GeneCopoeia CRISPR sgRNA Libraries For Functional Genomics

GeneCopoeia, Inc.

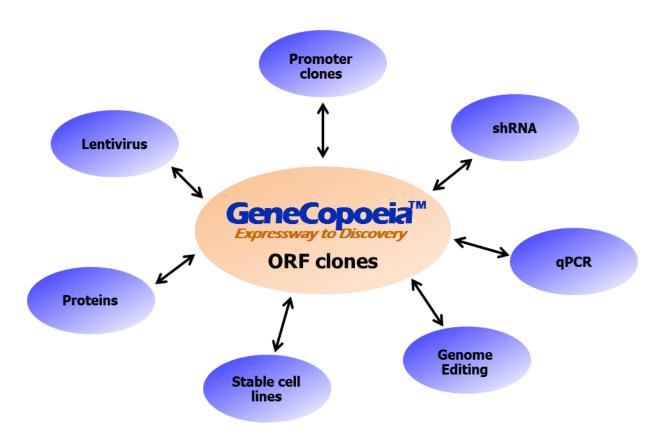
Presenter:

Ed Davis, Ph.D.
Senior Application Scientist
GeneCopoeia, Inc.

April 29, 2015

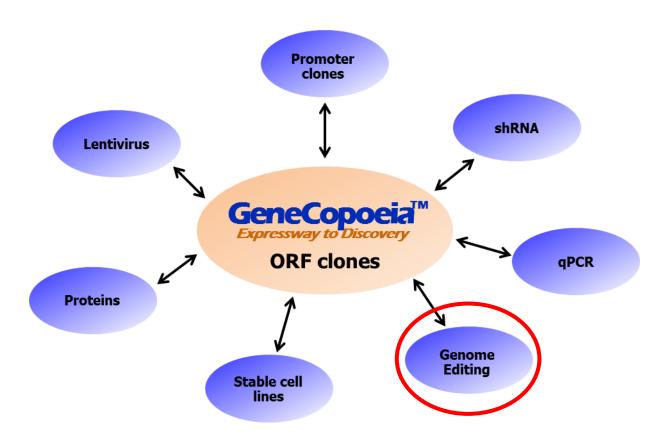


GeneCopoeia products & services



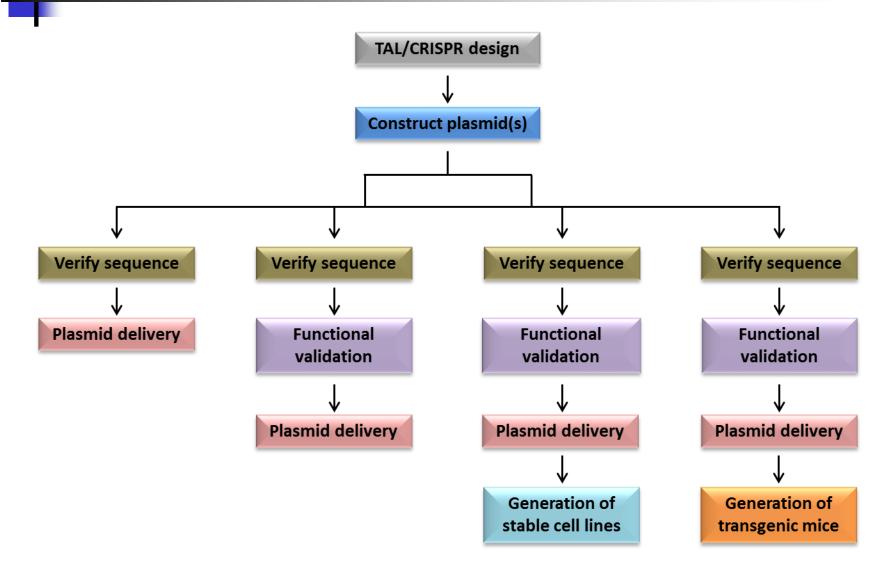


GeneCopoeia products & services





GeneCopoeia genome editing services





Outline

- Introduction: Functional genomics & CRISPR
- What are CRISPR sgRNA libraries?
- Development of previous libraries
- GeneCopoeia pathway- & gene groupspecific CRISPR sgRNA libraries



