

# 大阪大学共同研・ゲノム編集センター 使用説明会 ～明日から始めるゲノム編集～

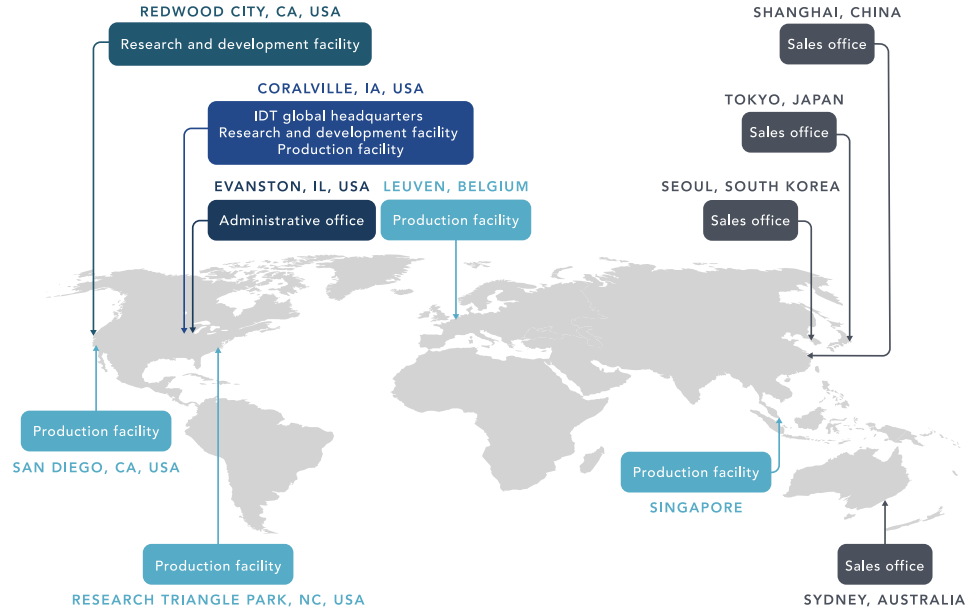
インテグレートッドDNAテクノロジーズ株式会社

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04-21-2026

# TODAY, IDT IS THE LEADING BRAND IN DNA AND RNA SYNTHESIS

- Founded in 1987 by Dr. Joseph Walder
- Largest custom oligonucleotide manufacturer worldwide
- >1500 employees in 9 locations
- >130,000 active customers
- >95% of ordered products are manufactured and shipped in less than 24 hours

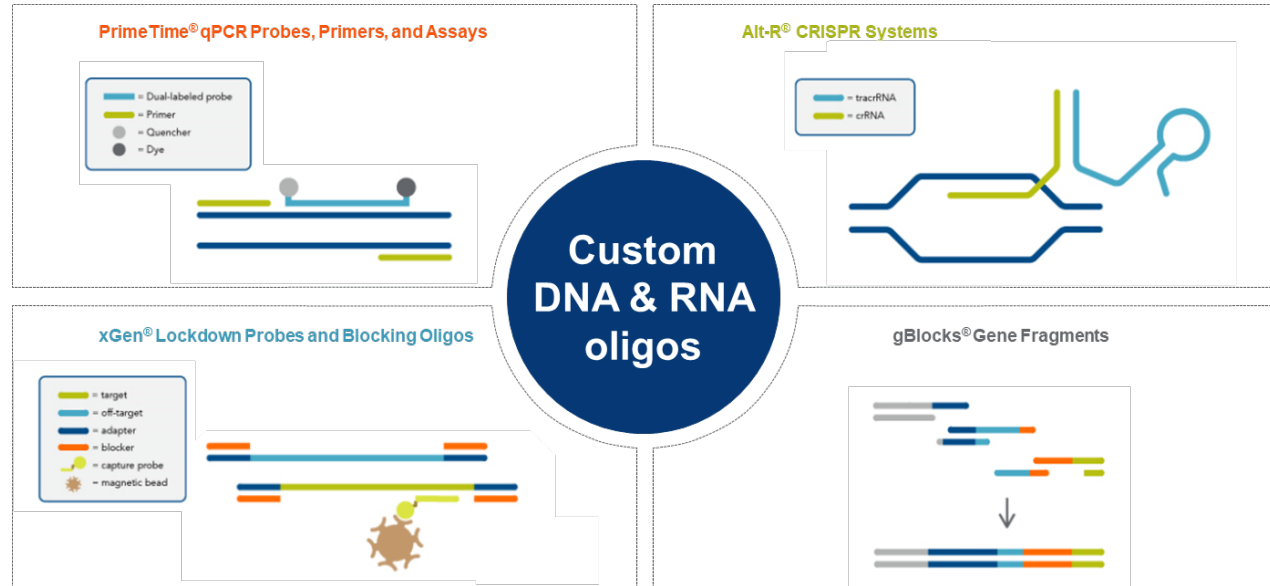


# THE WORLD'S LARGEST SUPPLIER OF CUSTOM NUCLEIC ACIDS

>64,000 oligos synthesized every day

- Major R&D teams (~80 employees)
- CRISPR
- NGS
- Genotyping
- qPCR
- Synthetic Biology
- RNAi
- Bioinformatics

## Major Product Lines



# ゲノム編集技術について

これまでと比べて、画期的である理由

- 汎用性: 動物、植物、昆虫、微生物、、、各種の生物に適用可能
- 高効率: 改変効率が極めて高く、改変個体や細胞の同定、単離が容易
- 多用途: どのゲノム領域でもターゲットにできる、いろいろな改変可能
- 特異的: ターゲットのみを改変し、ほかに痕跡を残さない
- 簡便: 試薬・機器を比較的簡単に安価に入手し短期間で行える

既存の技術では、特定の生物でのみ、低い効率で、長い時間と大きな費用をかけてしか実現できなかった。

ゲノム編集がその制約を大きく乗り越えた。

(医療応用実用などには倫理面を含め、越えなければならないハードル未だ多し)

# Alt-R™ CRISPR SYSTEM: A COMPLETE WORKFLOW

ターゲットのデザイン → 試薬の選定 → → → → → → → → → → 解析

Design

CRISPR-Cas  
reagents

HDR  
reagents

Analyze

## Alt-R Cas9 gRNA design tool

- Predesigned guides
- Custom designs
- Design checking

## Alt-R Cas9 HDR design tool

- Friendly UI
- Empirically defined design rules
- Integration with Cas9 gRNA designs

## rhAmpSeq design tool

### Alt-R gRNAs

- Cas9 crRNA:tracrRNA
- Fluorescently labeled tracrRNAs
- Cas9 sgRNA
- Cas12a crRNA
- Custom ordering for any gRNA (e.g. pegRNA, Cas13)

### Alt-R CRISPR proteins

- WT Cas9
- HiFi Cas9
- Cas9 nickases
- A.s. Cas12a *Ultra*
- L.b. Cas12a *Ultra*
- Fluorescently labeled Cas9

### Alt-R Electroporation Enhancers

### Alt-R HDR Donor Oligos

- Up to 200 nt
- Modified ssODNs

### Alt-R HDR Donor Blocks

- Modified to reduce blunt integration
- Up to 3000 nt
- Sequence-verified via NGS

