



Weatherall
Institute of
Molecular
Medicine

Radcliffe Department of Medicine

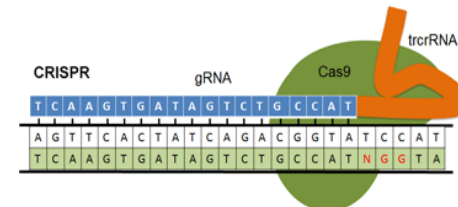
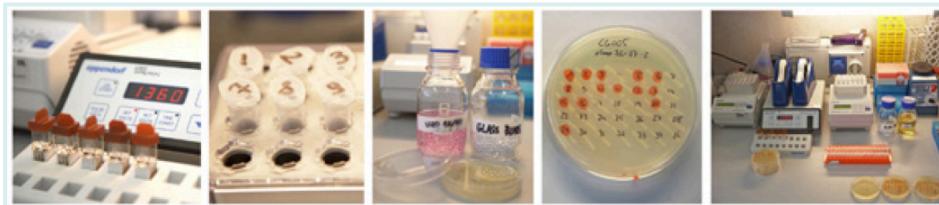


Engineering Genomes

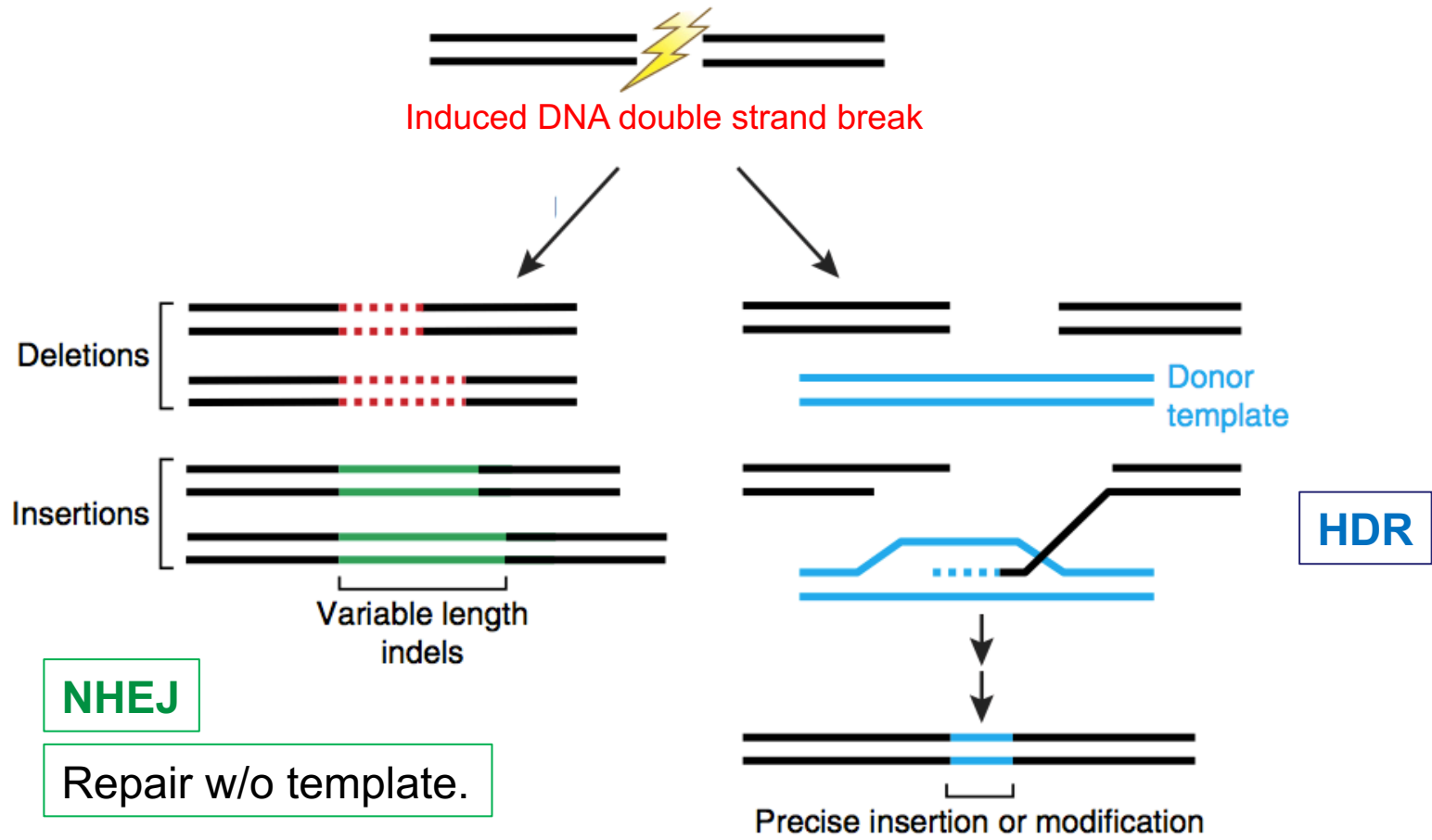
An Introduction to CRISPR/Cas9 Technology