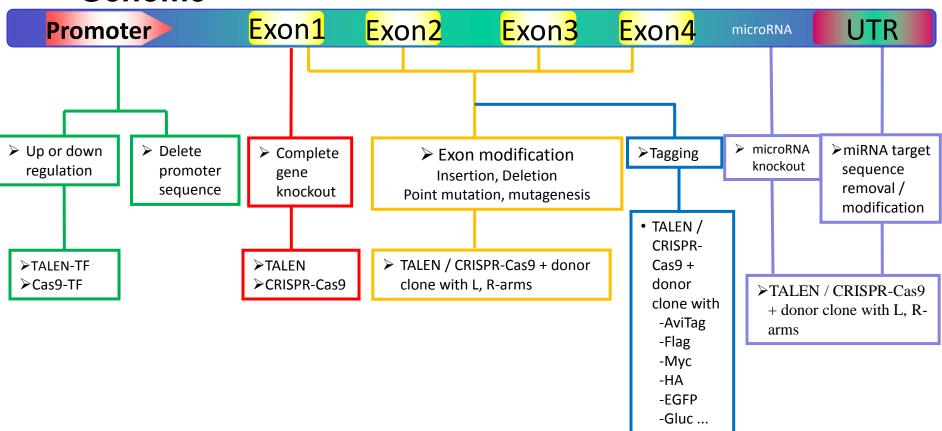
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Functional genomics

Genome



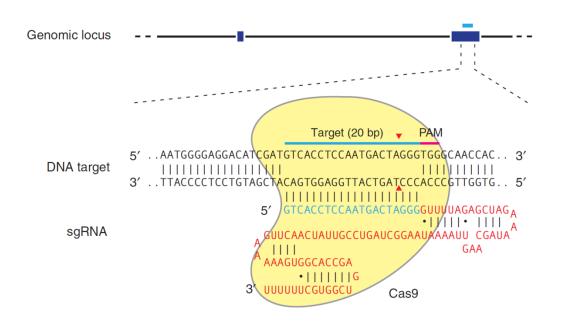
- 1) Safe harbor knock-in ORF clones are used for transgene insertion at the human AAVS1 or mouse ROSA26 genomic sites using CRISPR/Cas-9 or TALEN. Safe harbor site integration ensures that transgenes will be transcriptionally active and expressed at consistent levels, and presents no known adverse effects on cells caused by disruption of the sites.
- 2) IndelChek™ kit for TALEN and CRISPR/Cas-9 functional validation.





CRISPR genome editing technology

CRISPR-Cas9: RNA-guided endonuclease

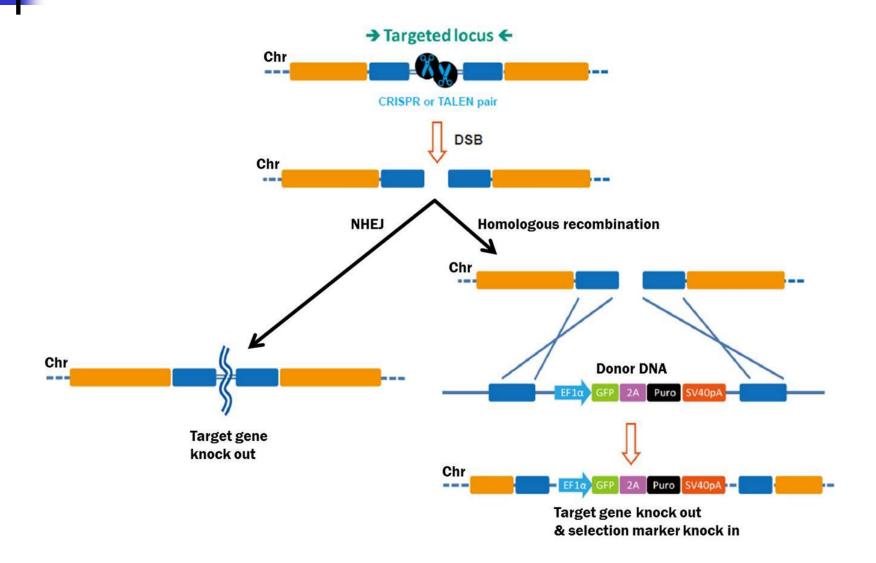


Ran, et al. (2013). Nature Protocols 8, 2281

- 20 nt single guide RNA (sgRNA) guides Cas9 nuclease to target site.
- Requires NGG "PAM" site immediately downstream of sgRNA target sequence.
- Cas9-RNA complex makes DSB 3-4 nt upstream of PAM.
- Target almost any gene in any cell