### Alt-R HDR Enhancer

- V2

### Alt-R HDR Enhancer

- Protein

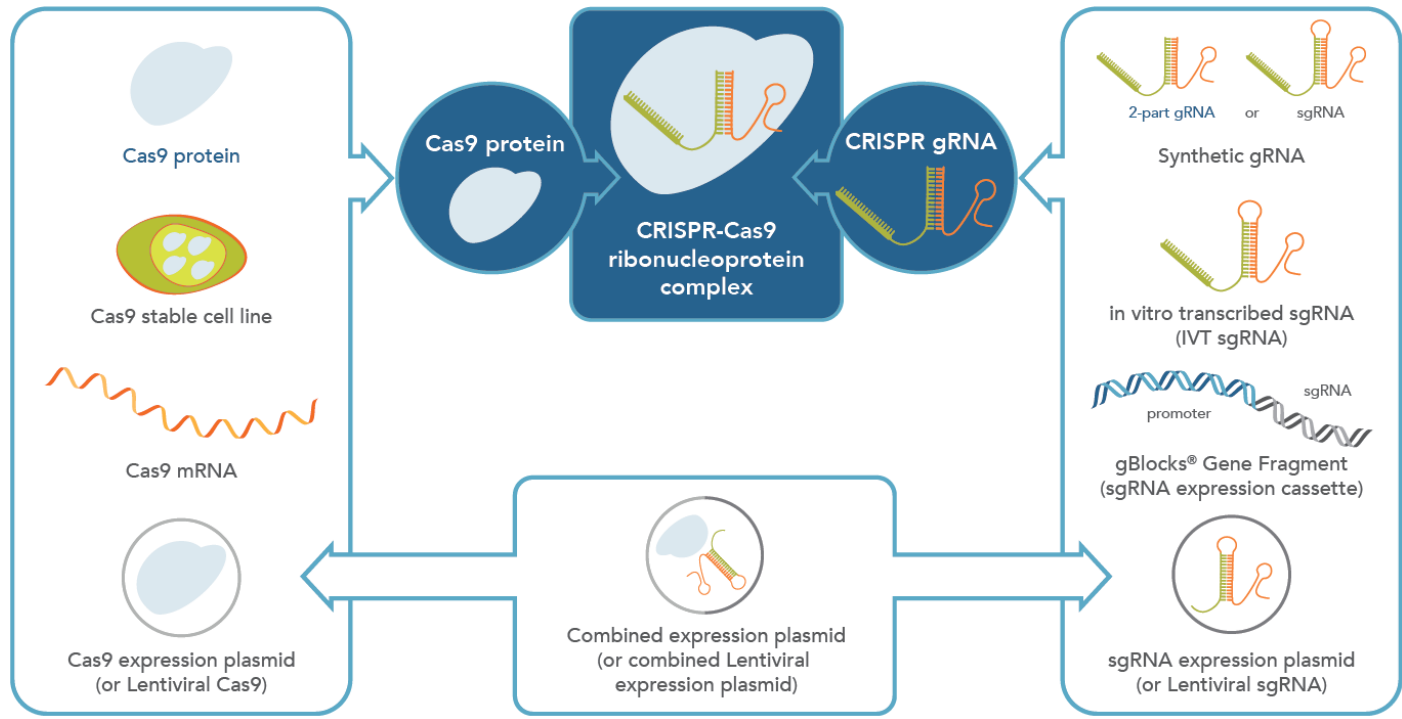
## rhAmpSeq system for CRISPR

- Multiplexed amplicon sequencing

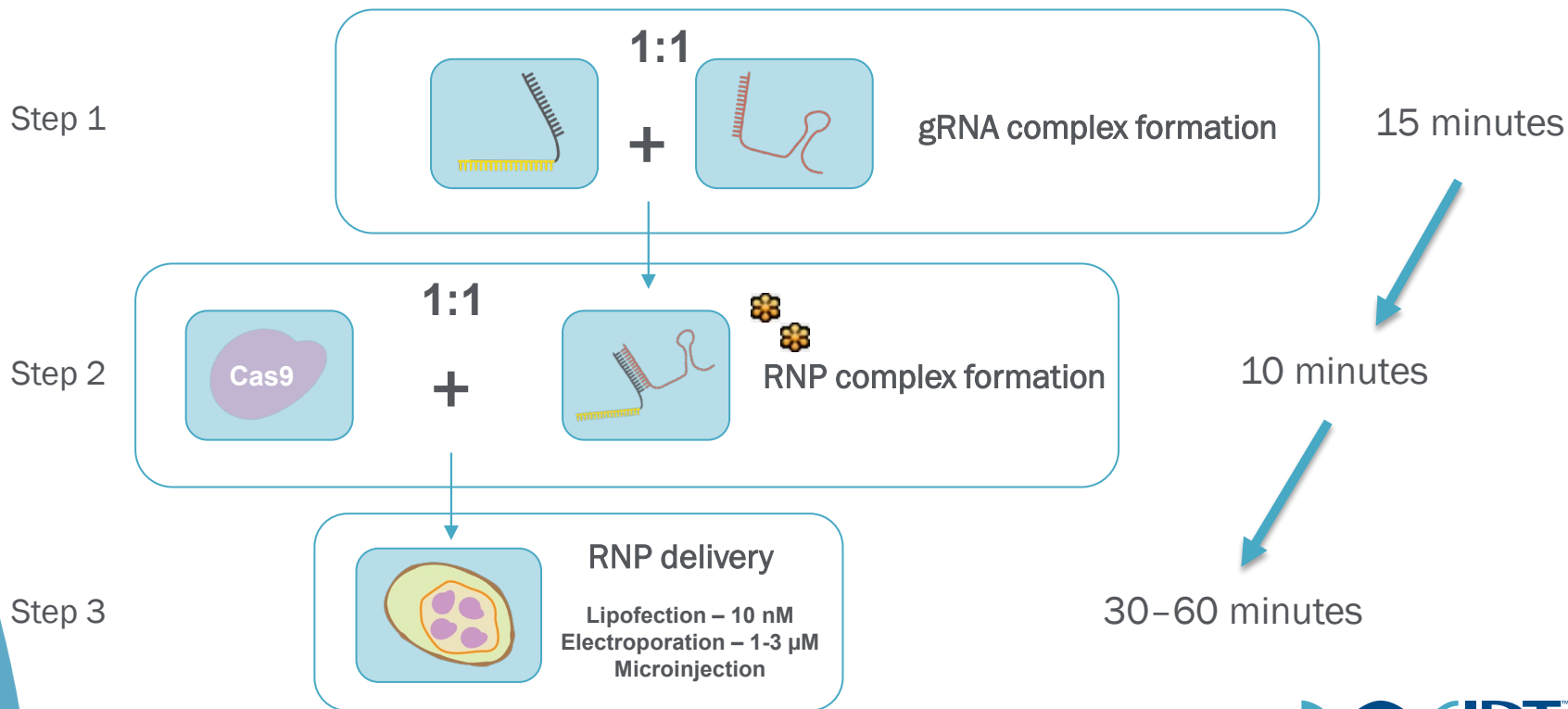
## CRISPRAltRations NGS analysis tool

- Cloud-hosted UI for analysis of CRISPR on- and off-target editing

# ゲノム編集に必要な 構成試薬 (GRNAとCAS9タンパク質)

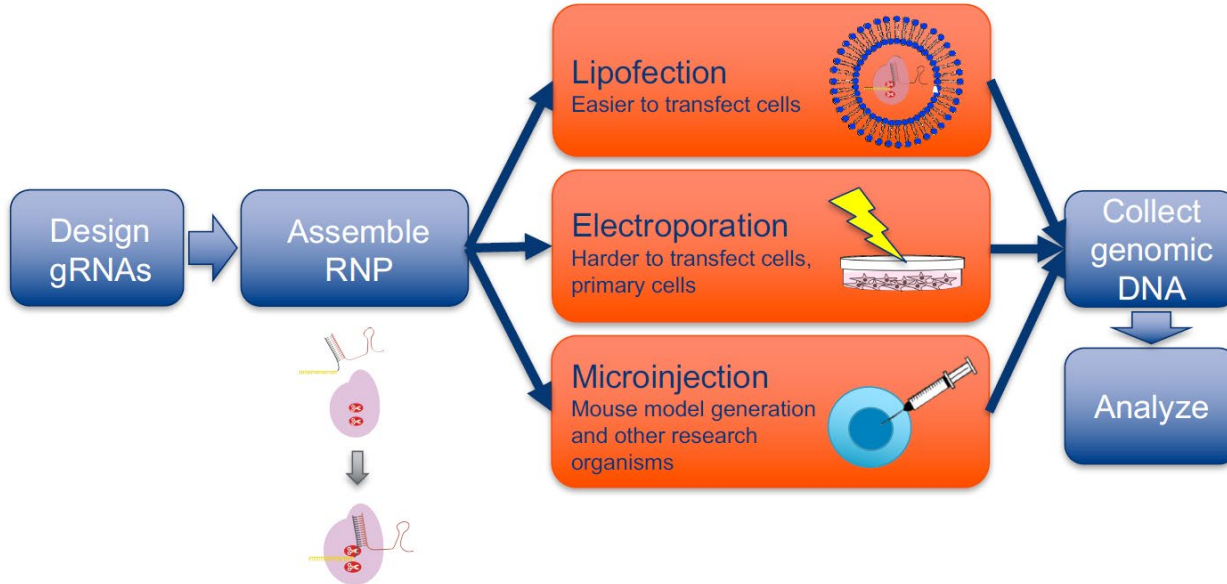


# 簡便なプロトコール

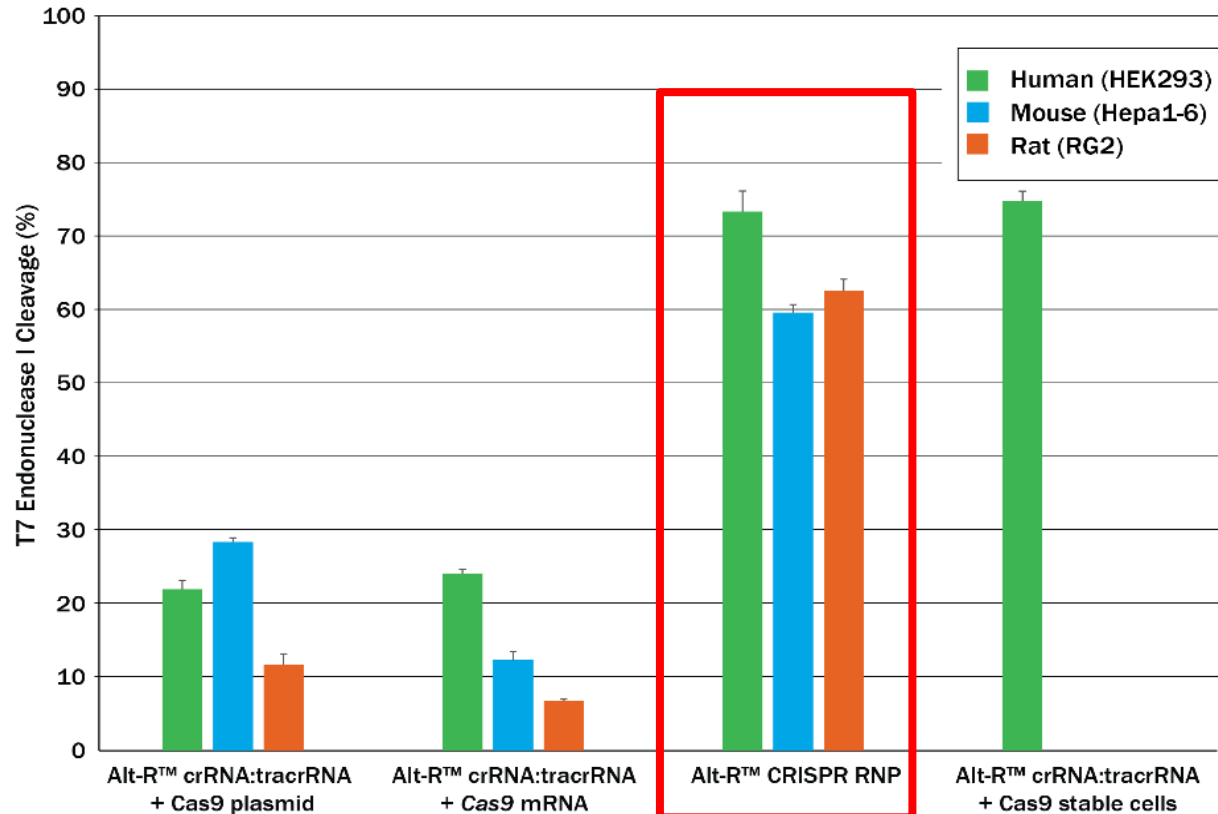


# CAS9 RIBONUCLEOPROTEIN(RNP)導入法

## CRISPR workflow







# リコンビナントCAS9タンパク質利用による編集効率UP



Confidential - Company Proprietary

# Alt-R gRNA OPTIONS FOR *S.p.* Cas9 & *A.s.* Cas12a

Guide RNAs	Cas9 guide RNAs			Cas12a guide RNAs	
<b>Structure</b>	Alt-R 2-part 	Alt-R 2-part XT 	Alt-R sgRNA 	Alt-R Cas12a crRNA 	
<b>gRNA format</b>	Alt-R CRISPR-Cas9 crRNA & tracrRNA	Alt-R CRISPR-Cas9 crRNA XT & tracrRNA	Alt-R CRISPR-Cas9 sgRNA	Alt-R CRISPR-Cas12a crRNA	
<b>Components</b>	crRNA tracrRNA	crRNA XT tracrRNA	sgRNA	crRNA	
<b>Size (nt)</b>	36 67	36 67	100	40-44 (41 nt recommended)	
<b>Annealing required</b>	Yes	Yes	No	No	
<b>Stability</b>	++	+++	++++	+++	
<b>Applications</b>	<ul style="list-style-type: none"> <li>• Cas9-expressing cells</li> <li>• RNP in most cell types</li> </ul>	<ul style="list-style-type: none"> <li>• Co-delivery with Cas9 plasmid/Cas9 mRNA</li> <li>• RNP under difficult experimental conditions (e.g., high nuclease environments)</li> </ul>		<ul style="list-style-type: none"> <li>• KO/KI, RNP in most cell types</li> <li>• Cas12a-expressing cells</li> </ul>	

# IDT CRISPR-CAS9 GRNA DESIGN TOOL

- 3種類のツールがあります。
- **1) Search for Predesigned CRISPR: プレデザインを検索**
  - ヒト、マウス、ラット、ゼブラフィッシュ、線虫の遺伝子を対象
- **2) Design Custom crRNAs: カスタムで設計**
  - お客様が指定するターゲット配列情報から設計
  - 全生物種の遺伝子を対象
- **3) CRISPR-Cas9 crRNA Checker: 既存のターゲット配列の検証**
  - 論文などに記載されているproto spacer 配列のデザインを検証する

# Alt-R Cas9 CRISPR LIBRARIES

## Alt-R™ CUSTOM CRISPR gRNA LIBRARIES

Chemically modified guide RNA libraries for CRISPR screening research



Fast turnaround time, flexible formulation



Adaptable for different CRISPR systems



Complete CRISPR workflow solution from custom design to analysis

Alt-R Custom CRISPR gRNA libraries are available for all CRISPR nucleases, including Cas9, Cas12a, Cas13, prime editing enzymes, and others. These libraries were developed to address the need for better CRISPR screening solutions. They are chemically modified guide RNAs (gRNAs) synthesized on the IDT proprietary high-fidelity RNA manufacturing platform to provide high quality, reliable gRNA libraries with fast delivery.