Weatherall Institute of Molecular Medicine,
D.Phil. Course Oxford
November 14th, 2019

Philip Hublitz, PhD
WIMM, Genome Engineering Services

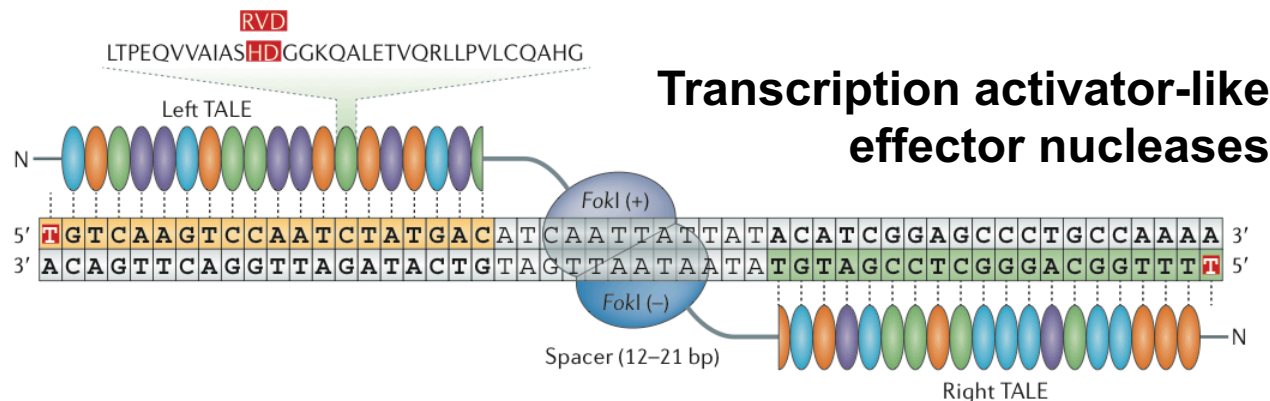
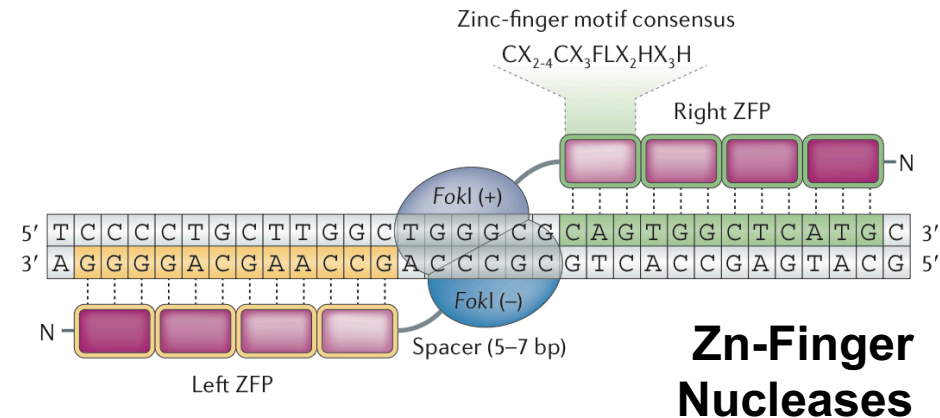


