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Improved Efficiency of CRISPR/Cas9 Genome Editing in Rodent Models Through Optimized Microinjection Components Validation Huanyu Jin, Jinping (James) Luo, Padmaja Tummala, Henrike Siemen, Nga Nguyen, Anthony Thomas, Ronald Basco, Kathy Bang, Jinling Li, Ruby Yanru Chen-Tsai

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### Introduction

- •Animal Models are crucial to understanding genetic mechanisms in vivo settings.
- CRISPR/Cas9 genome editing technology has offered a highly-efficient method for the development of animal and cell models.
- CRISPR/Cas9 technology is highly precise for generating knock-in, knockout and point mutation rodent models.
- However, the efficiency and success rates vary dramatically among different laboratories due to variability in protocols, designing and construction of CRISPR elements.

## Comparing Efficiency of Point Mutation Model Generation Using Cell Line Vs. Embryo Validation Methods

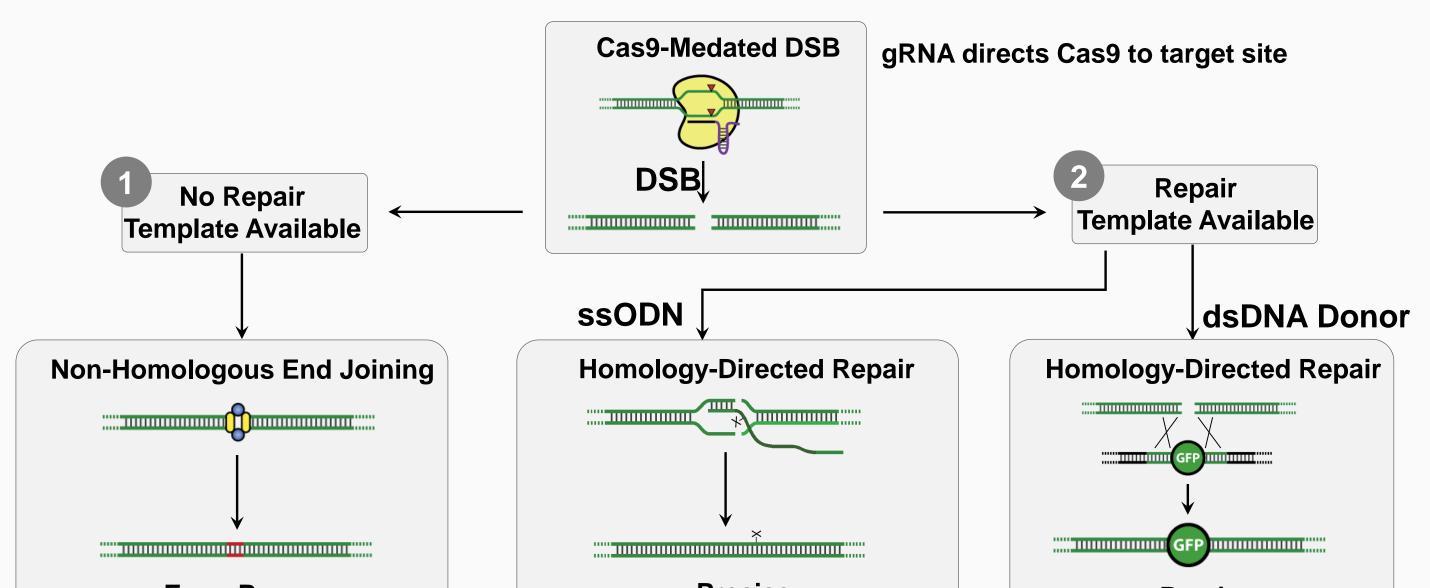
	Cell Line Validation				Embryo Validation			
Gene: Mutation	# Injections	# Total mice born	# Positive founders	% Mod. Efficiency	# Injections	# Total mice born	# Positive founders	% Mod. Efficiency
Autism gene: A350V	3	97	1	1.0%	1	33	13	39.3%
Neuronal gene: S338A	4	73	0	0.0%	1	34	15	44.1%

 Here, we demonstrate that using gRNA validated in embryos (*in vivo*) as opposed to in cells (*in vitro*) offers a consistent and efficient outcomes of CRISPR/Cas9 technology in producing rodent models:

•We compare efficiency of generating two mouse models using our optimized embryo-validated (blastocysts) gRNAs and cell line-validated (N2A cells) gRNAs.