### BENEFITS

1. Innovative solution offered by a global leader in RNA synthesis and CRISPR innovation
2. Reliable, consistent, and fast delivery, with custom formulations available to suit a variety of project needs
3. Adaptable for alternative CRISPR systems such as Cas12a, Cas13, and prime editing
4. Enhanced nuclease resistance for maximal editing in Cas9-expressing cells or via ribonucleoproteins (RNP)
5. Optimized RNA synthesis processes to mitigate cross-contamination risk

### PRODUCT SPECIFICATIONS

Features	Options
Design	Pre-designed, custom, user-provided
CRISPR systems	Cas9, Cas12a, Cas13, prime editing, and other alternative systems
Guaranteed yield	0.5 nmol, 2 nmol, 5 nmol, and custom normalized deliverables
Cas9 gRNA formats	2-part crRNA:tracrRNA complex and sgRNA
Custom lengths supported	30-150 nt
Chemical modifications	2'-O-methyl RNA, PS linkages, end-blocking Alt-R modifications
Plate types	96- & 384-well PCR, Deep-well, V-bottom, ECHO, custom options available
Formulation options	Multi-guide per well, pooled by gene Arrayed (single gRNA/well) Custom formulations upon request
QC	Individual ESI/MS
Supporting reagents & functional analysis pipeline (optional)	WT Cas9, HiFi Cas9, Cas12a, and Cas12a Ultra Glycerol-free options available in tubes or plates (ideal for robotics) Electroporation Enhancers rhAmpSeq™ CRISPR Analysis System (NGS-based on-/off-target editing analysis)

> [WWW.IDTDNA.COM](http://WWW.IDTDNA.COM)

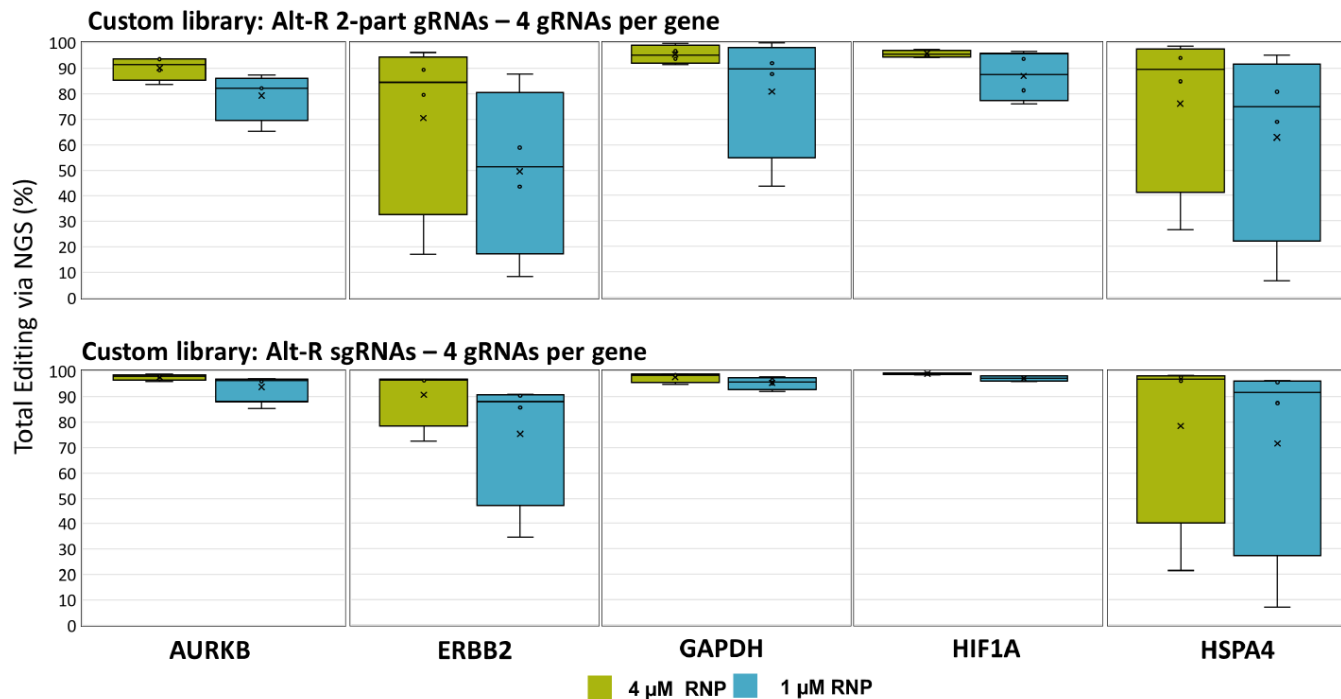


### Custom gRNA libraries:

- 2 part (crRNA+tracrRNA) and sgRNA
- Standard offering provides desalted gRNAs, but purified versions, with custom mod patterns, large scale deliverables and likewise gRNAs for alternative applications built on Cas9 (i.e. Prime Editing) can easily be accommodated
- **Scales** – 2nmol, 10nmol, 50nmol, 100nmol
- Available in tube or plates
- **Custom formats** – IDT can support any format to suit your needs
  - Arrayed – individual guides targeting different genes in different wells
  - Pooled – multi-guide pooled by gene
- **Custom Plates** – IDT has a variety for you to choose from including ECHO plates
- **Need help designing?** We can assist.



