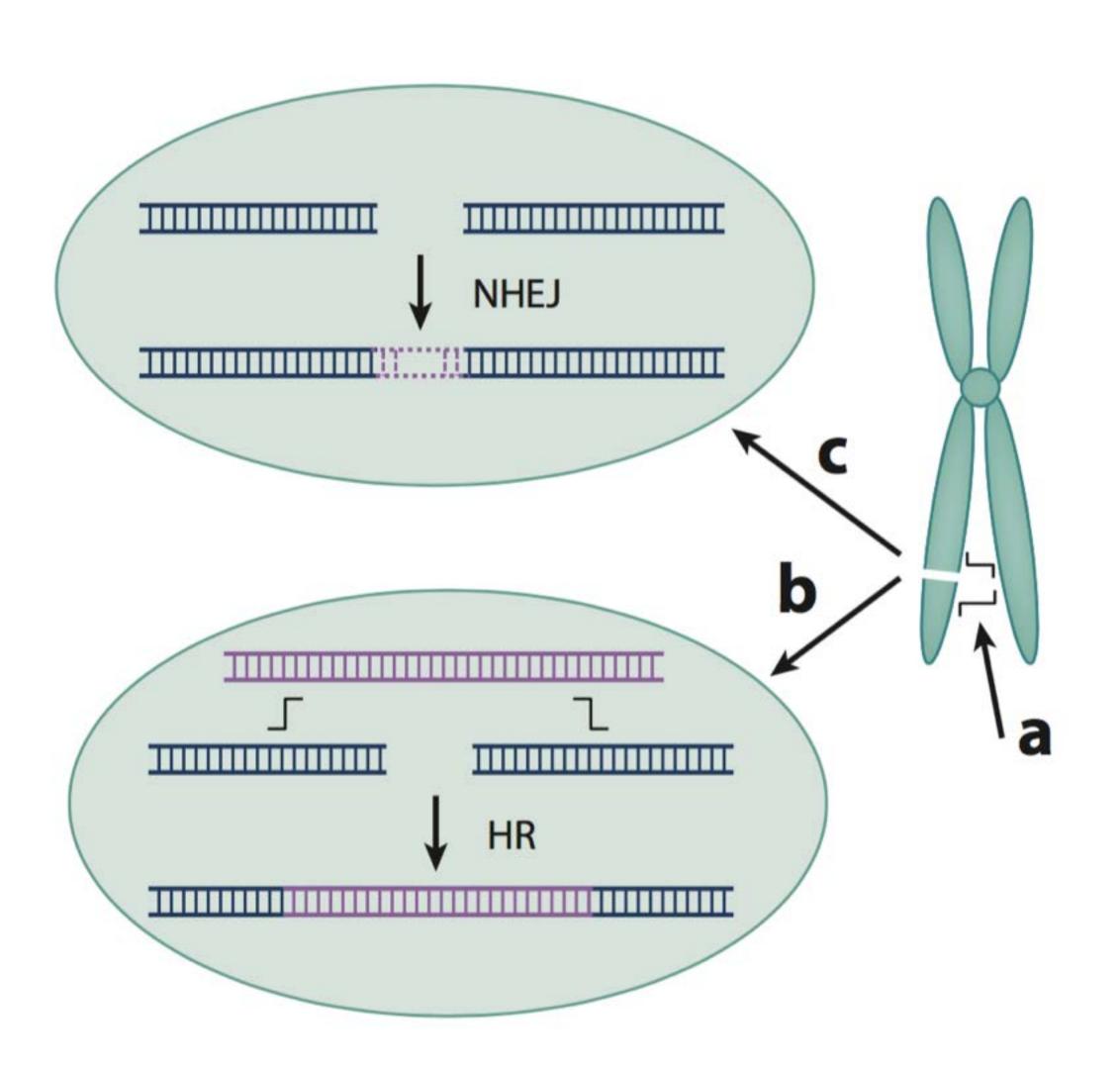
# Genome editing: clinical applications



### Double-strand break repair



D. Carroll

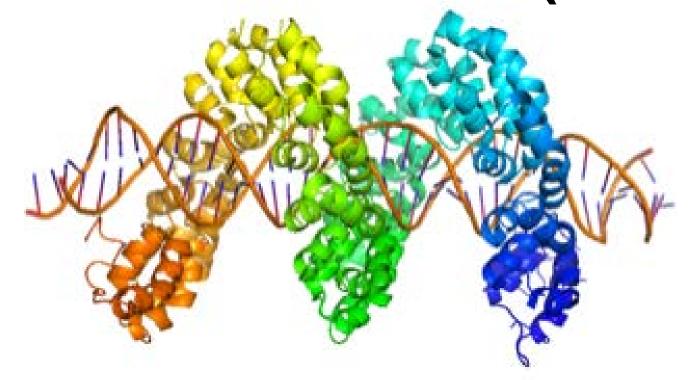
#### Genome-Editing Nucleases

#### Zinc Finger Nucleases (ZFNs)

**TALE Nucleases (TALENs)** 

Sangamo / Biogen



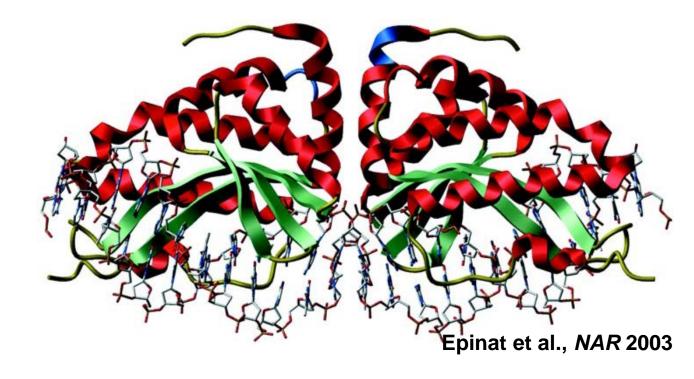


Cellectis / Pfizer

#### Meganucleases

#### **CRISPR RNA-Guided Nucleases**

Bluebird



Intellia / Novartis Editas CRISPR / Bayer

Slide courtesy of J. Keith Joung, Mass General / Harvard

#### Genome editing

nuclease > break > edited allele

- 1. indel KO
- 2. SNP correction of mutation
- 3. transgene targeted integration

### Basic science origins of editing

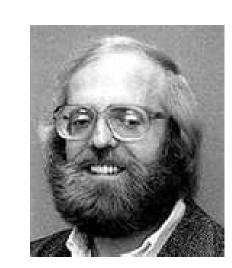
Maria Jasin - I-Scel



Aaron Klug and Carl Pabo – zinc finger proteins

Dana Carroll – zinc finger nucleases





Ulla Bonas – TAL effector

Jennifer Doudna and Emmanuelle Charpentier – RNA-guided nature

of Cas9



### Genome editing: clinical record (Oct 2016)

#### Target -> IND -> clinic:

### Case study 1: CCR5 / HIV / T cells

Target validation: human genetics; Berlin patient

Nuclease: ZFN (Sangamo)

Preclinical efficacy:

No adverse effect on T cell in vitro post-editing

Mouse model efficacy

Preclinical safety:

Patient cell dose NSG mouse tumorigenicity study

CMC:

GMP ex vivo Ad5/35 or ex vivo mRNA electroporation

#### Clinical record (2009 - )

80 subjects treated

Treatment well-tolerated to date

Edited cells persist >4 yrs

Stable increase in CD4 T cell count ~150 cells / ul

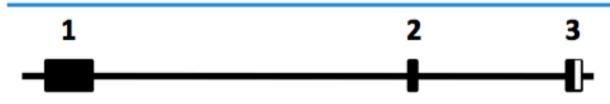
Decrease in viral reservoir

T cells home to the GALT and retain function

>60% subjects in recent cohort control virus in absence of ART

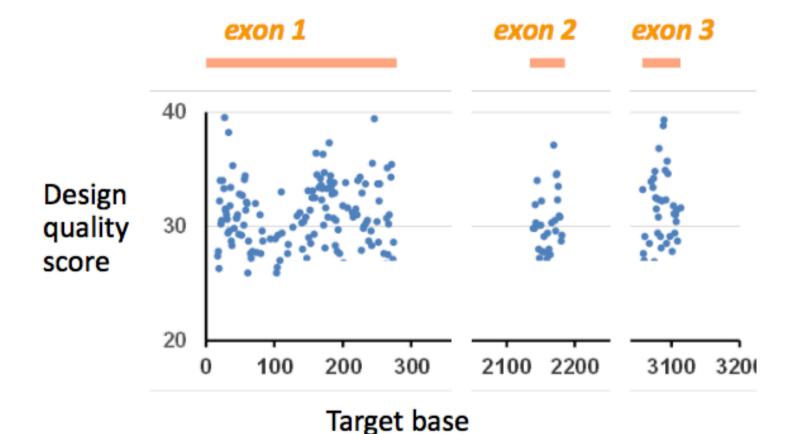
### Ninety one percent knockout in primary T cells

#### ZFNs Targeting TCRα Gene Are Highly Active and Well-Tolerated in Transformed and Primary Human Cells



**Example: TCRα constant region** 

- 3 exons / 374 bp
- 374 potential locations for cleavage
- → 218 locations are targetable with ZFNs (58%)
- → One ZFN pair per every 1.7 bp



T cells (10 days post transfection)

#### K562 cells

ZFN	Modification level (% indels)	
pair#	250 ng RNA	500 ng RNA
1	<b>77</b> ± 1.3	88 ± 1
2	<b>91</b> ± 1.5	<b>94</b> ± 1.7
3	<b>82</b> ± 0.6	<b>92</b> ± 1.3
4	95 ± 1.8	<b>96</b> ± 1.2
5	<b>88</b> ± 1.3	93 ± 1
6	<b>85</b> ± 1.4	89 $\pm 0.4$
7	<b>77</b> ± 1.7	87 ± 1.1
8	<b>54</b> ± 3.2	<b>75</b> ± 1.3
9	90 ± 1.6	<b>90</b> ± 1.5
10	<b>77</b> ± 0.8	<b>84</b> ± 2.2
GFP	0	0



ZFN	% TCR negative
pair#	cells
1	78.4%
2	88.1%
3	85.3%
4	85.2%
5	56.1%
6	91.4%
7	69.4%
8	81.7%
9	90.9%
10	85.6%
mock	1.8%

8/23/2016 7
Sangamo Confidential



# Genome-edited T cells in the clinic: general lessons

Targeted knockout ex vivo – single and multiple loci – clinical scale efficiency and specificity.