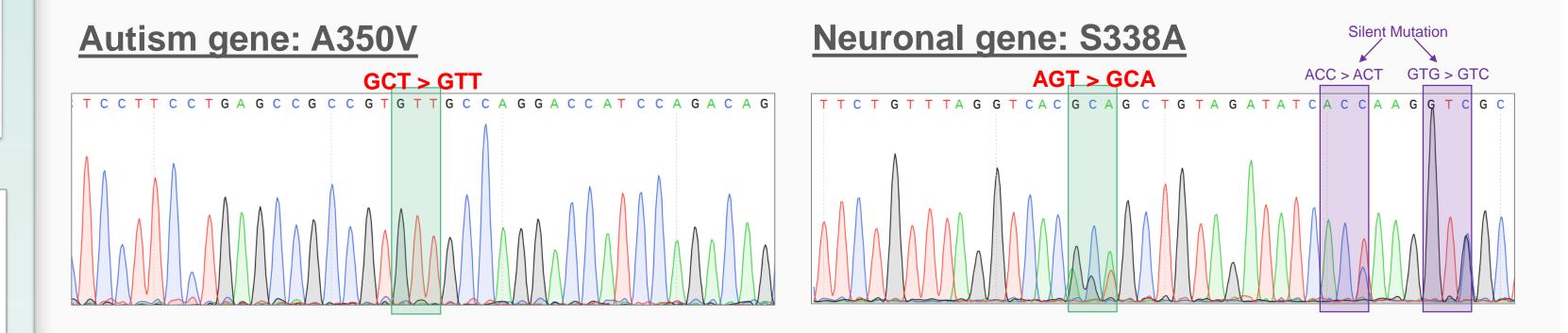
•Since application of this optimized procedure, we have been achieving consistently high efficiency in generating knockout (KO), conditional knockout (cKO), point mutation (PM) and knock-in (KI) mouse (and rat) models.

## **Cas9-Mediated Genome Engineering**



% Modification efficiency = # of positive founders/ # of pups born

 % Modification efficiency in projects using cell line validated gRNAs is significantly lower compared to projects using embryo-validated gRNAs



**Figure.** Sequence chromatogram of representative founder mice generated using embryovalidation gRNAs

### CRISPR/Cas9-Mediated Gene Editing in Mouse Models Using Embryo-Validated gRNAs

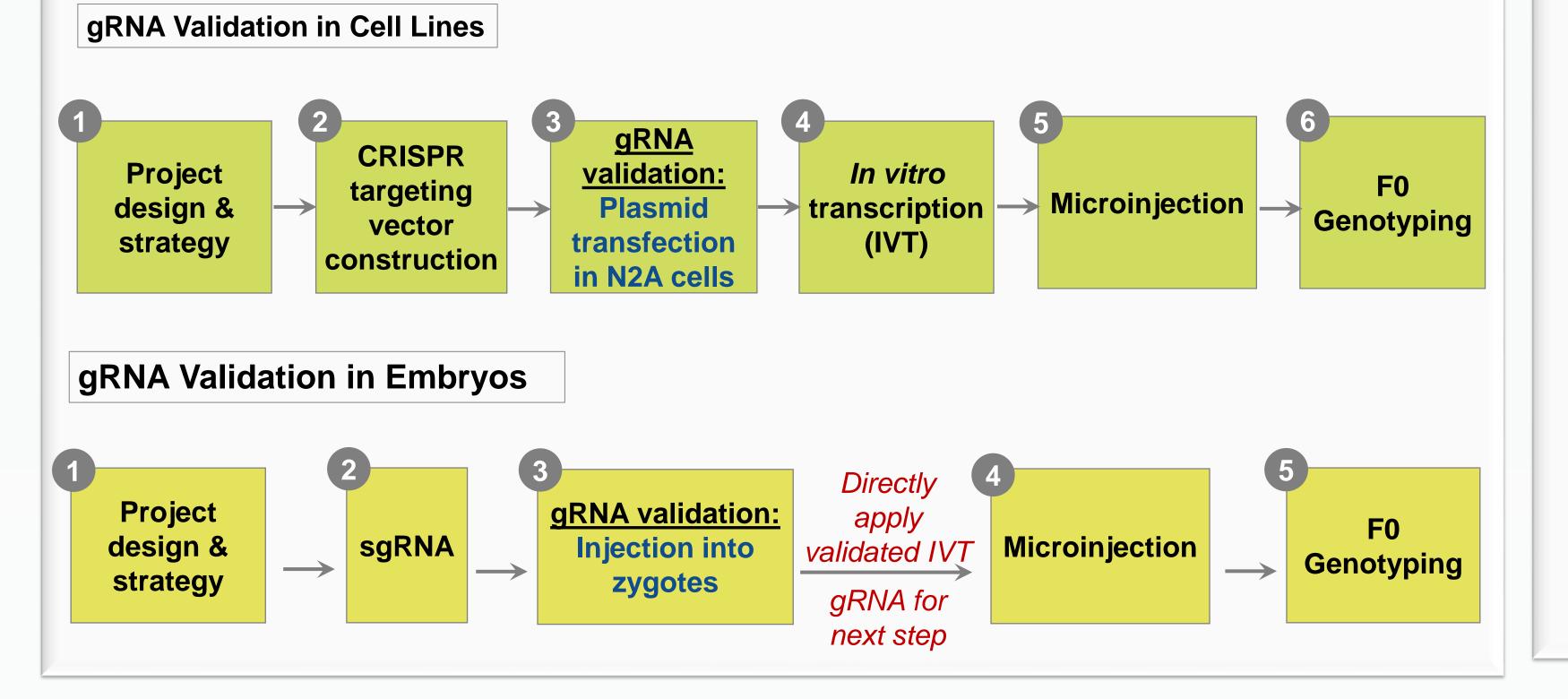
# Embryo # Embryo # Pups Born # Positive