# CRISPR SCREENS WITH CUSTOM Alt-R gRNA LIBRARIES



4 μM RNP Nucleofector™ Nucleofection (Lonza) into HAP1 cells, 4 gRNAs for 5 genes as 2-part or sgRNA gave >85% avg editing at all target loci

# ALT-R™ CRISPR CUSTOM GUIDE RNAS

## CUSTOM ORDERING TOOL FOR ANY gRNAs

**ORDER MENU** You are using the IDT KK portal.

### Custom CRISPR gRNA ordering

Select All ACTIONS: ▾ # of Items: 1 GO BULK INPUT 📄

# 1  ✖ ℹ 🗑

**Scale** ℹ

Custom Alt-R™ gRNA, 2 nmol ▾

**Sequence** ✖ (5' → 3')

5' MOD ▾ INTERNAL ▾ 3' MOD ▾ BASES ▾

/AltR1/rArU rGrCrA rUrArG rCrUrA rGrArC rUrArG  
rArUrC rArGrA rCrUrC rUrCrU rCrGrA rUrCrA

# Bases: 38 (Min:30 Max:150)

GC: 44.7% Tm: 64.8°C ⚙ DeltaG: -62.25 kcal/mole

CONVERT TO RNA

Step 1: Enter Sequences (1 item)

Step 2: Order Custom CRISPR Essentials (0 items)

CONTINUE >

Show CRISPR Help

- Online ordering of any custom gRNA: including Cas13, pegRNA, gRNAs for novel nuclease systems
- Lengths: 30 to 150 nt
- Scales: 2, 10, 50, & 100 nmols
- Supports: RNA bases, Alt-R modifications, 2'OMe bases, and phosphorothioate linkages

# IDT CRISPR PROTEIN ENGINEERING

- Cas9 vs. Cas12a (Cpf1)
- Alt-R WT Cas9 vs. **Alt-R HiFi Cas9**
  - Problem with Cas9: off-target editing
  - Bacterial mutagenesis to evolve high-fidelity Cas9
- Alt-R WT Cas12a (Cpf1) vs. **Alt-R Cas12a (Cpf1) Ultra**
  - Problem with Cpf1: low on-target editing
  - Bacterial mutagenesis to evolve enhanced activity A.s. Cas12a
  - Actively working on developing mutants of *L.b.* Cas12a with increased activity
- Fluorescent CRISPR fusion proteins:
  - **Alt-R™ S.p. Cas9-GFP V3, 100 µg**
  - **Alt-R™ S.p. Cas9-RFP V3, 100 µg**

IDT-exclusive mutants

Confidential - Company Proprietary

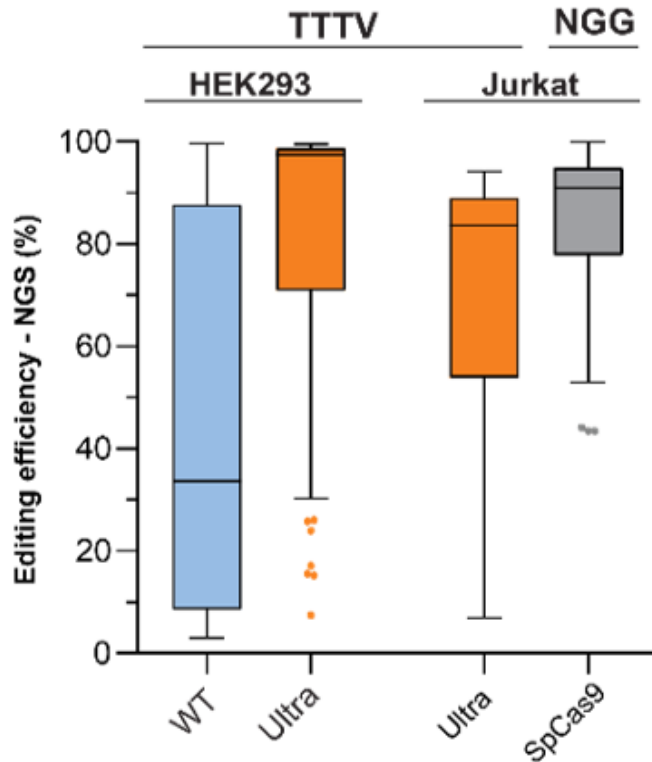
# NUCLEASE

Product Name	Catalog #
Alt-R™ S.p. Cas9 Nuclease V3, 5 mg	10000735
Alt-R™ A.s. Cas12a (Cpf1) Ultra, 5 mg	10007804
Alt-R™ S.p. HiFi Cas9 Nuclease V3, 5 mg	10007803
Alt-R™ Cas9 Elec Enhancer, 100 nmol	10007805
Alt-R™ S.p.Cas9 V3, glycerol-free, 100µg	10007806
Alt-R™ S.p.Cas9 V3, glycerol-free, 500µg	10007807
Alt-R™ S.p.Cas9 V3, glycerol-free, 5 mg	10007808
Alt-R™ L.b. Cas12a (Cpf1) Ultra, 100 µg	10007922
Alt-R™ L.b. Cas12a (Cpf1) Ultra, 500 µg	10007923
Alt-R™ L.b. Cas12a (Cpf1) Ultra, 5 mg	10007924

## Large Scale Cas9 Proteins:

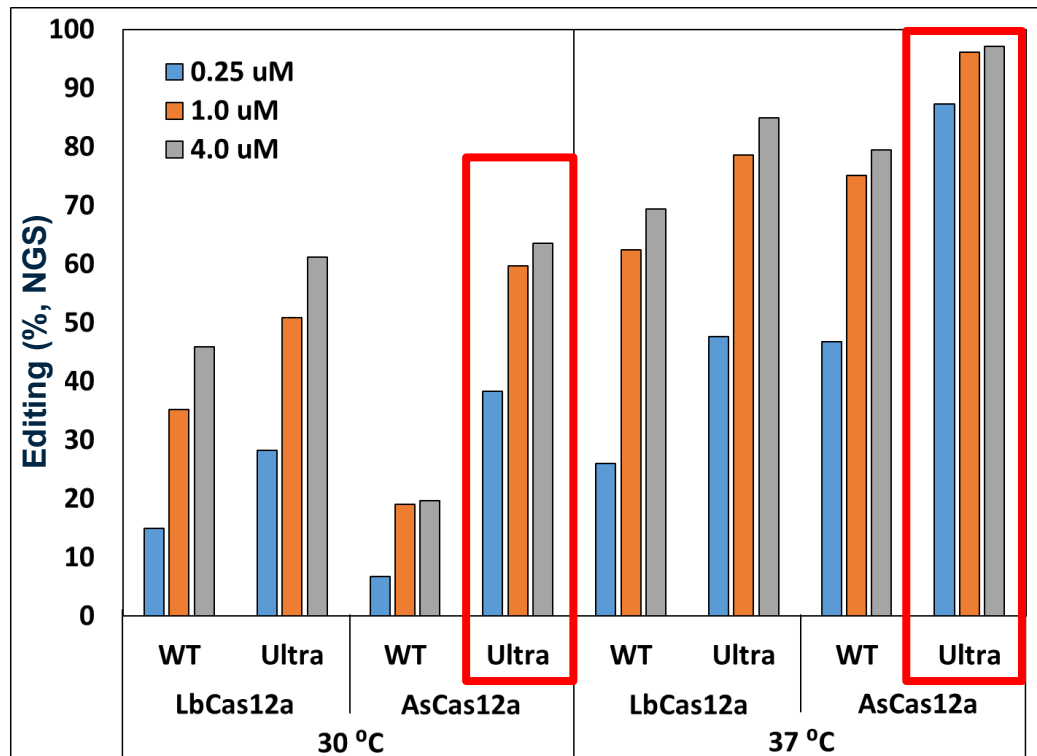
- Glycerol-free WTCas9v3 – ideal for sensitive cell lines
- As and Lbcas12a (Ultra) – useful for targeting AT-rich regions without available Cas9-specific PAM sequences. Much higher on-target potency than wild-type A.s. Cas12a (Cpf1). Can recognize many TTTT PAM sites in addition to TTTV motifs, increasing target range. Active at room temperature, making them flexible tools for applications requiring delivery at lower temperatures.