CMC GMP path established

Preclinical pharm-tox path established

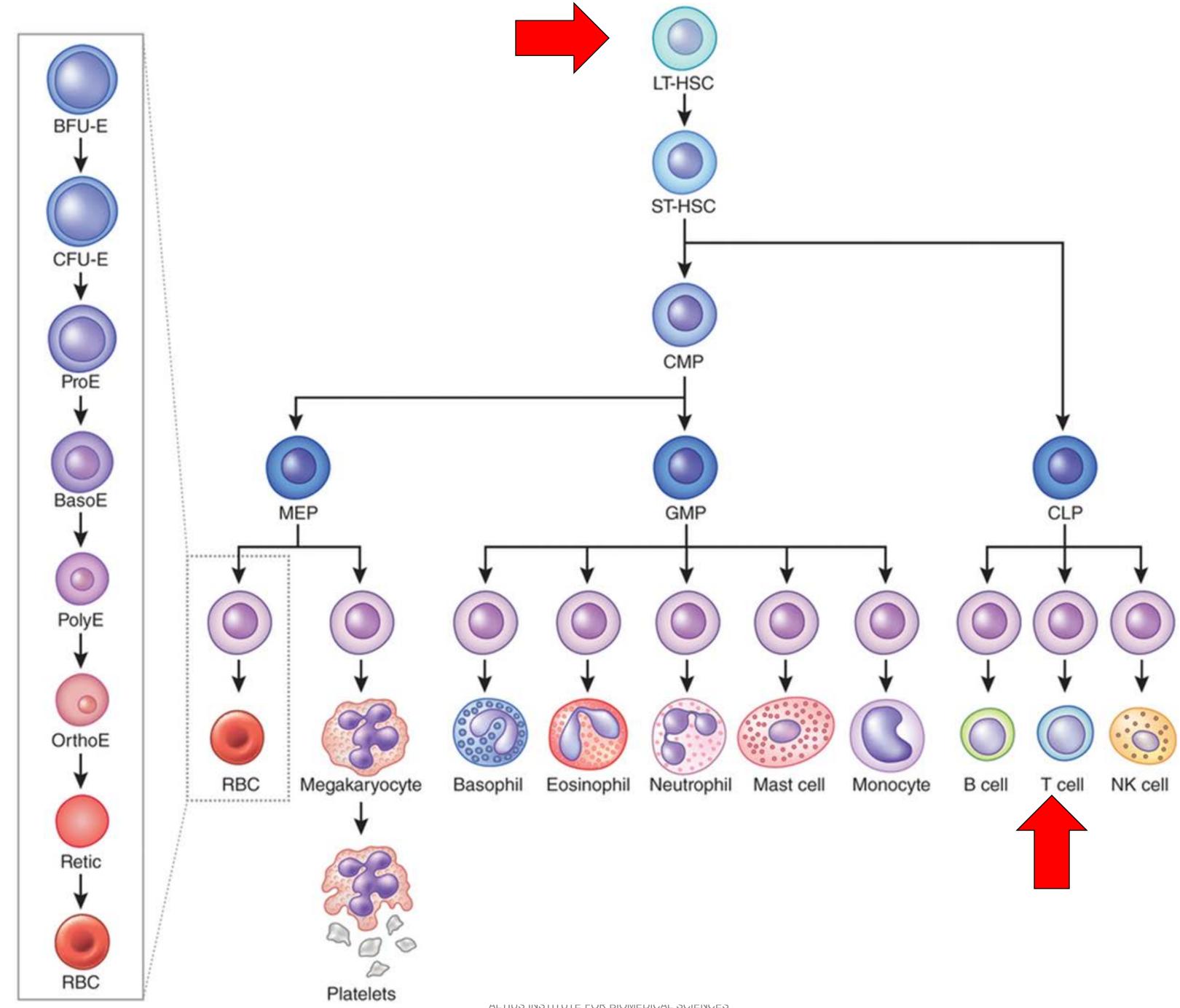
Regulatory framework (OCGCT CBER FDA) in place

Avenues for improvement

- 1. cell processing
- 2. allogeneic "off the shelf" rather than autologous product

# Efficient targeted addition in primary human T cells using mixed mode delivery

Wang and Holmes, 2015 Α PGK - AAVS1-R **GFP** B □Indel % Genome modification 8 0 0 0 ■GFP+ AAVS1-GFP AAVS1-GFP + ZFN 000 4000 6000 8000 SSC 0.52 53.91 4000 AAVS1 ZFN: 104 AAVS1-GFP: o (vg/cell) **GFP** 



## Target -> IND -> clinic: Case study 2: CCR5 / HIV / HSPCs

Target validation: human genetics; Berlin patient

Nuclease: ZFN (Sangamo)

Preclinical efficacy:

No adverse effect on CD34 cell in vitro post-editing Mouse model efficacy

Preclinical safety:

Patient cell dose NSG mouse tumorigenicity study

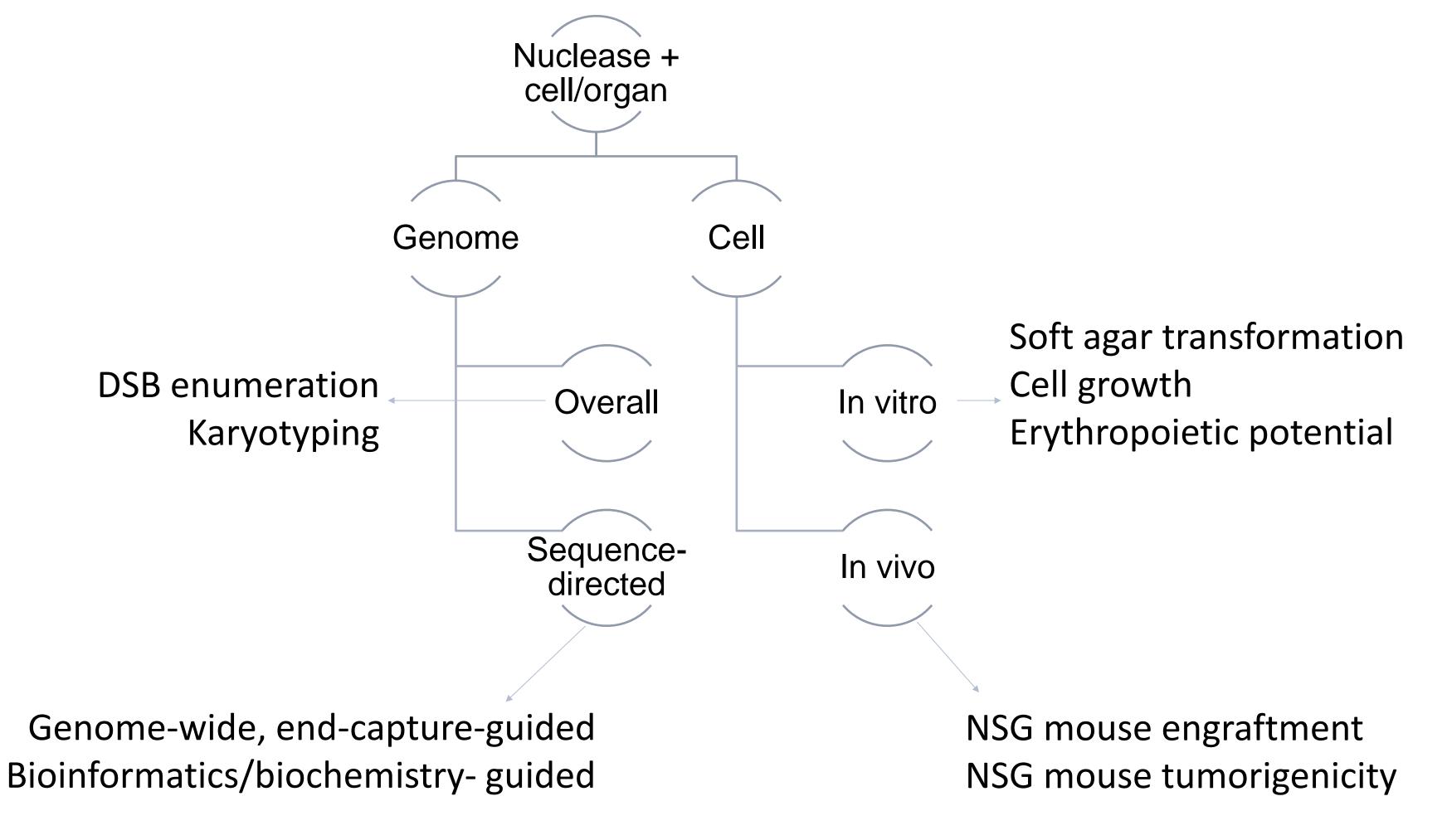
CMC:

mRNA electroporation

Clinical record

Clinical trial open

# Readouts of Genome Editing Safety: Charted Path Based on Two Open INDs For ZFN-Driven HSPC Editing



# Genome-edited CD34+ cells in the clinic: general lessons

Targeted knockout ex vivo – single loci – clinical scale efficiency and specificity.