Targeted DNA editing by DSB induction





Applications for genome editing

Knockout via NHEJ

indels in human *EMX1* locus

```
WT 5'-..GGAGGAAGGGCCTGAGTCCGAGCAGAAG-AAGAAGGGCTC..-3'
D1 GGAGGAAGGGCCTGAGTCCGAGCAGAAG-AAGAAGGGCTC
+1 GGAGGAAGGGCCTGAGTCCGAGCAGAAGAAAGAAGGGCTC
D2 GGAGGAAGGGCCTGAGTCCGAGCAGAAG---GAAGGGCTC
D3 GGAGGAAGGGCCTGAGTCCGAGCAGAAG----AAGGGCTC
GGAGGAAGGGCCTGAGTCCGAGCAGAAG-----AGGGCTC
```

Cong, et al. (2013). Science 339, 819



Pathway & gene group sgRNA libraries

Library name	Number of genes	
Innate kinases & ubiquitin ligases	239	
Nuclear hormone receptors	118	
Tumor metastasis genes	57	
Oncogenes	288	
Tumor suppressor genes	231	
Protein kinases	658	
Key genes in 50 pathways	139	
Custom	Made-to-order	

Available as bacteria, DNA, or lentiviral particles either:

- A. Pooled
- B. Individually arrayed





What are CRISPR sgRNA libraries?

- Collections of hundreds-to-thousands of plasmids encoding sgRNAs
- > In presence of Cas9, create DSBs intended to knock genes out
- Knock out many genes simultaneously
- Target whole genome or smaller groups of genes



Applications for CRISPR sgRNA libraries

Previously carried out using shRNA libraries

- Drug target discovery (e.g. Cooper & Brockdorff, 2013. Development 140, 4110)
- Reporter assays (e.g. Sethi, et al., 2012. PLoSONE 7.)
- Loss-of function phenotypic screening (*e.g.* Zhang, *et al.*, 2014. *Proc Natl Acad Sci U S A* **110**, 12361.
- > Others (Drug target validation, *etc.*)



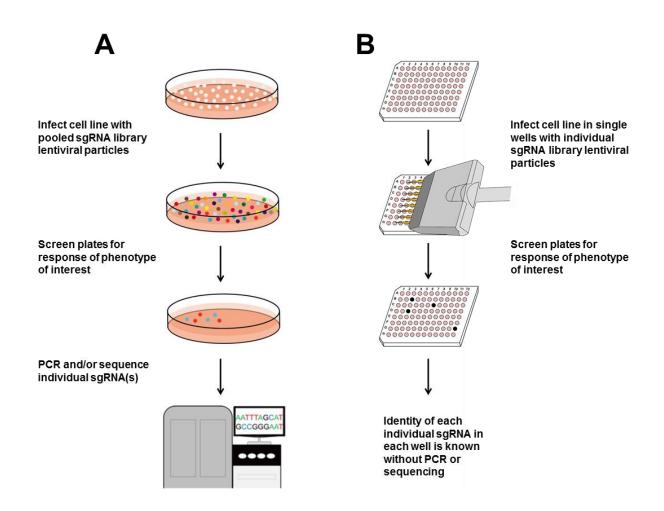
CRISPR or RNAi?