Cellular DNA Repair Pathways

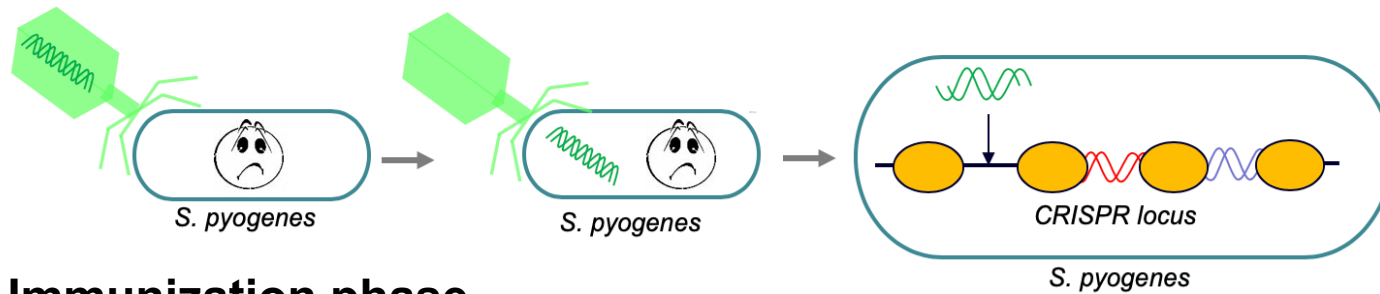


Pre-CRISPR Times

	ZFNs	TALENs
DNA targeting specificity determinant	Zinc-finger proteins	Transcription activator-like effectors
Nuclease	FokI	FokI
Success rate[‡]	Low (~24%)	High (>99%)
Average mutation rate[§]	Low or variable (~10%)	High (~20%)
Specificity-determining length of target site	18–36 bp	30–40 bp
Restriction in target site	G-rich	Start with T and end with A (owing to the heterodimer structure)
Design density	One per ~100 bp	At least one per base pair
Off-target effects	High	Low
Cytotoxicity	Variable to high	Low
Size	~1 kb × 2	~3 kb × 2

