BNORC Transgenic Core

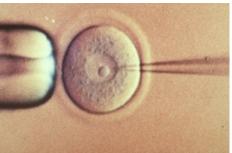
Mouse Genetic Engineering made "Easi"

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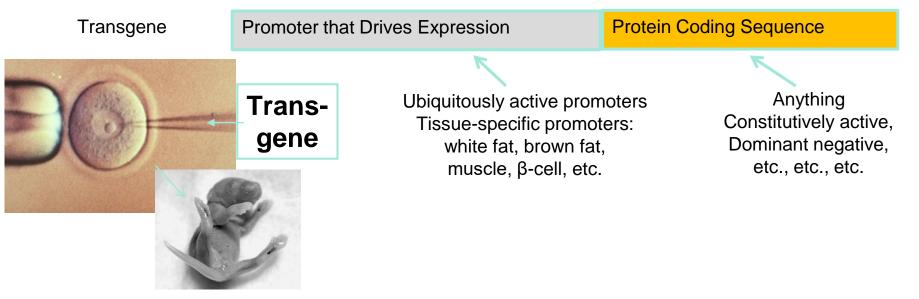
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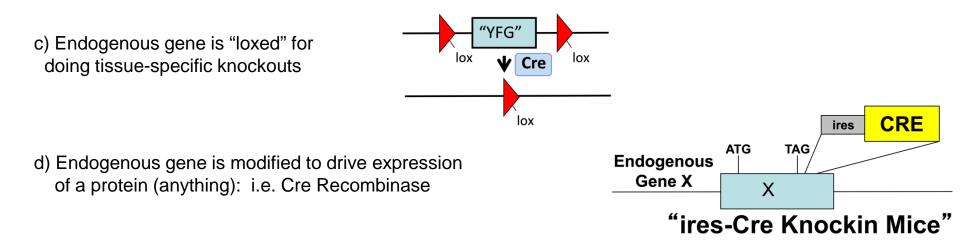
Two general types of mouse genetic engineering