Knock out vs. Knock down

Method	Change genetic code	Change expression level	Knock down	Knock out
CRISPR	✓	✓		✓
RNAi		✓	•	



Important: Need "readout" (phenotype or assay)







Example readout # 1: Cell death

- Step 1: Infect cell line with pooled sgRNA lentiviral particles
- Step 2: After a few days of cell division, prepare DNA from mixed population of living cells and either:
 - Perform Sanger sequencing on selected sgRNAs

OR

- Perform deep sequencing, analyze representation of all sgRNAs before and after infection
- > sgRNAs that are under-represented in living cells represent candidates for genes necessary for biological process of interest.
- Step 3: Test individual candidate sgRNAs in assay to validate hits





Example readout # 2: Drug resistance

- Step 1: Infect cell line with pooled sgRNA lentiviral particles
- Step 2: Apply drug of interest to selected cells and allow cell division. Then either:
 - Perform Sanger sequencing on selected sgRNAs

OR

- Perform deep sequencing, analyze representation of all sgRNAs before and after infection
- > sgRNAs that are over-represented in pool represent candidates for genes necessary for drug mechanism of action.
- Step 3: Test individual candidate sgRNAs in assay to validate hits





Example readout # 3: Reporter assay

- > Step 1: Infect cell line with pooled sgRNA lentiviral particles
- > Step 2: FACS sort cells to enrich for those expressing a fluorescent reporter (e.g. GFP). Then either:
 - Perform Sanger sequencing on selected sgRNAs

OR

- Perform deep sequencing, analyze representation of all sgRNAs before and after infection
- > sgRNAs that are under- or over-represented in pool represent candidates for genes necessary biological process of interest.
- Step 3: Test individual candidate sgRNAs in assay to validate hits



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Previously-developed libraries

Genetic Screens in Human Cells Using the CRISPR-Cas9 System

Tim Wang, 1,2,3,4 Jenny J. Wei, 1,2 David M. Sabatini, 1,2,3,4,5*† Eric S. Lander 1,3,6*†

Wang, et al. (2014). Science 343, 80

Genome-Scale CRISPR-Cas9 Knockout Screening in Human Cells

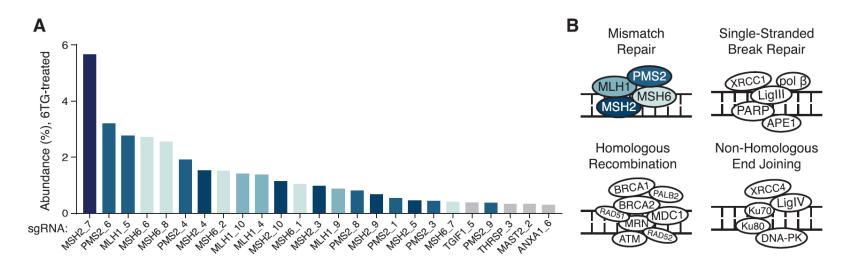
Ophir Shalem,^{1,2}* Neville E. Sanjana,^{1,2}* Ella Hartenian,¹ Xi Shi,^{1,3} David A. Scott,^{1,2} Tarjei S. Mikkelsen,¹ Dirk Heckl,⁴ Benjamin L. Ebert,⁴ David E. Root,¹ John G. Doench,¹ Feng Zhang^{1,2}†

Shalem, et al. (2014). Science 343, 84

Used pools of lentiviral-delivered sgRNAs for genome-scale forward mutagenesis screens in human cells



Previously-developed libraries



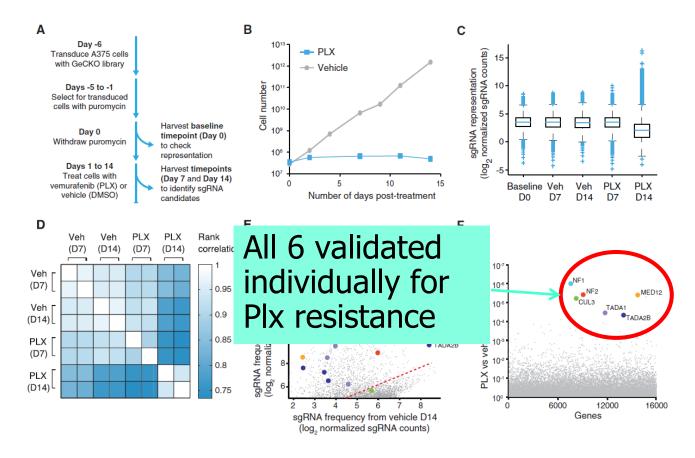
Wang, et al. (2014). Science 343, 80

Pool of sgRNAs targeting 7,300 genes



Previously-developed libraries

Plx drug resistance screen in A375 melanoma cells



Shalem, et al. (2014). Science 343, 84



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Available as bacteria, DNA, or lentiviral particles either:

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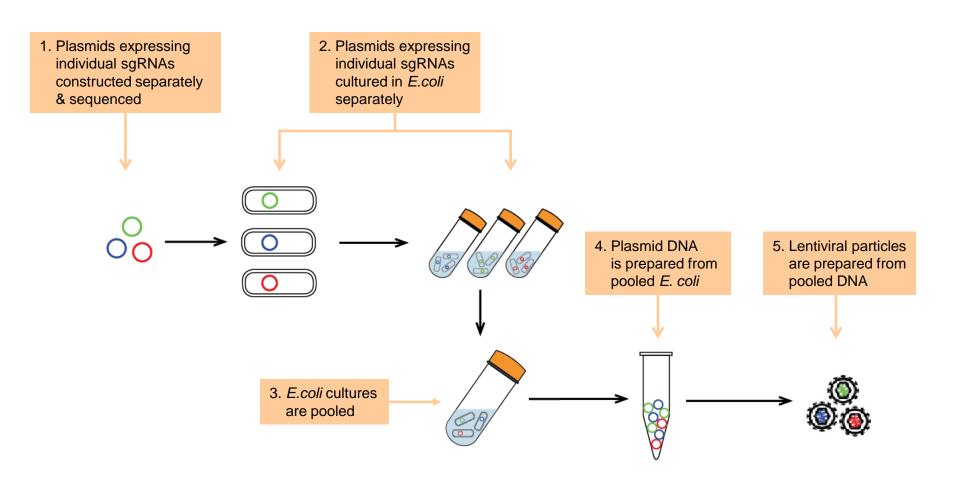


Advantages of the GeneCopoeia sgRNA libraries

- Individually constructed and cultured in *E. coli* before pooling. Makes possible use in pools or as individual sgRNAs.
- Pools limited to 150 sgRNAs, ensuring excellent representation of each sgRNA
- Sequence verification provides high quality of each sgRNA
- Small library sizes: Reduces time and cost of screening