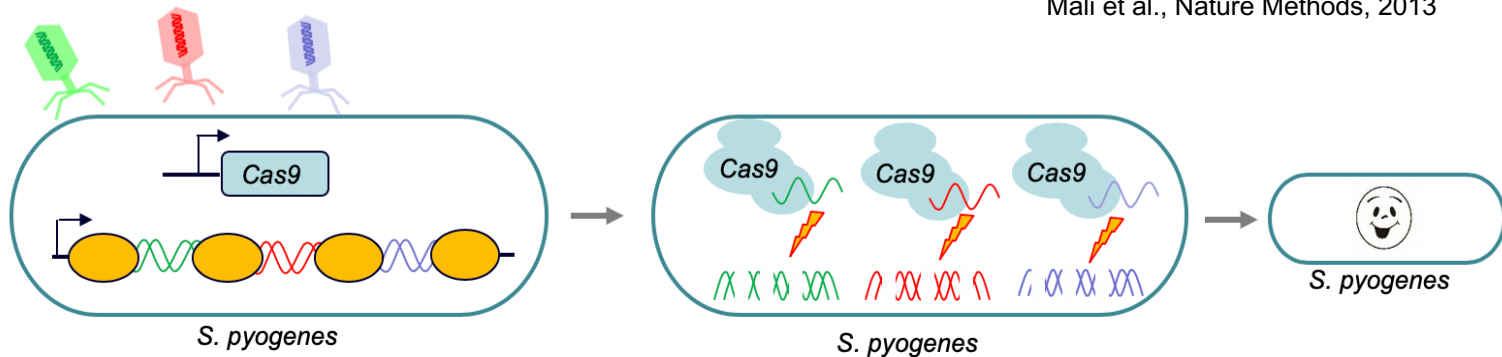


CRISPR: an Innate Immune System



Immunization phase

Ben Davis and Addgene
Mali et al., Nature Methods, 2013



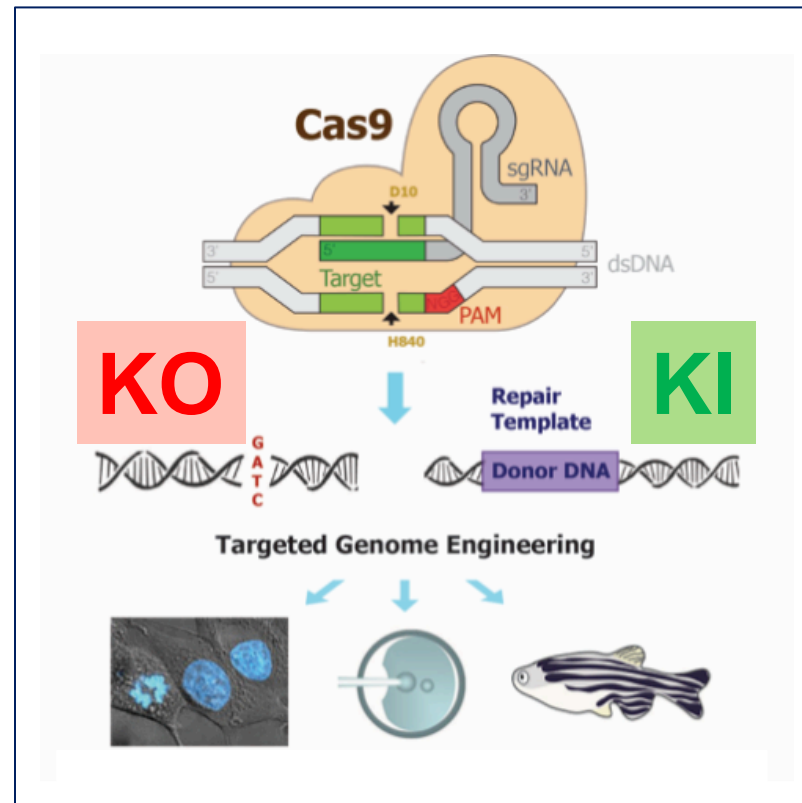
Immunity

Clustered **R**egularly **I**nterspersed **S**hort **P**alindromic **R**epeats

CRISPR/Cas9



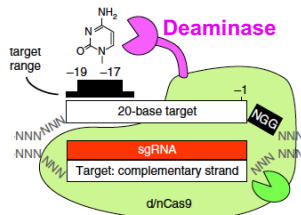
Nature, March 6th, 2014



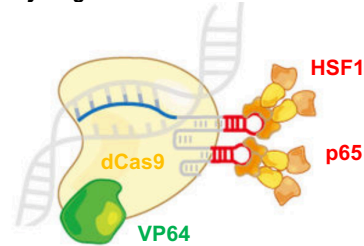
WIMM Genome Engineering, Website, 2018

CRISPR/Cas: What is it good for?