1. Addition of exogenous DNA to the mouse genome – a.k.a. "Transgenics".

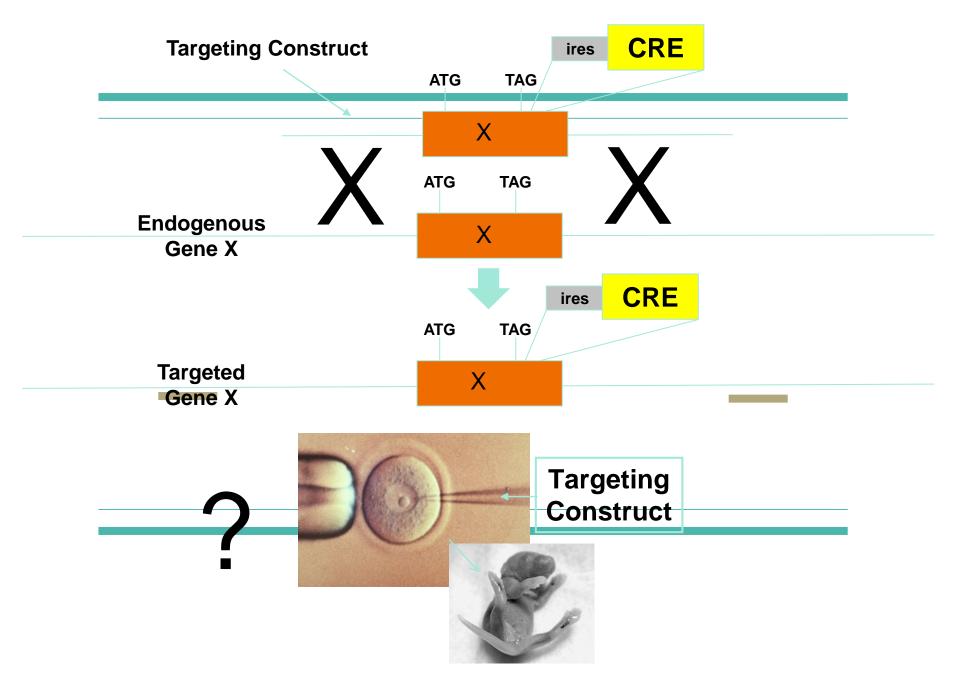


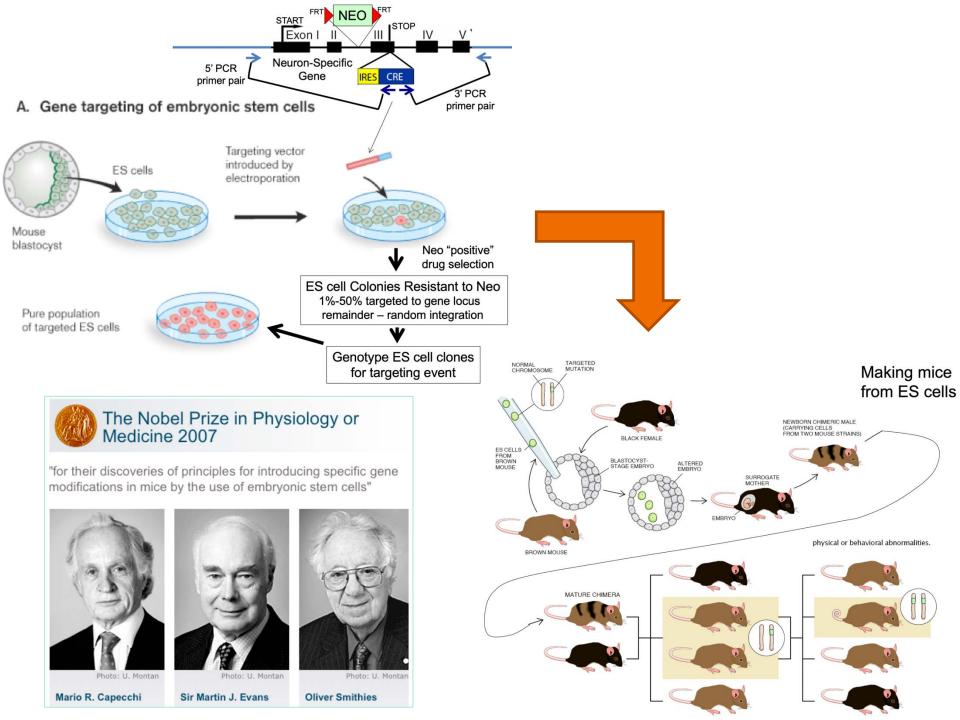
2. Targeted manipulation of the mouse genome – a.k.a. "Gene Targeting".

- a) Endogenous gene is made nonfunctional "global gene knockout"
- b) Endogenous gene is subtly altered i.e. mimic a human point mutation, etc.

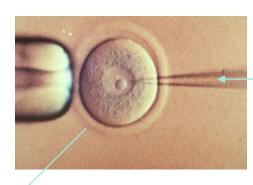


Gene Targeting – key role of Homologous Recombination





Easi-CRISPR: rapid, easy Gene Targeting

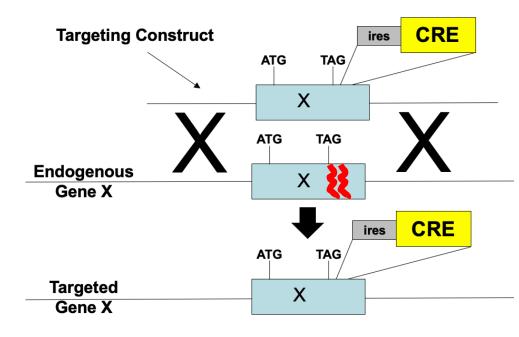


Targeting Construct and CRISPR "reagents"



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- 1. CRISPR is used to create break in DNA:
- \uparrow 's rate of homologous recombination (HR).
- 2. A **ssDNA** targeting construct is used:
 - no random integration.
 ↑'s rate of HR.