# A.s. Cas12a *ULTRA* HAS EDITING ACTIVITY COMPARABLE TO *S.p.* Cas9



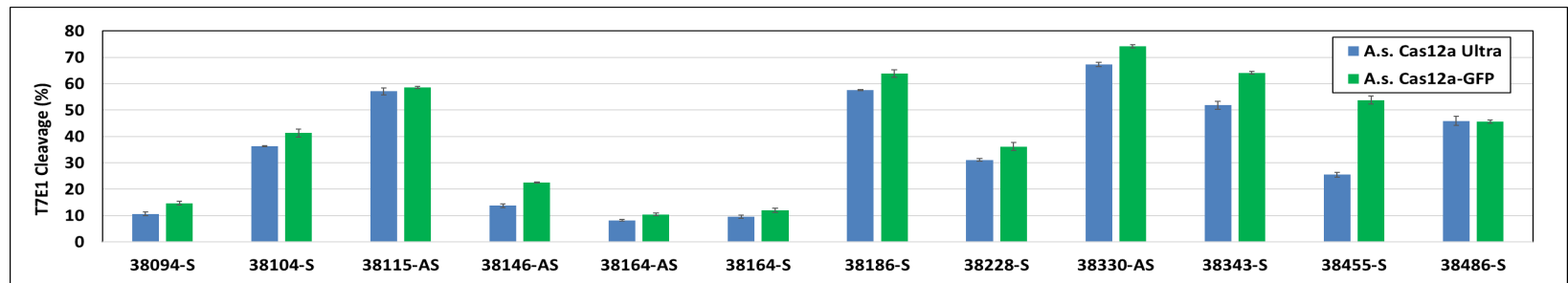
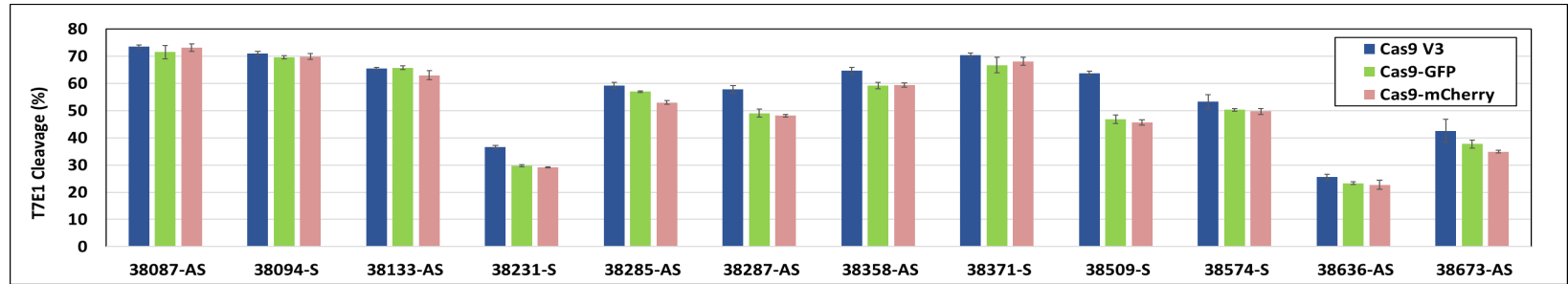
71 Cas9/Cas12a RNP (TTTV PAMs) targeting multiple loci delivered by electroporation in HEK293 and Jurkat cells

# A.s. AND L.b. Cas12a *ULTRA* HAVE INCREASED TEMPERATURE TOLERANCE



# IDT FLUORESCENT CRISPR PROTEINS RETAIN HIGH ON-TARGET ACTIVITY

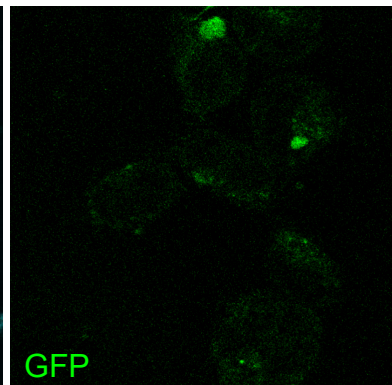
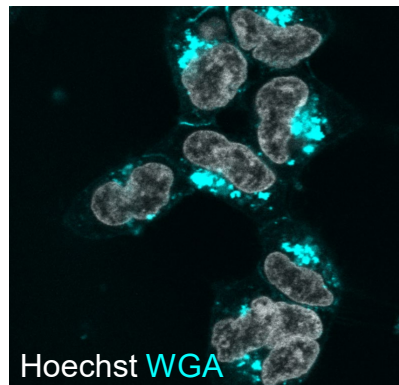
- *S.p.* Cas9 (Cas9 V3, Cas9-GFP, or Cas9-mCherry) was delivered at 2.0  $\mu$ M RNP into HEK293 cells by Nucleofector™ Nucleofection (Lonza) targeting sites within the HPRT gene
- *A.s.* Cas12a RNP (Cas12a *Ultra* or Cas12a-GFP) was delivered at a suboptimal dose of 50 nM RNP into HEK293 cells by nucleofection to achieve a range of editing across the tested HPRT sites



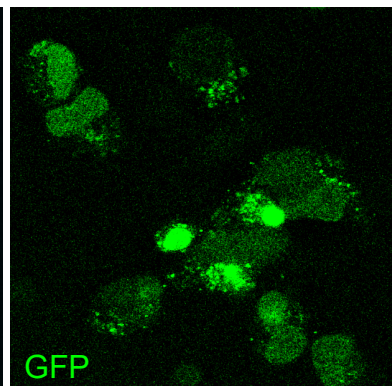
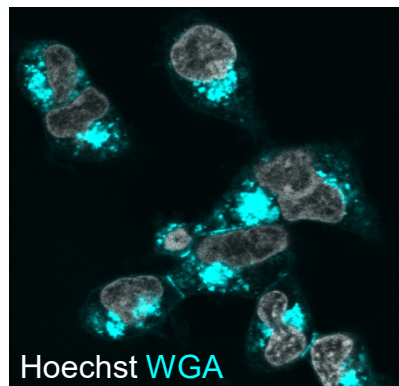
# Cas9-GFP ALLOWS FOR VISUALIZATION OF DELIVERY

2  $\mu$ M RNP delivery by Nucleofector™ Nucleofection (Lonza) into HEK293 cells, imaged ~18 hours after delivery using a confocal microscope

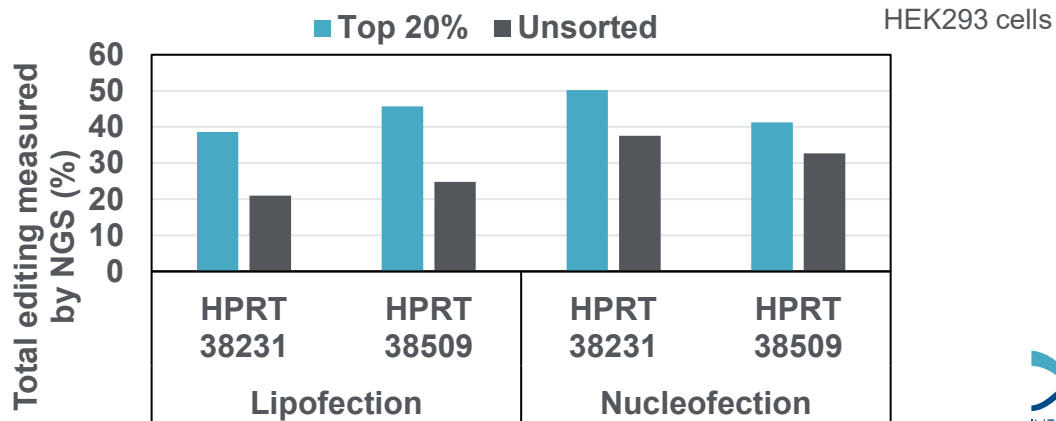
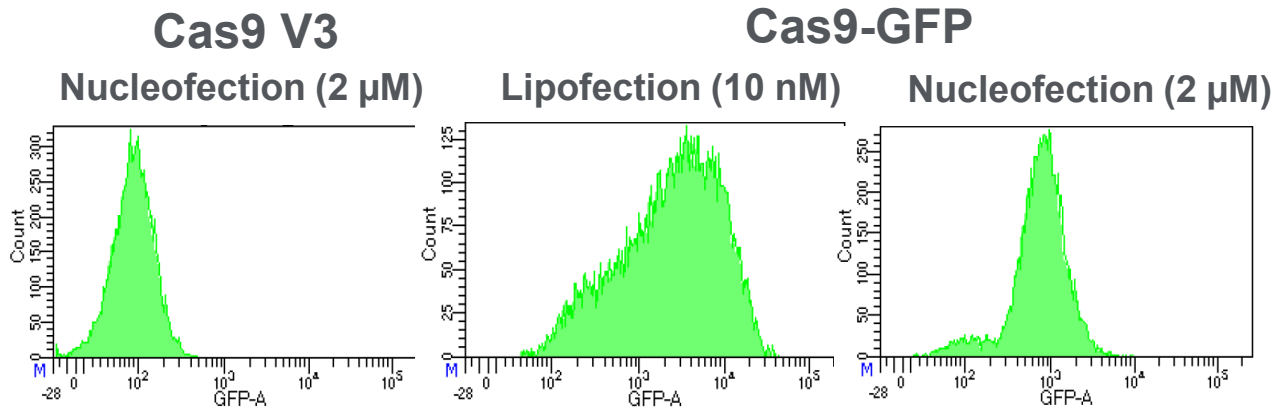
**Cas9  
V3**