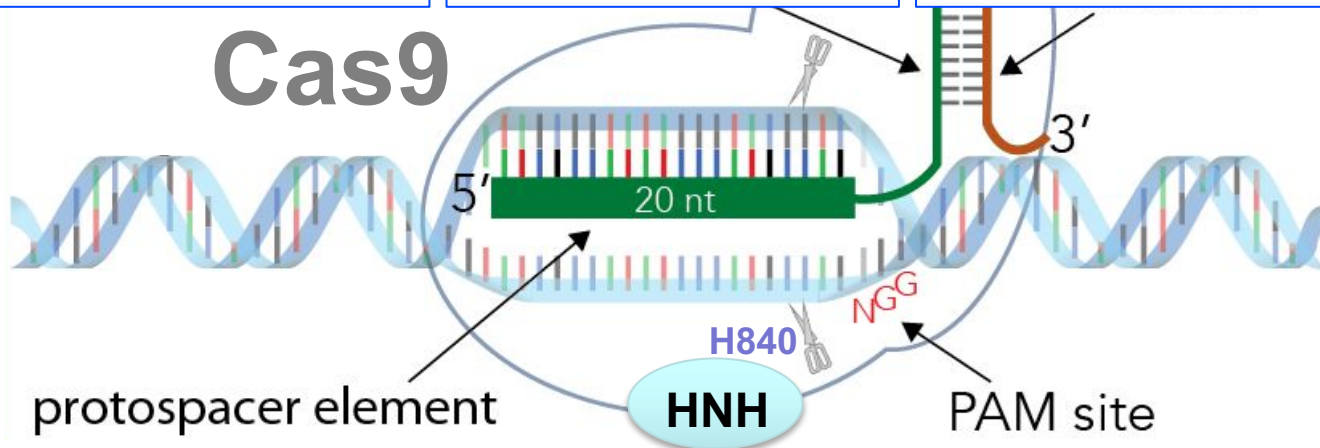
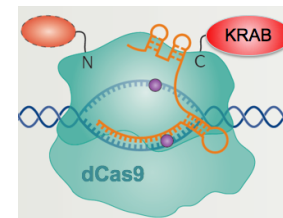
Base Editors Targeted SNP Generation



Assembled SAM complex Synergistic Activation Mediators



CRISPRi Active Repression



CRISPR/Cas: What is it good for?

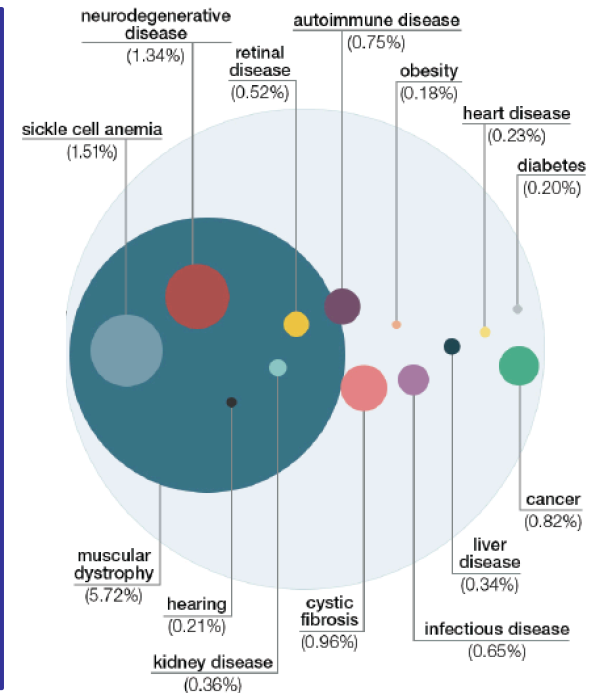
Generation of **LOF/GOF models**: engineering cells and organisms, attempting to cure diseases.



