3bpΔ	Mosaic
142	3bpΔ/5bpΔ	3bpΔ/200bpΔ	Mosaic
143	3bpΔ/10bpΔ	3bpΔ/10bpΔ/5bpΔ	Mosaic
144	1bp (insertion)/4bpΔ	N/A	Knockout
146	4bpΔ/150bpΔ	N/A	Knockout
147	3bpΔ/5bpΔ	N/A	Heterozygous
148	1bpΔ/3bpΔ	1bpΔ/3bpΔ	Heterozygous
251	1bpΔ/21bpΔ	1bpΔ/3bpΔ	Mosaic

Park et al., 2017, Sci Reports

Germline Transmission is Key!



- ❖ Not all founders with edits genotyped in soma transmit through the germline
- ❖ Mosaicism in germline also occurs
- ❖ Litter bearing vs singleton consideration
- ❖ Future is direct germline editing

Knock-in	Method	Embryos Transferred	Offspring	+ Founders	Germline Transmission
<i>Sox2-mCherry</i>	Plasmid	150	32	3 (9%)	8%
<i>Nanog-mCherry</i>	ssDNA	75	25	2 (8%)	47%
<i>Gata6-Halo</i>	ssDNA	60	3	3 (100%)	46%
<i>R26-CAG-H2B-miRFP703</i>	Plasmid	75	22	2 (9%)	41%
<i>Arrdc5-3X FLAG</i>	ssDNA	50	8	4 (50%)	62%
<i>Arrdc5-eGfp</i>	ssDNA	50	12	2 (17%)	100%

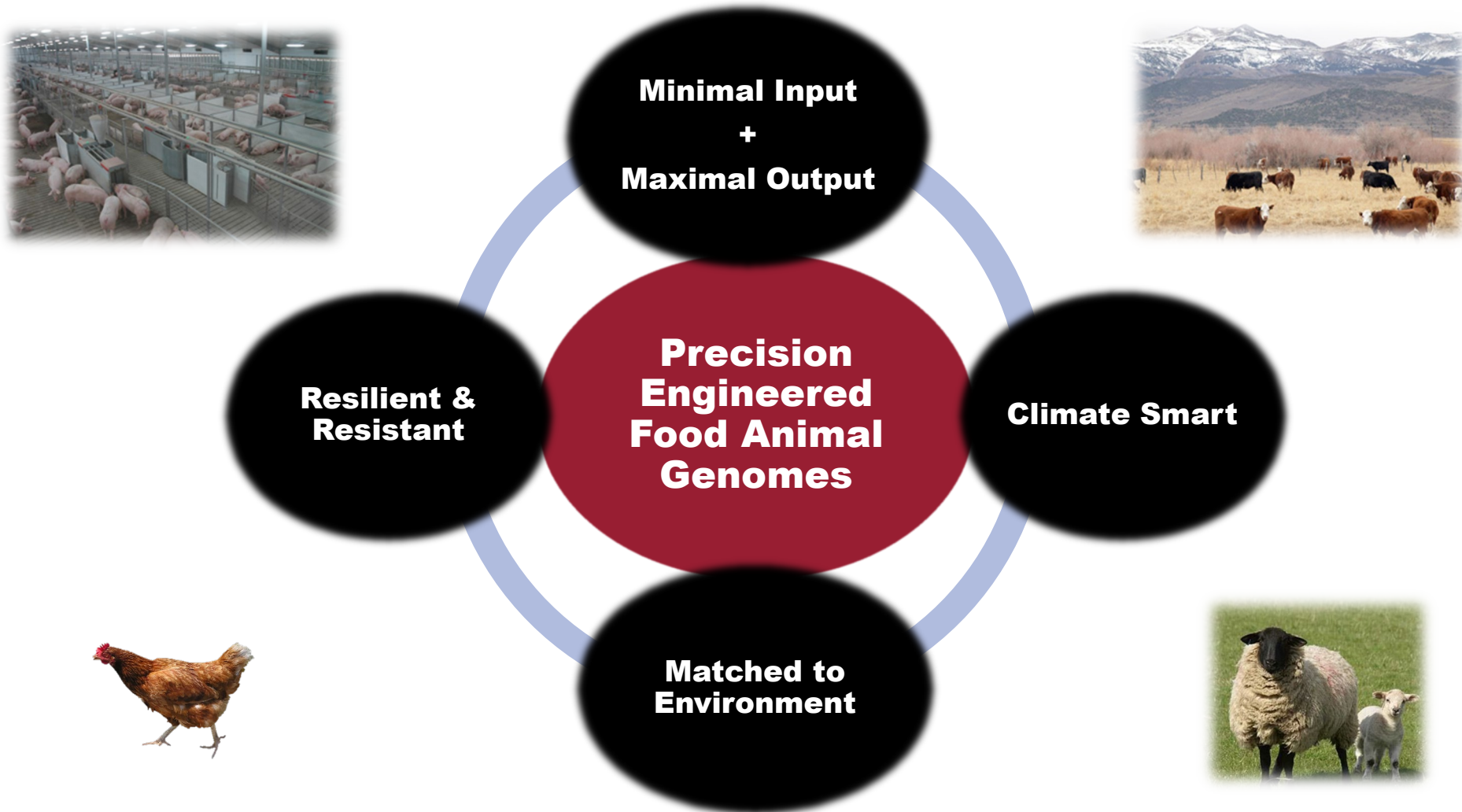
On-Target vs Off-Target Edits

- **Stability of on-target edits**
 - If it does change, how to account for natural evolution of DNA?
- **Potential for off-target editing to reduce animal welfare and compromise food safety**
 - ❖ Do we need to assess?
 - ❖ What will the standard practice be?
 - ❖ Can we really distinguish off-target from random mutations?
 - ❖ Is this an impossible box to check?
 - ❖ Should **phenotype** be the guide?
- **Enhanced strategies may mitigate concern**
 - Cas9 Variants like Cas9-HF1 are reported to reduce non-specific contacts, yet to be tested with livestock

Takeaways

- ❖ **The future of global food security **needs** genome engineering of livestock**
 - ✓ **Gene editing offers the means to engineer **resiliency** and **efficiency****
- ❖ **Primary strategies for livestock are conventional **TALEN** and **CRISPR-Cas9** to generate embryos with **INDEL** and **HDR** gene edits**
 - ❖ **Expected and unexpected DNA modifications can occur**
 - ✓ **Strategy of choice and research design matter**
- ❖ **Science is the **Guide** and the **Solution!****

The Future of Food Security



THANK YOU

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