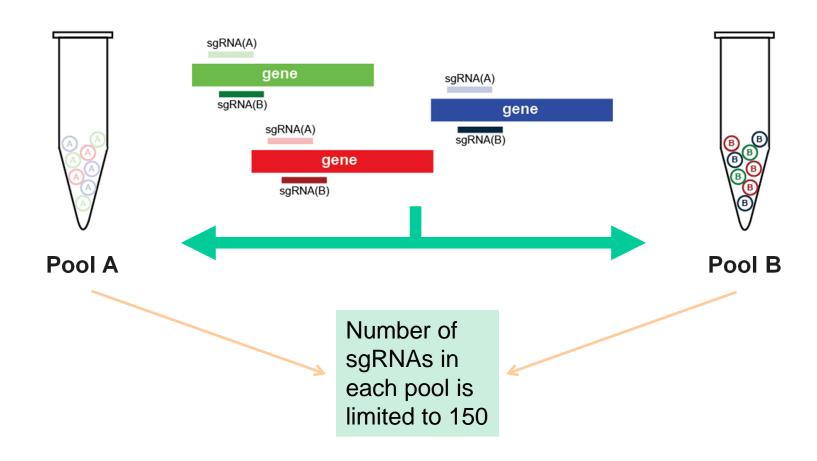


Library pooling strategy



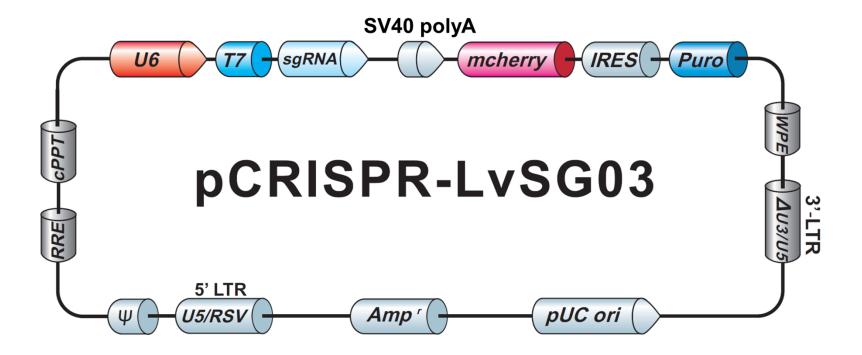


Library pooling strategy





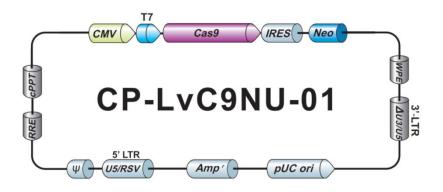
"Dual-use" lentiviral vectors

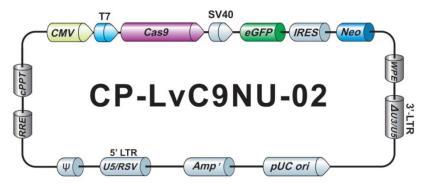




sgRNA library considerations

Best to use Cas9-expressing stable cell line for transducing with sgRNA libraries





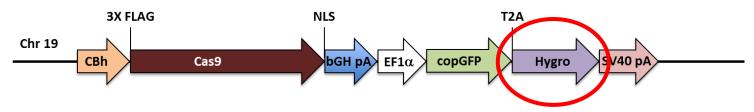




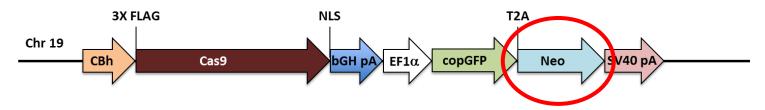
sgRNA library considerations

Best to use Cas9-expressing stable cell line for transducing with sgRNA libraries

Cas9 integrated at AAVS1 with Hygro:



Cas9 integrated at AAVS1 with Neo:



Can purchase pre-made Cas9 cell lines, order custom Cas9 cell line, or purchase DIY Safe Harbor clones

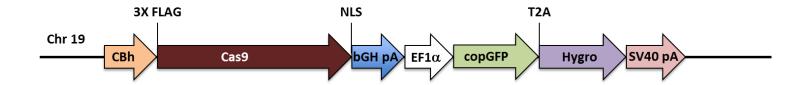




How do I order sgRNA libraries from GeneCopoeia?



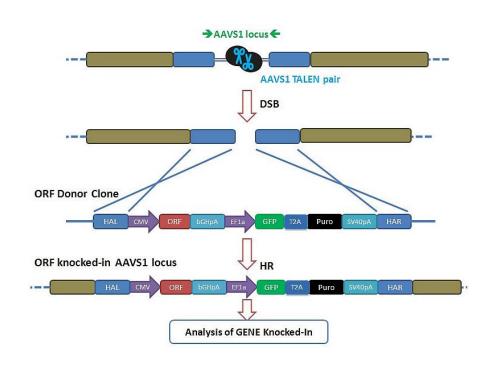
Premade Cas9-expressing stable cell lines



- > Stable cell lines with constitutively-inducible-expressing Cas9
- > Have pre-made lines, or can have us integrate Cas9 in your cell line
- Donor clones available for DIY stable cell line creation
- Cas9 integrated at Safe harbor locus for high expression and insertion without consequences
- Ideal for sgRNA library screening or validation



Safe Harbor cloning kits

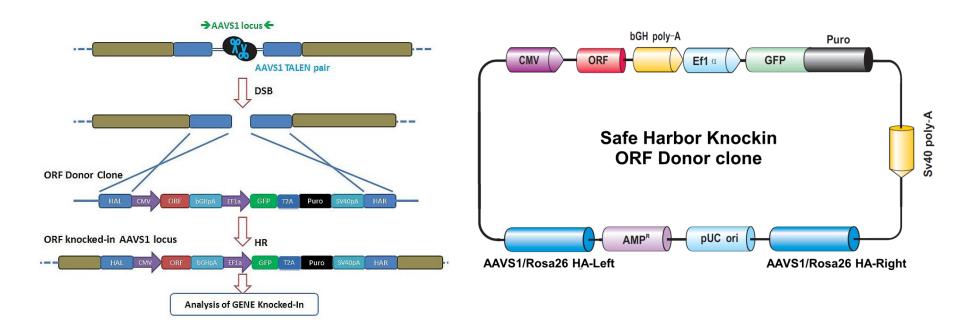


Features

- Human AAVS1 & mouse Rosa26 sites ensure transcriptioncompetency of the transgenes & present no known adverse effects on cells
- Safe Harbor integration provides low copy number of transgene & close to physiological-level expression.
- Can use with TALEN or CRISPR