CMC GMP path established

Preclinical pharm-tox path established

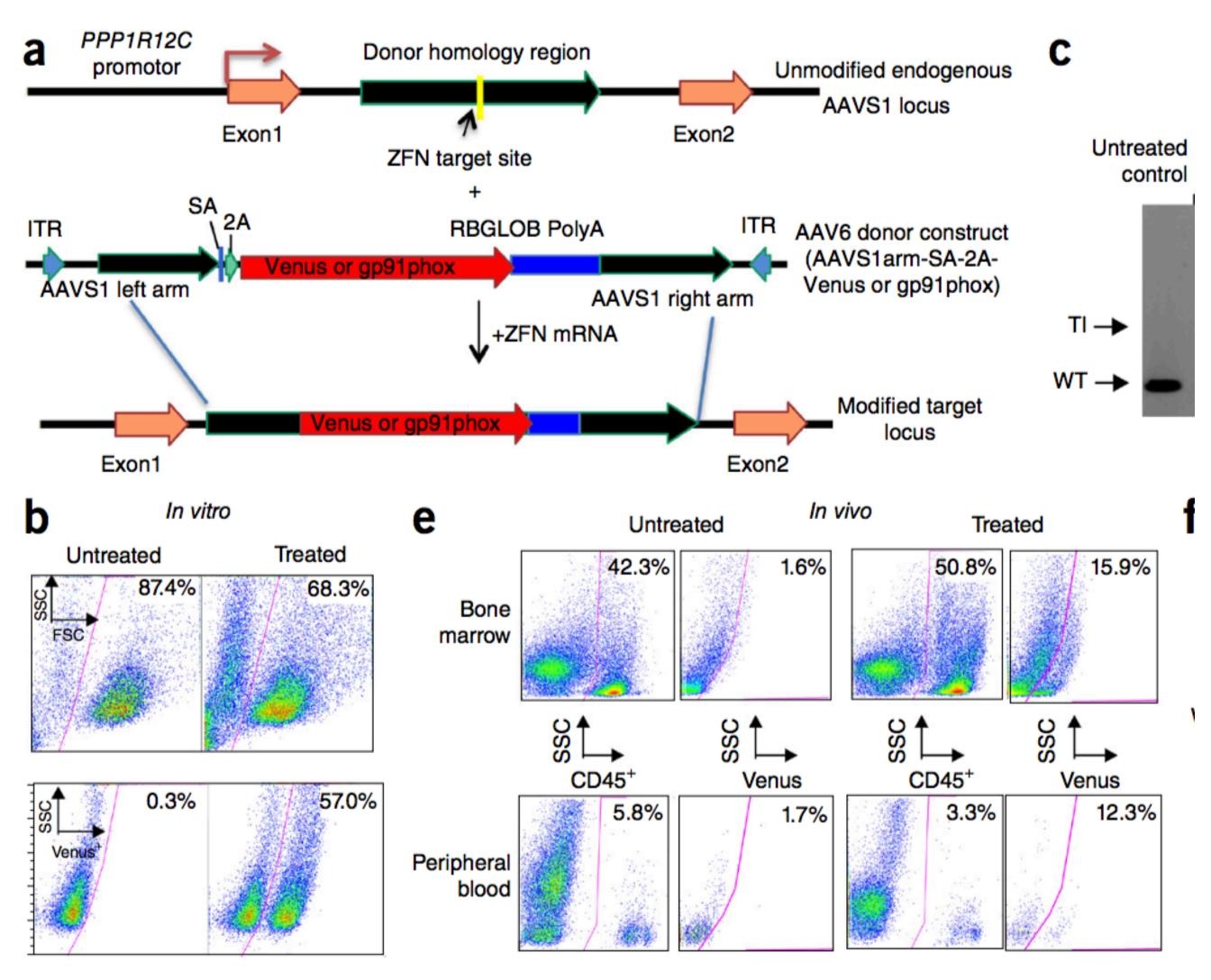
Regulatory framework (OCGCT CBER FDA) in place

Avenues for improvement

- 1. cell harvest ("sufficient numbers")
- 2. targeted correction/addition

# High efficiency targeted integration in human HSPCs using mixed-mode delivery

DeRavin NBT 2016



#### Near- and longer-term challenges

T cells (oncology): cell weaponized beyond a CAR/TCR

T cells (oncology): allogeneic (off the shelf) product

T cells (beyond oncology and HIV): editing for improved potency – targets!

HSPCs: harvest and expansion (the iPSC route?)

HSPCs: higher efficiency of targeted integration

HSPCs: rare diseases of the blood

### Next-generation challenge

In vivo delivery (note three open INDs for in vivo editing in the liver – F9 deficiency, MPS I, MPDs II, and multiple ongoing clinical trials by Alnylam and Ionis on RNAi and antisense).