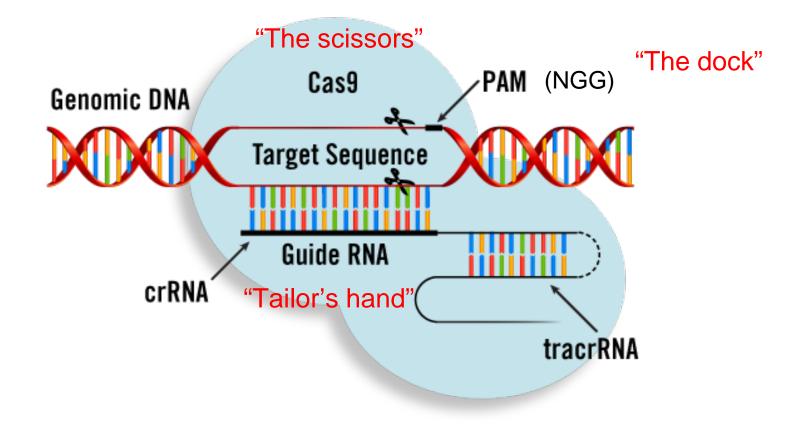
Easy muse gene targeting with Easi-CRSPR

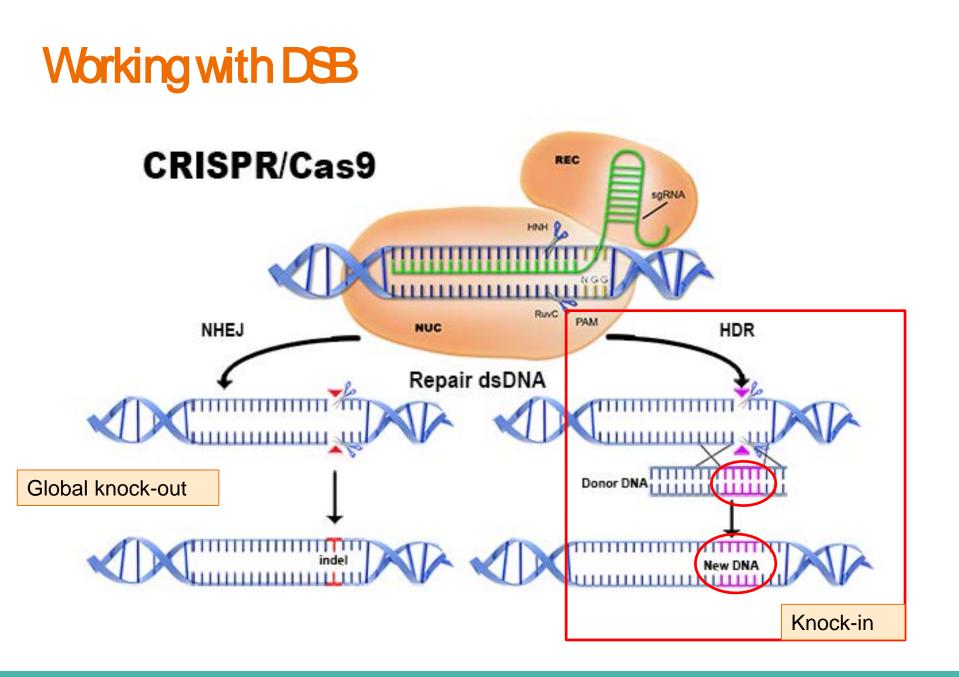
Chen Wu, PhD Co-director, BNORC Transgenic core Instructor in Medicine, BIDMC <u>cwu9@bidmc.harvard.edu</u>, 617-735-3259