Nature, 2015



Nature, 2016

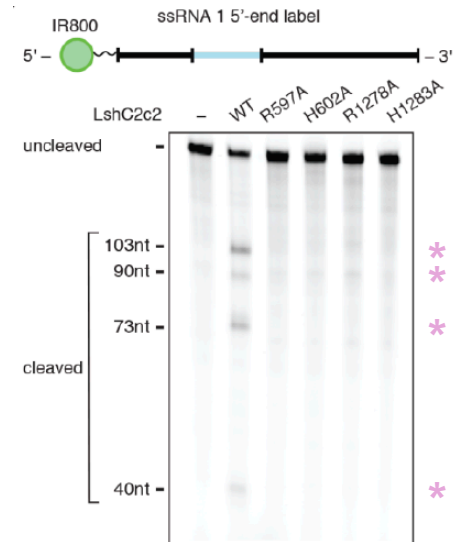
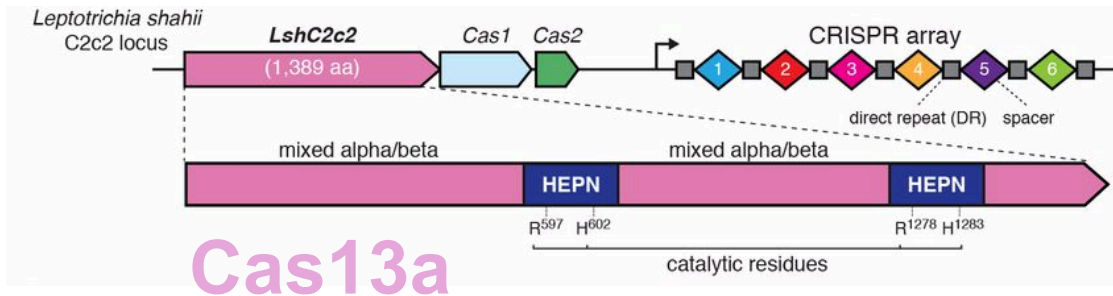


Research Arc Cell Press, 2018

CRISPR/Cas: What is it good for?

Generation of LOF/GOF models.

mRNA editing:



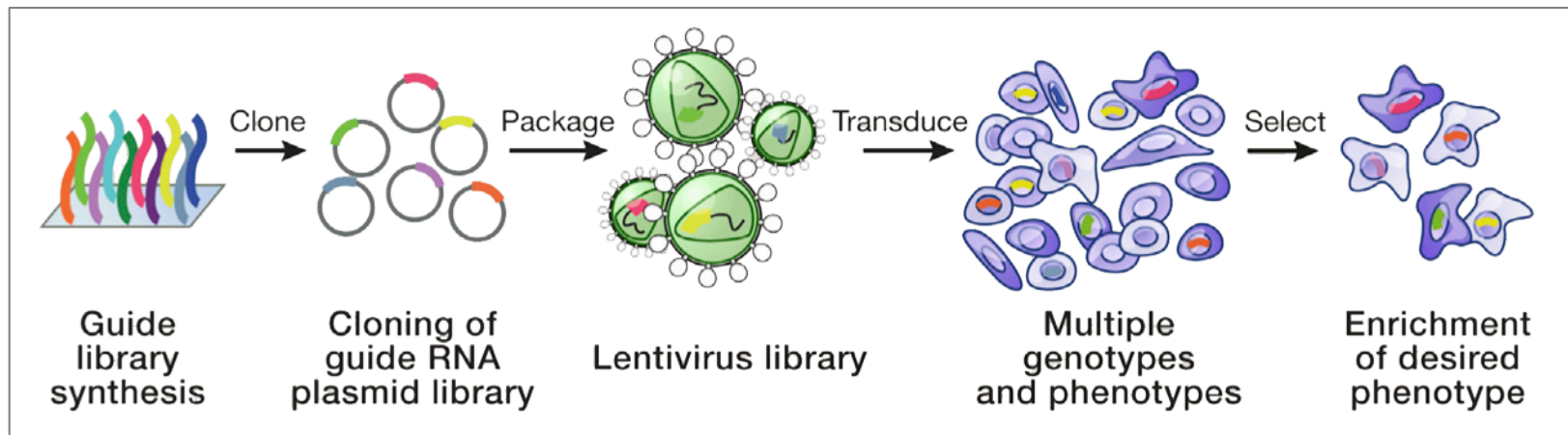
Abbudayyeh *et al.*, Science, 2016

CRISPR/Cas: What is it good for?