**Cas9-  
GFP**

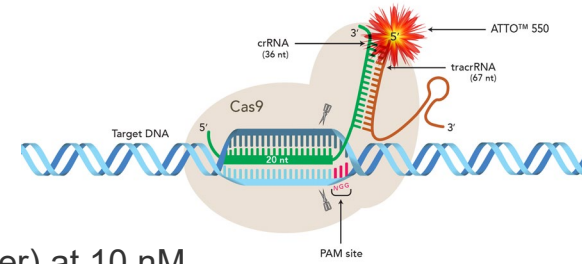


# DELIVERY PROFILE IS REFLECTIVE OF EDITING

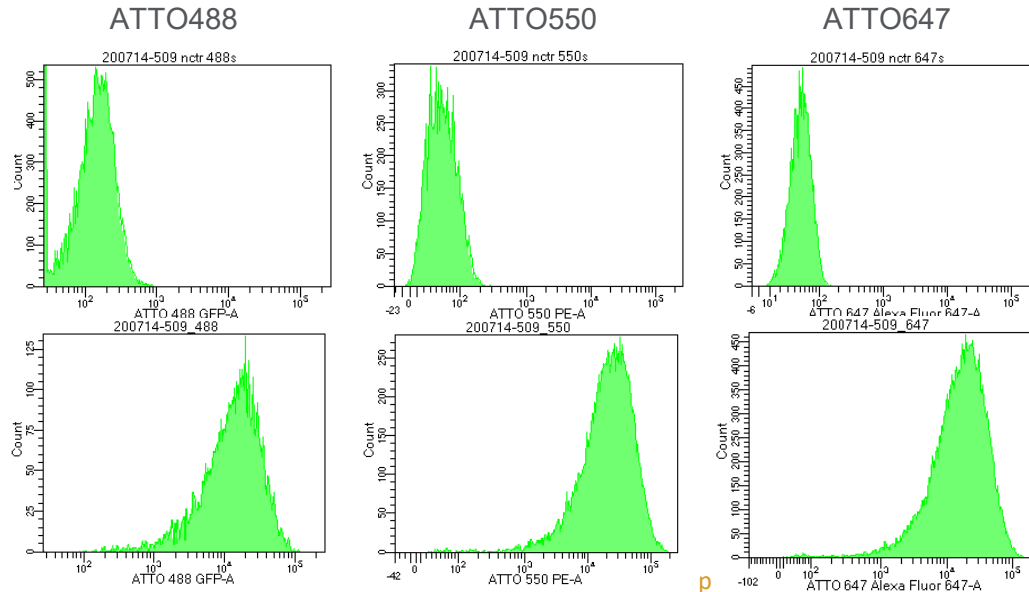


# FLUORESCENTLY-LABELED tracrRNA

- Labeled tracrRNA was duplexed with Alt-R™ crRNA XTs
- crRNA:tracrRNA duplex complexed with Alt-R™ Cas9 V3
- RNP was delivered into cells using Lipofectamine™ RNAiMAX (Thermo Fisher) at 10 nM
- HEK293 cells were incubated for ~18 h at 37°C.
- Cells were washed twice with PBS and run on a flow cytometer



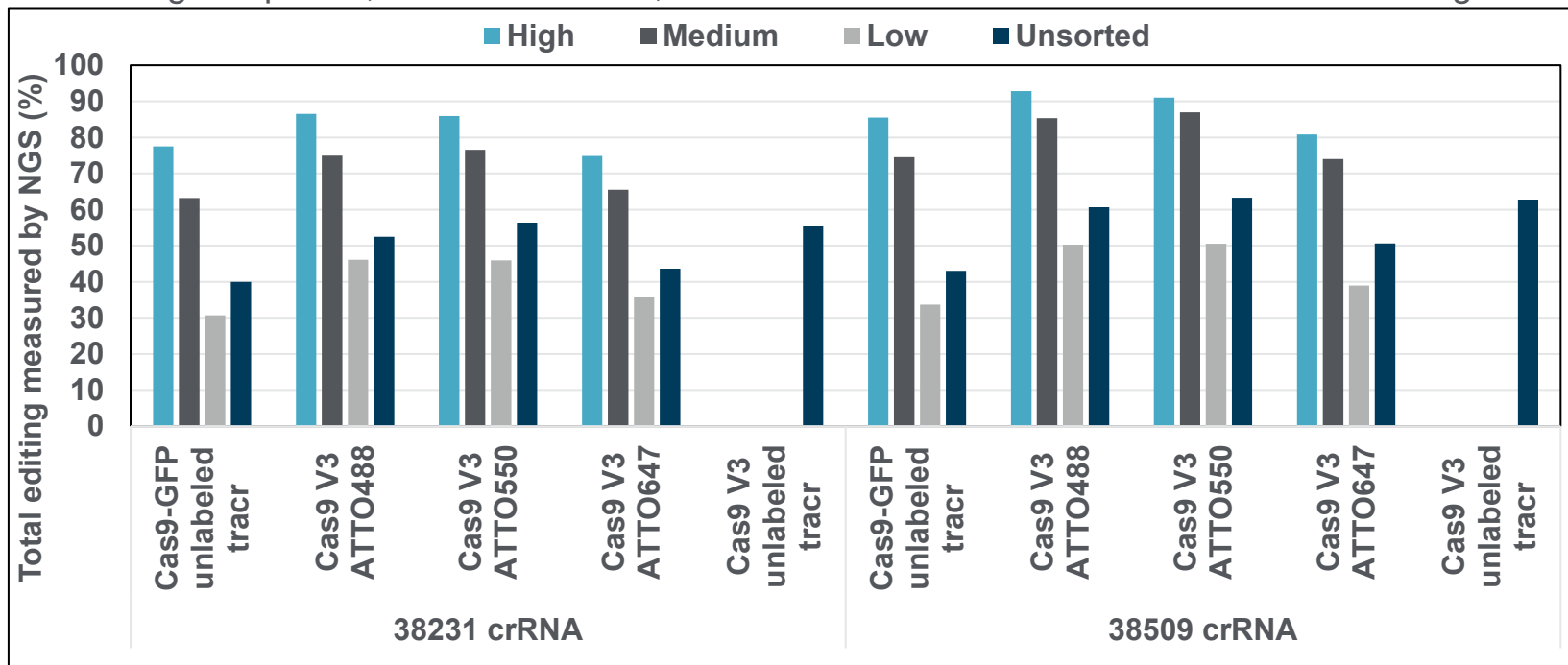
Cas9 V3  
+ Unlabeled  
tracrRNA



Cas9 V3  
+ Labeled  
tracrRNA

# ENRICHMENT OF EDITED CELLS USING LABELED tracrRNA BY FACS

High: Top 20%, Medium 80–60%, Low: Bottom 60% of cells based on fluorescent signal



(10 nM RNP delivery by Lipofectamine™ RNAiMAX (Thermo Fisher) into HEK293 cells, sorted after ~18 hours)

# IDT SOLUTIONS TO IMPROVE HOMOLOGY-DIRECTED REPAIR

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# USE THE Alt-R HDR DESIGN TOOL FOR OPTIMAL DESIGN

## Alt-R HDR Design Tool

Design and order homology-directed repair (HDR) donor templates and associated Cas9 guide RNAs for genome editing human, mouse, rat, zebrafish, or C. elegans targets.

[← BACK](#) **1. PICK TARGET(S)** **2. SPECIFY DESIGN** [WATCH VIDEO ▶](#)

Visualize & specify your mutation by opening the sliders on the interactive map or input fields below. Need additional assistance? Click 'Watch video' to learn more.