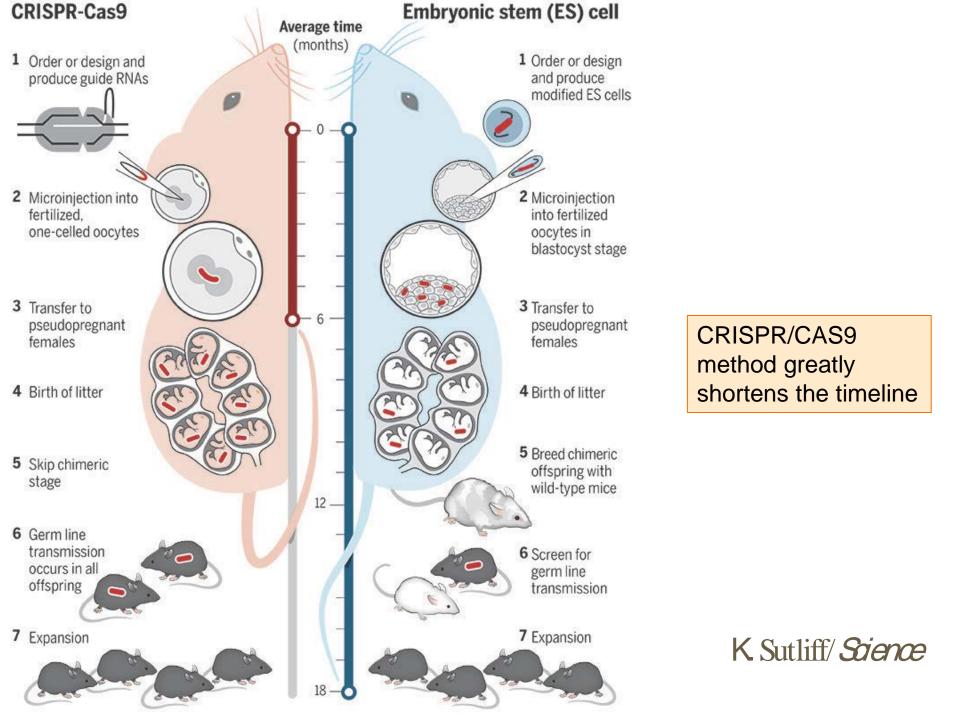
An overview of CRSPR technology

Abrief overview of CRSPR technology

To create double-stranded DNA break









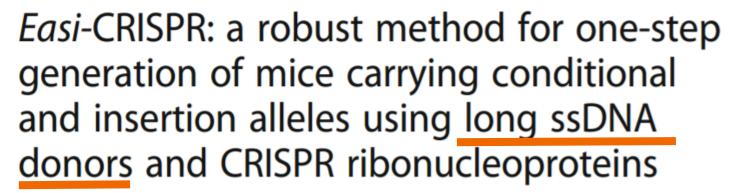
Quadros et al. Genome Biology (2017) 18:92 DOI 10.1186/s13059-017-1220-4

Genome Biology

RESEARCH



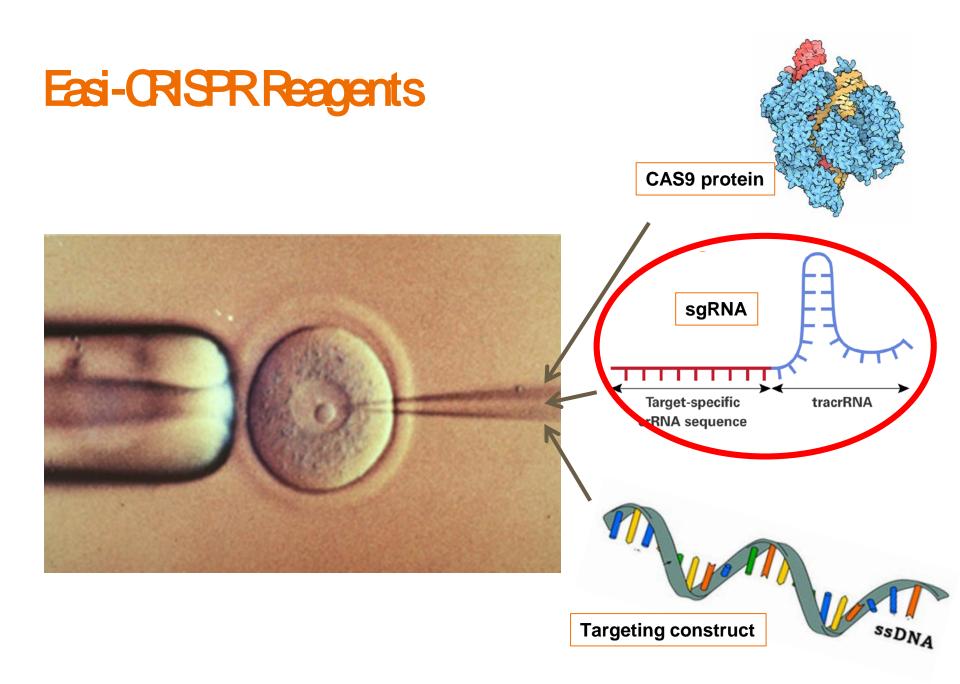
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Rolen M. Quadros^{1†}, Hiromi Miura^{2,3†}, Donald W. Harms¹, Hisako Akatsuka^{2,4}, Takehito Sato⁴, Tomomi Aida^{5,6,7}, Ronald Redder⁸, Guy P. Richardson⁹, Yutaka Inagaki^{3,10,11}, Daisuke Sakai^{10,12}, Shannon M. Buckley^{13,15}, Parthasarathy Seshacharyulu¹⁴, Surinder K. Batra^{14,15}, Mark A. Behlke¹⁶, Sarah A. Zeiner¹⁶, Ashley M. Jacobi¹⁶, Yayoi Izu¹⁷, Wallace B. Thoreson¹⁸, Lisa D. Urness¹⁹, Suzanne L. Mansour^{19*}, Masato Ohtsuka^{2,3,10*} and Channabasavaiah B. Gurumurthy^{1,20*}