Generation of LOF/GOF models.

mRNA editing.

High-throughput screens: KO, CRISPRi, CRISPRa.



Hsu, Cell, 2014

CRISPR/Cas: What is it good for?

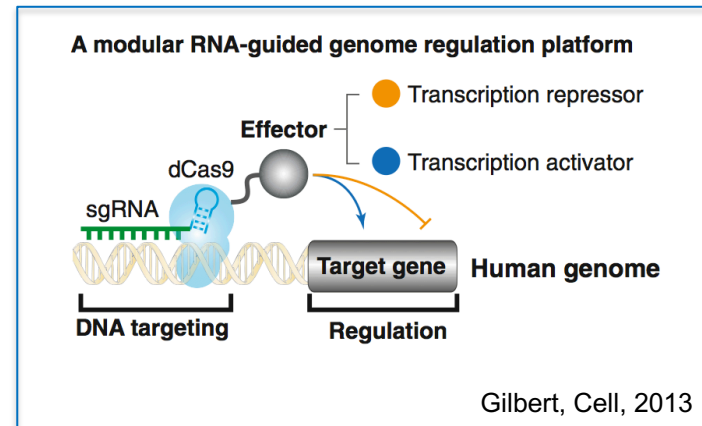
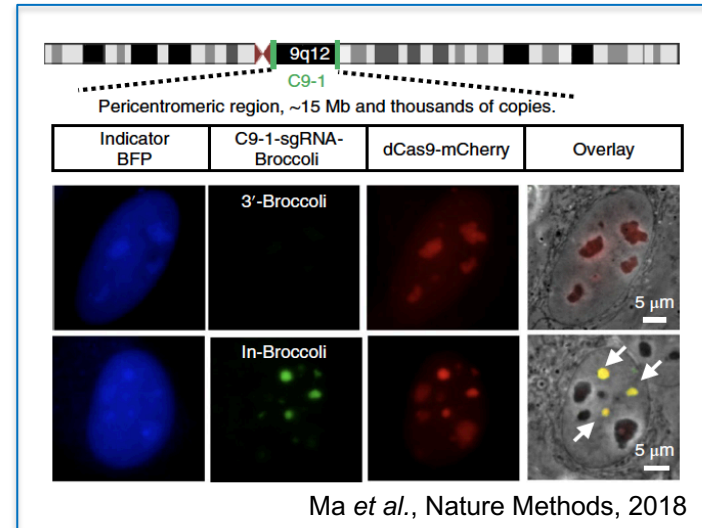
Generation of LOF/GOF models.

mRNA editing.

High-throughput screens.

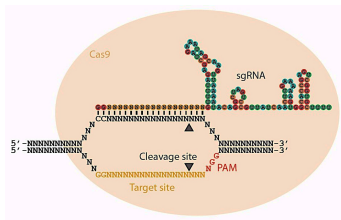
Targeting fusion proteins:

- Labeling
- Gene activation
- Gene repression
- Epigenetic remodeling
- **Base editing**