Safe Harbor knockin ORF clones

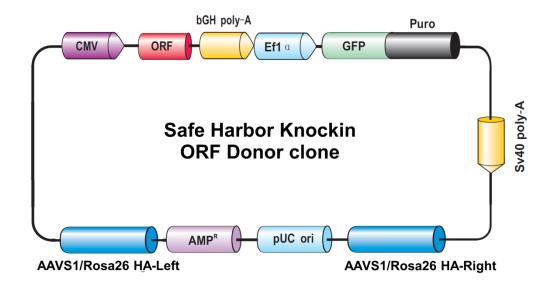


Features

- Over 40,000 sequence-verified human & mouse ORFs available
- Inserted between AAVS1 or Rosa26 sites for ready safe harbor integration using TALEN or CRISPR



Safe Harbor knockin ORF clones



Some applications for Safe Harbor knockin ORF clones:

- Rescue of a knockout or mutagenesis phenotype
- Overexpression of a fusion tagged protein
- Expression of a gene from a different species (e.g. human gene in a mouse)



Lentiviral production service & packaging kits

- > Packaging of lentiviral-based clones into ready-to-use particles
- > Ideal for CRISPR lentiviral clones
- Can package individual CRISPR clones, or CCI/outside libraries



GeneCopoeia Technical Note: sgRNA libraries



TECHNICAL NOTE

Genome Editing: Applications For GeneCopoeia CRISPR sgRNA Libraries

Ed Davis, Ph.D.

Biomedical researchers are enjoying a Renaissance in functional genomics, which aims to use a wealth of DNA sequence information—most notably, the complete sequence of the human genome—to determine the natural roles of the genes encoded by the genome. As a result, biochemical networks and pathways will be better understood, with the hope of leading to improved disease treatments.

A major approach of functional genomics is to ablate gene function, by either "knockdown" (reduction) or "knockout" (complete elimination). Since 2012, researchers have turned increasingly to CRISPR (clustered, regularly interspaced, short palindromic repeats) for functional genomics studies. CRISPR's simple RNA-guided mechanism provides a quick, convenient, and relatively low-cost method for many applications, from gene knockout, in-frame fusion tagging, mutagenesis, and transgene knockin. Several groups recently adapted CRISPR for high-throughput knockout applications, by developing large-scale CRISPR sgRNA libraries. GeneCopoeia recently launched a number of smaller, pathway- and gene group-focused CRISPR sgRNA libraries, which offer several key advantages over the whole-genome libraries. In

Download from:

http://www.genecopoeia.com/wp-content/uploads/2015/04/GeneCopoeia-Technical-Note-CRISPR-sgRNA-Libraries-05-2015.pdf



Summary

- CRISPR sgRNA libraries have tremendous potential for functional genomics screening
- Applications for CRISPR sgRNA libraries include novel drug target discovery, drug target validation, phenotypic screening, reporter assays, & more
- GeneCopoeia sgRNA libraries advantages over previously developed libraries include sequence validation, individual clone construction, and individual culturing in *E. coli*
- GeneCopoeia CRISPR sgRNA libraries are available in a range of formats, and have many powerful companion products and services





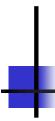
GeneCopoeia CRISPR & TALEN Technology For Genome Modification

Wednesday, May 13, 2015 9:00 am EDT (GMT-0:400)

Register here:

https://attendee.gotowebinar.com/register/4429962 400977907202





Another upcoming webinar!

GeneCopoeia Genome Editing Stable Cell Line Services

Wednesday, May 27, 2015 12:00 pm EDT (GMT-0:400)

Register here:

https://attendee.gotowebinar.com/register/7724986 543678467842





Any questions?

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Or visit us on the web:

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