Comparison with conventional CRISPR

	Conventional	Easi-CRISPR
Targeting construct form	Plasmid (dsDNA)	ssDNA
Targeting construct availability	Cloning	Chemical synthesis
Homology arm length	> 2.5 kb	100 nt
Genotyping	Long range PCR / Southern blot (difficult)	Regular PCR (easy)
Success rate	< 10%	25% (and counting)



sgRNA design: in silico

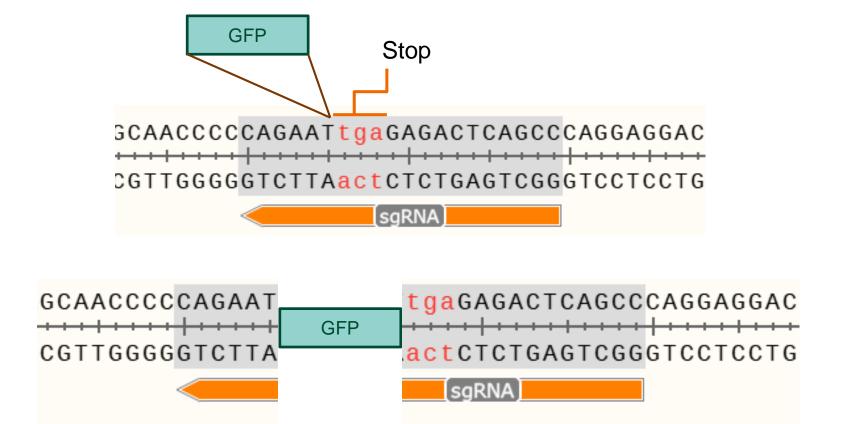
- Institutional sites: Broad, MIT, crispor.org
- Commercial sites: IDT, Synthego, etc.
- Cloud-based bioinformatics sites: Benchling.com

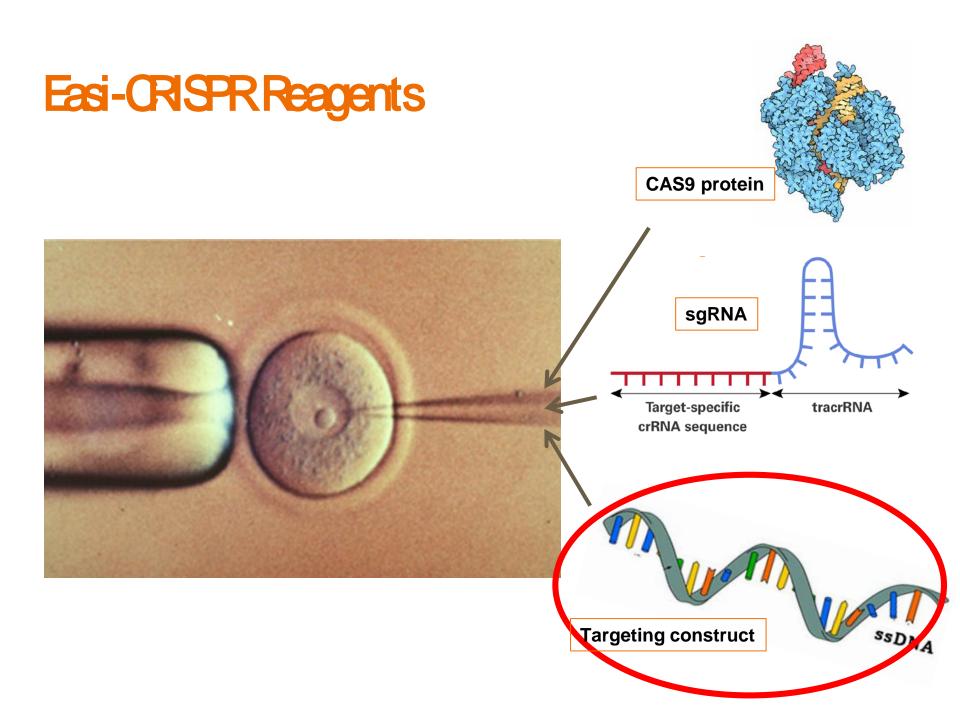
Online predicting algorithms are based on analysis of 3000+ sgRNAs (Doench et al. 2014 and 2016)