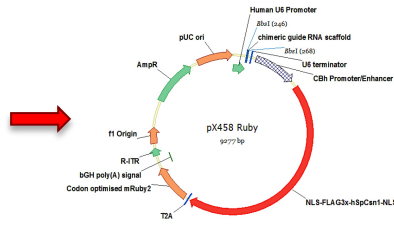


Pipeline

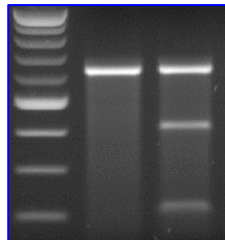
sgRNA design



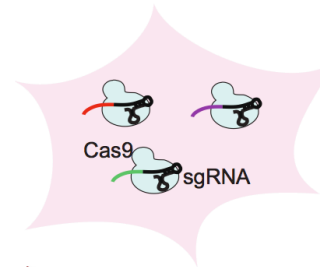
Vector preparation RNP assembly



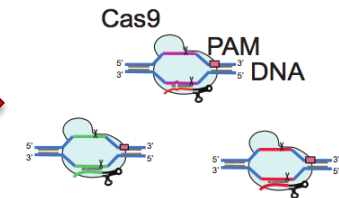
Guide validation



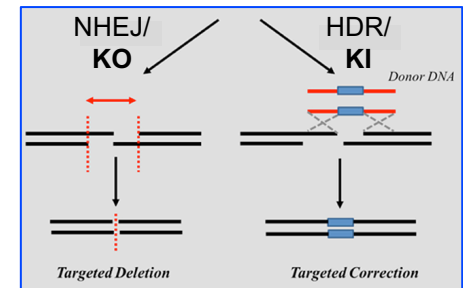
Delivery



Target Cleavage



Repair



SCREEN

HELP and ADVICE: come and see me... ;-)

Thoru Pederson: **U6**, **H1** & **7SK** need a G at start (3 in WT).

Feng Zhang: add a G at 0, it won't disturb. Replacing 20 and 19 with G's (might) affect Cas9 specificity. Keith Joung: shorter guides (18bp) are better and have less off-target cleavage. Jin-Soo Kim: double G at start is perfect.



Fei-Ann Ran, Nature Protocols, 2014

CACC G GNN NNN NNN NNN NNN NNN NN
C CNN NNN NNN NNN NNN NNN NN **CAAA**

19nt

“Standard design” for guides to be used in transient transfections into cells.

Hublitz and many others, unpublished

Consideration: Cas9 Source

mRNA as plasmid

Txn/TIn takes time.
No go in oocyte.
Good for cell lines.

Gradual action.
Cheap and easy.

Stable for several cell
divisions: **OFF-targets.**

Pretty efficient.

mRNA as mRNA

TIn takes time.
Risk of mosaicism
in PNI.

Gradual action,
double modification?
(e.g. double LoxP).

Rapidly degraded.

Variable efficiency?

Protein

Immediate action.
Good for NHEJ and
HDR in oocytes.

Multiplexing difficult:
deletion vs. double
repair.

Rapidly degraded.

Highly efficient.

Consideration: sgRNA Source

Plasmid

Cheap.

Persists for several cell divisions: **OFF-target issues**.

Selection possible by 2A-GFP *et al.* coupling.

Good efficiency.

SygRNA (RNP)

Slightly more expensive.
1 or 2 components.

Rapidly degraded,
Immediate action.
High efficiency.

Labeled **tracrRNA** not functional → labeled Cas9 might work sooner or later...

Good efficiency.

SygRNA /Cas9 protein **RNPs** as the reagent of choice.

It all depends on the experimental approach, though.

CRISPR/Cas9 in Cell Lines

Test **3 sgRNAs** per target site (genome and epigenome).

Test cells thoroughly for **ploidy** and eventual **SNPs**. Find out if they can be **cloned** (crucial for establishment of cell lines).

Establish **optimum transfection conditions**: efficiency vs. viability, delivery method (LPF, Amaxa, Neon), allow recovery for 48 to **72h** (test for maximum clonal survival rate).

Selection by Cas9-2A-**eGFP/BFP/Ruby**/etc. by FACS or use of Cas9-2A-**Puro/Bla/Gen** for enrichment culture. Isolation of single cell clones.

Screen: Analyze experimental clones by method of choice.

N.B.: **All off-target modifications are contained in the cell.**
Establish ≥ 3 independent lines for phenotype comparison.