NCBI Transcript Accession: NM\_000518.4

Exon Intron guide RNA Reference Sequence SNP/MNP Insertion Deletion

5226994 5227026

5' 3'

Start edit: 5226994 SNP/MNP/Deletion length: 0 [ZOOM TO EDIT](#) [PREVIEW TRANSLATION](#) [← PREVIOUS](#) [NEXT →](#)

Mutation:  [Select tag \(optional\)](#) [INSERT](#)

Homology arms: Left  Right   Use default homology

Number of designs: 5 Add silent mutations:  No  Yes

Guide: GTAACGGCAGACTTCTCCTC

[← BACK](#) [DESIGN >](#)

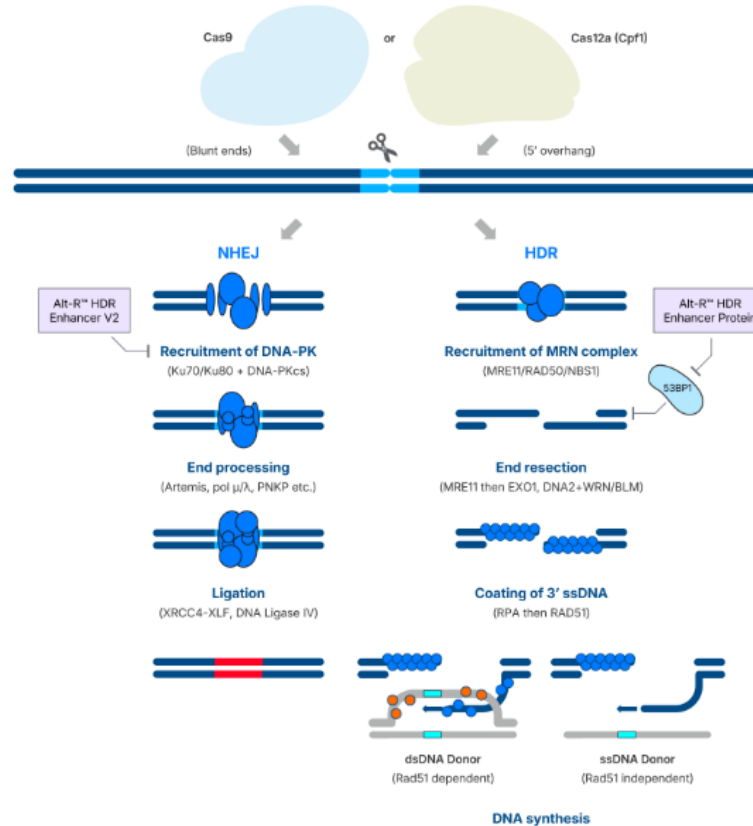
For S.p. Cas9 designs, guidelines based on empirical data are used to recommend donors that are associated with guide RNAs that are near the desired mutation and have both high on-target and off-target scores. For S.p. Cas9 D10A nickase designs, guidelines based on empirical data were used to recommend paired guide RNAs that flank the desired mutation, have PAM sites oriented away from the mutation, and have cut sites 37–70 nucleotides apart.

## Alt-R HDR Design tool: a novel bioinformatics tool for ssDNA HDR template design

- Human, mouse, rat, zebrafish, nematode, or custom input
- gRNA selection using IDT Alt-R gRNA design tool
- Addition of silent mutations improves rates of HDR
- Supports WT and Cas9 Nickase (D10A) strategies
- Single or multiple (batch) design
- Alt-R modified donors available in output

# HDR ENHANCER PROTEIN

## Alt-R HDR Enhancer V2



CRISPR-induced double-strand breaks are repaired via NHEJ or HDR. The Alt-R™ HDR Enhancer Protein promotes HDR by inhibiting 53BP1, making it specific to the HDR pathway, in contrast to DNA-PK inhibitors that enhance HDR by blocking the NHEJ pathway directly.

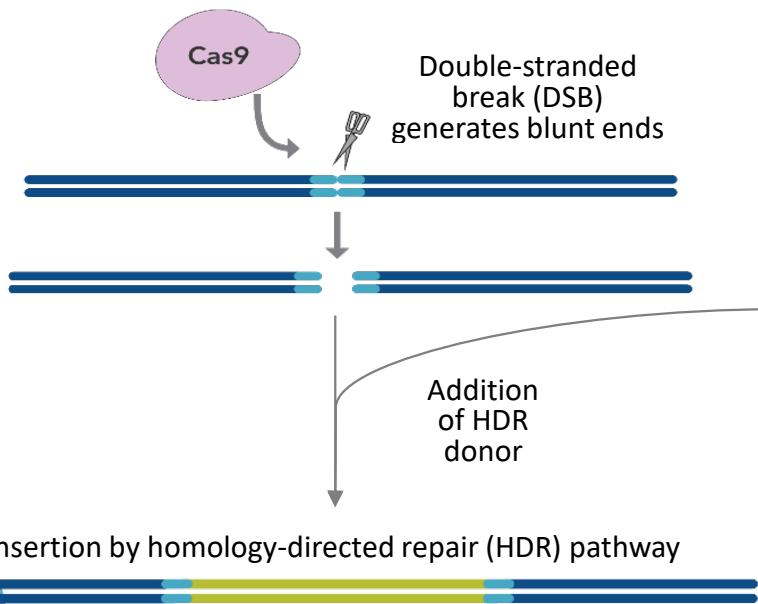
New

Alt-R™ HDR Enhancer Protein

# HDR ENHANCER V2 OR PROTEIN

項目	HDR Enhancer V2	HDR Enhancer Protein
分類	小分子	タンパク質
作用の考え方	NHEJ を弱める	HDR のブレーキ (53BP1) を外す
作用点	経路レベル	分子レベル (53BP1)
性格	出力強め	精密・安全寄り
プロトコール上の位置	核酸導入後 (培養中)	RNPと同時導入
「効く時間帯」	DSB修復が進行している間	DSB発生直後

# ノックイン: 大きなサイズのテンプレートの選択肢 (120塩基～)



- ① ssDNA donor with ends homologous to DSB
- ② dsDNA donor with ends homologous to DSB
- ③ Plasmid donor with ends homologous to DSB

▪ Ultramer®  
Oligonucleotide:ssODNドナーテンプレート(200mer)

▪ gBlocks® Gene Fragments  
(2本鎖DNAフラグメント 125～3000bp)

▪ 人工遺伝子合成  
(Genes 40円/bp)

# HDR ドナーテンプレート まとめ

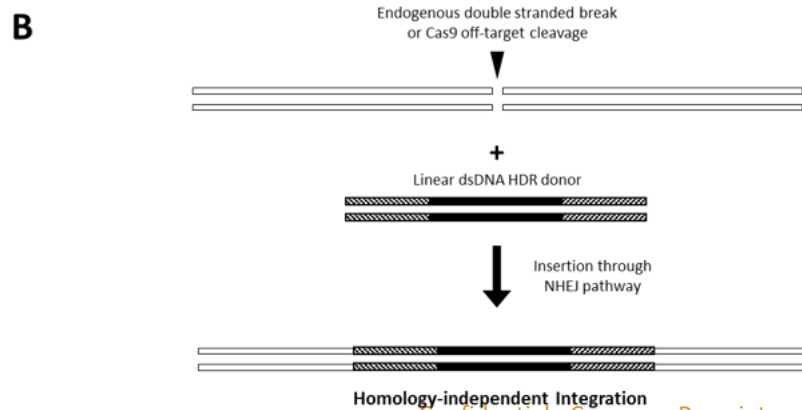
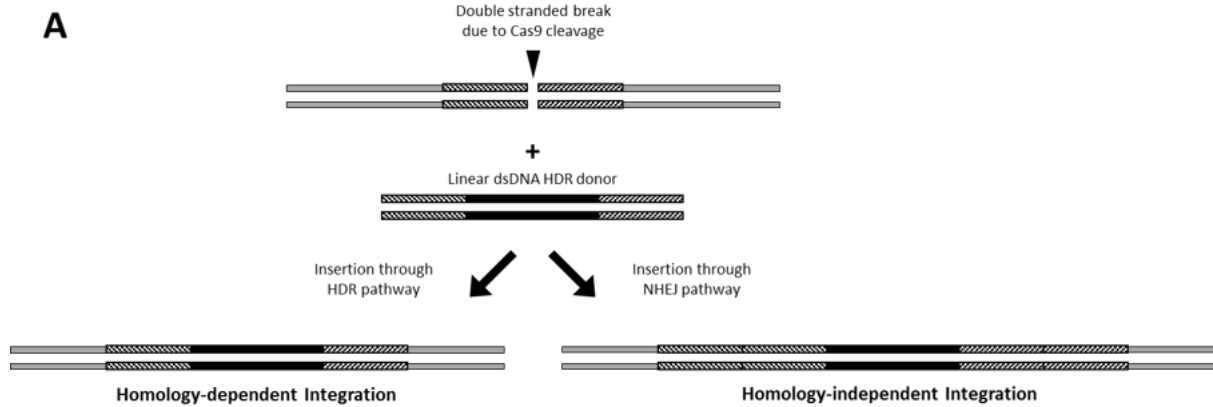
タイプ	製品	サイズ	毒性	HDR効率	コスト	納期	HDR Enhancer 併用時の効率 *
ssODN	Alt-R HDR Donor Oligo/ PSmod	45- 200mer	低い	++++	安価	5-10営業 日	✓✓
					¥17,560~		
dsDNA (直鎖二本鎖)	Alt-R™ HDR Donor Blocks	201- 3,000bp	低い~中 程度	+++	比較的安価	3週間~	✓✓
					¥34,620 or ¥72/bp~		
dsDNA (環状)	Genes	25- 3000bp、 それ以上	より高い	+	高価	数週間	✓
					¥40/bp~		