Error-Prone Often Results in Indel Formation

#### Precise Point Mutation, Precise INDEL, Tag

Precise Reporters, Cre, Splice Isoforms

## Experimental Design for Embryo-Validation & Cell Line-Validation of gRNAs



Models	Injections	Transfers (%)	(%)	Founders (%)
KO*	144	83 (57.6)	17 (20.5)	9 (52.9)
PM	156	126 (80.8)	37 (29.4)	11(29.7)
cKO-1 <sup>st</sup> LoxP	120	88 (73.3)	26 (29.5)	13 (50.0)
cKO-2 <sup>nd</sup> LoxP (retargeting**)	213	147 (69.0)	29 (19.7)	6 (20.7)***
KI	256	178 (69.5)	33 (18.5)	8 (24.2)

\* Knockout (KO) models are generated using a donor-free targeting scheme.

- \*\* Conditional knockout (cKO) mouse models are generated by inserting two LoxP sequences using a sequential targeting strategy: (1) The 1<sup>st</sup> LoxP site is inserted either upstream (5') or downstream (3') of the region that needs to be floxed; (2) The single LoxP+ mice thus generated are used to produce zygotes for re-targeting and inserting the 2<sup>nd</sup> LoxP; (3) If heterozygous LoxPs are present at both locations, it is necessary to breed the founders with WT mouse to confirm the LoxPs are on the same allele.
- \*\*\* Efficiency of generating cKO mice with two LoxP sites inserted at targeting site (double positive)

### Comparing gRNA Validation in N2A Cells Vs. Blastocysts

### **Conclusions and Discussion**

Criteria	N2A Cells	Embryos/ Blastocysts	
gRNA used for validation	Plasmid	In vitro transcribed sgRNA	
gRNA used for injection	sgRNA (different from validation)	sgRNA (same as validation)	
Background mouse	Mouse neuroblastoma cells	Same as final mouse model	
Validation timeline	~4-6 weeks	1-2 weeks	
Correlation between validation and final mouse model	No direct correlation because of different types of cells	Direct correlation resulting in ~100% gRNA/Cas9 modification	
Work procedure	More steps needed: build plasmid	Fewer steps	
# Injections	Unpredictable; More injections needed	Highly predictable	
Efficiency	Lower efficiency	Higher efficiency	

- Validation of gRNAs is a key parameter for successful genome modification in mouse (and rat) models when using the CRISPR/Cas9 technology.
- Validation of gRNA should be done in the embryos/blastocyst isolated from the same strain in which the desired animal model is to be generated.
- For generating the final animal model, successful gRNAs should be in the same form as that used for the validation assay. This embryo-based validation method also serves as a quality control step for the reagents being used for microinjection.
- With embryo-validation there is a strong correlation between gRNA activity and modification efficiency: higher gRNA activity results in higher efficiency of genome modification.
- Validation of gRNA in blastocysts is a fast and efficient method to provide consistent and higher efficiencies for genetically engineering mouse models.

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