Criteria of a good sgRVA

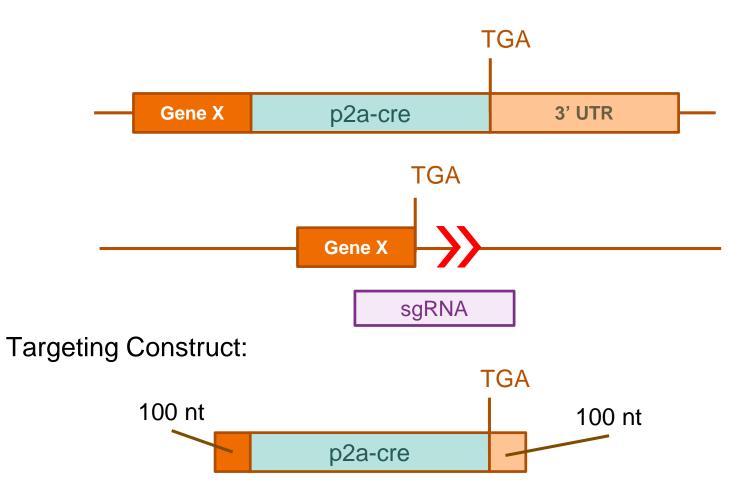
- Efficacy or on-target score cuts where it should
- Specificity or off-target score does not cut where it shouldn't
- Close to the insertion site

The ideal sgRVA site: contains the insertion site

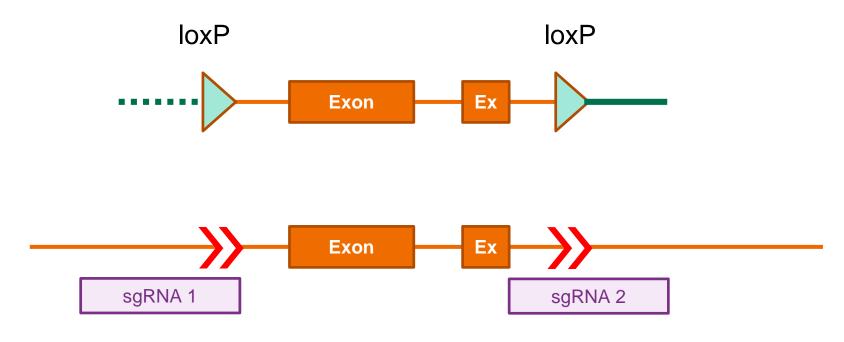




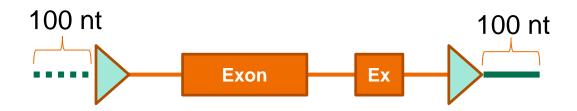
Project design (I) - reporter or driver



Project design (II) - floxed exon(s)



Targeting construct:



Easi-CRSPR limitation: template availability

- 1. size: 2kb (IDT), 5kb (Genewiz)
- 2. Sequence composition: GC content, repetitive sequence, etc.
- 3. Cost: 90 cents/nt
- 4. Time: ~ 1 month

Easi-CRISPRWorkflow

- 1. Contact us
- 2. Design sgRNA and targeting construct
- 3. Order the reagents
- 4. Send the reagents to us
 - 4 weeks later...
- 5. Pick up the tail snips and genotype
- 6. Breed the founder(s)

Project summary



- Turn-around time: 3 months or sooner (from reagents to founder)
- Average success rate (Easi-CRISPR): 25% (71% the highest)
- Number of successful projects : 50+ within 2.5 years
- Including 4 co-injections: two targeting constructs with one sgRNA
- We provide free consultation on strategy and design

Questions and comments?

Thank you!!



Joel Lawitts, PhD Director, Transgenic core