# 大きいサイズのドナーテンプレートの特長

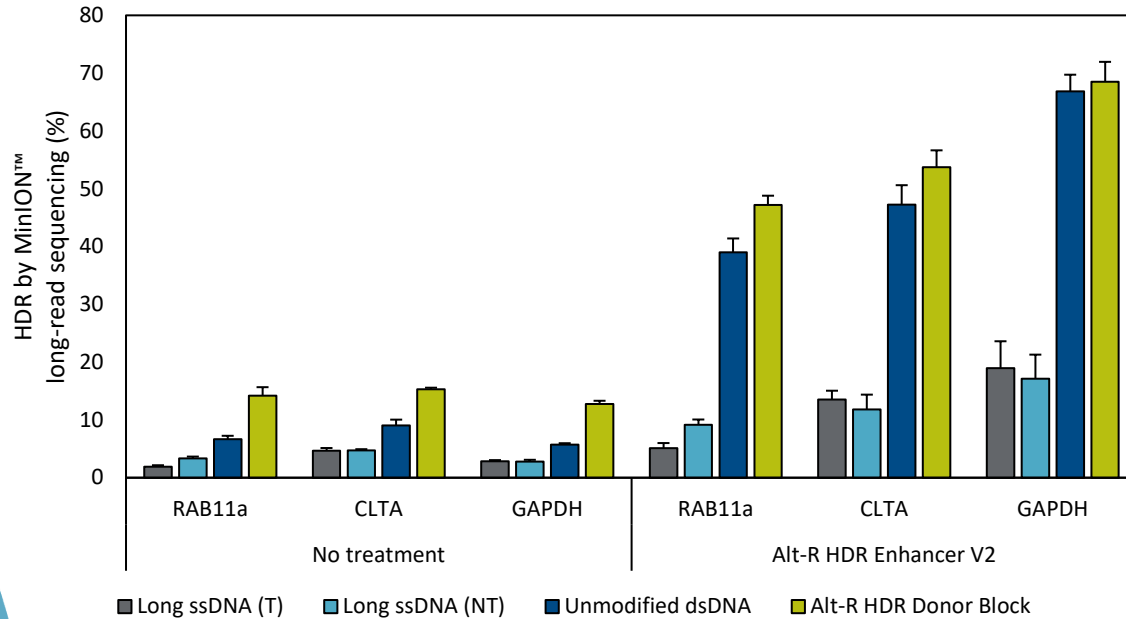
	ドナータイプ		
	Plasmid	Linear ssDNA	Linear dsDNA
HDR効率	低	高	高
細胞毒性	高	中	中
コストと納期	低	高	低～中
Off-target挿入リスク	中	低	中

Can modifications reduce the risk of blunt integration with dsDNA?

# SOLUTIONS TO REDUCE BLUNT INTEGRATION WITH dsDNA HDR TEMPLATES

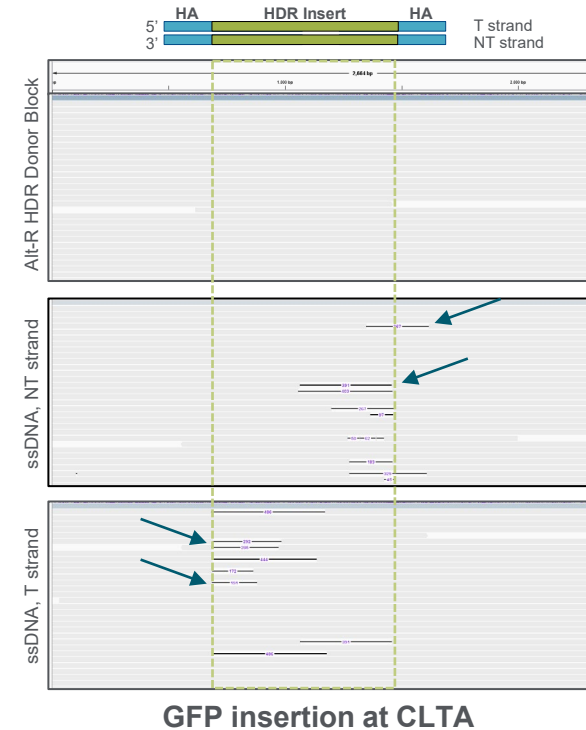


# Alt-R HDR Donor BlocksとAlt-R HDR Enhancer V2の併用が large HDR insertions効率を改善



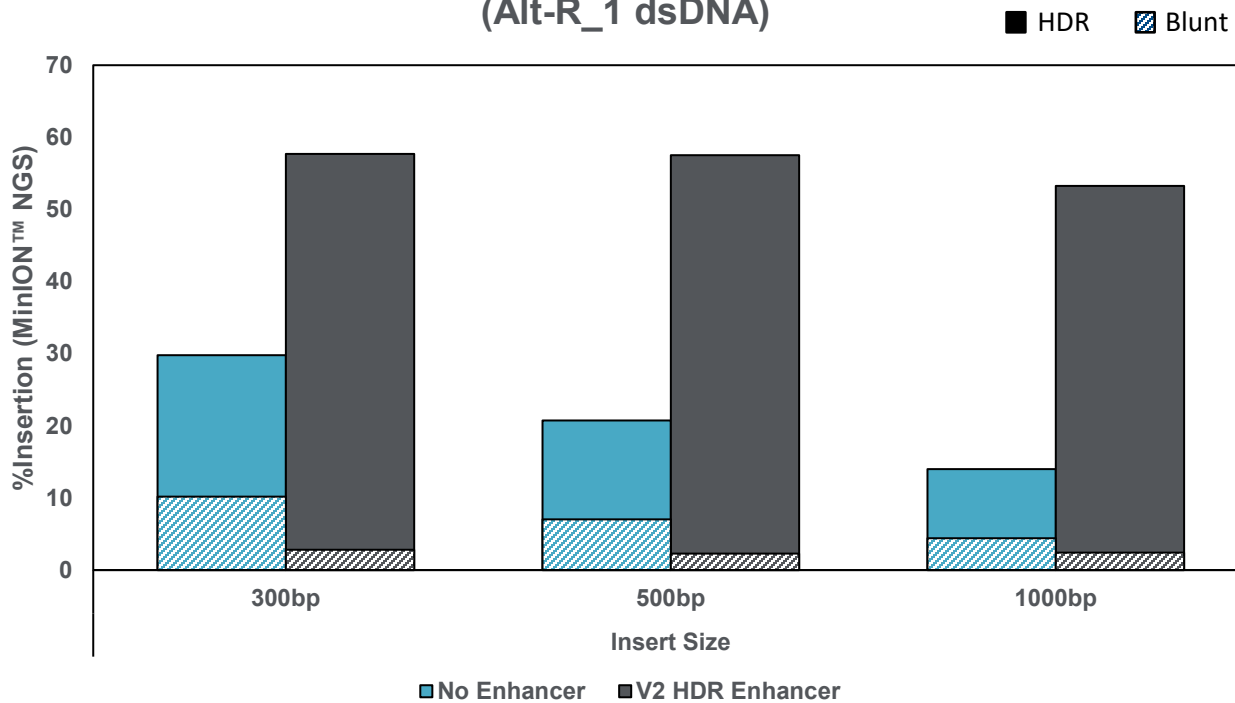
*Lonza Nucleofector™ delivery system:*

K562 or HEK-293 cells, 2  $\mu$ M Cas9 RNP, 50 nM ss or dsDNA (GFP tag, 200bp HA), 0 or 1  $\mu$ M Alt-R HDR Enhancer V2 (24 hrs) (n=4)



# HDR ENHANCER V2 ALSO IMPROVES HDR RATES FOR LARGE INSERTIONS

Use of V2 HDR Enhancer with modified dsDNA donors  
(Alt-R<sub>1</sub> dsDNA)



# NGSでのオフターゲット解析

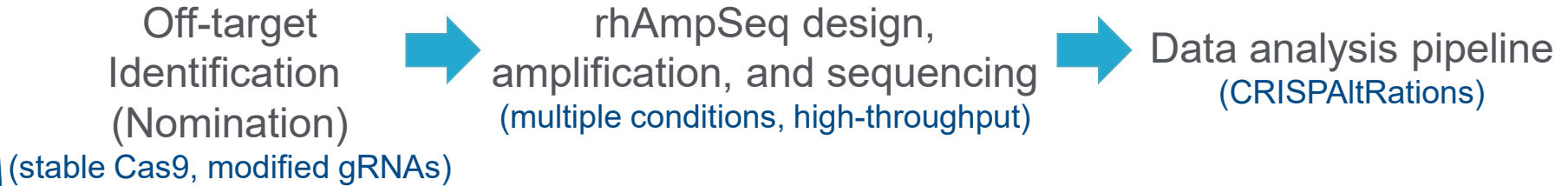
## rhAmpSeq™ MULTIPLEXED AMPLICON SEQUENCING FOR CRISPR

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# CRISPR 編集・オフターゲットの検出

- 多くの検証・予測されたオフターゲットサイト-ガイドRNAの特異性
  - In silicoの予測ツール
    - 予測の困難性、重要なサイトの見落とし、予測過剰
  - 可能性のあるオフターゲット効果を定義するIn vitro アッセイ
    - SITE-seq, CIRCLE-seq など.
    - 予測過剰傾向、シーケンスリード・スペースの無駄
  - GUIDEseqによる オフターゲット効果の識別
    - 目的の細胞株による、定量的
  - rhAmpSeq™ for CRISPR
    - マルチプレックス・アンプリコンベースのターゲットエンリッチメントNGS解析
    - オンターゲットサイトに対しての定量的評価、最大1000個までのオフターゲットサイトのマルチプレックス反応が可能

# EXAMPLE CRISPR EXPERIMENT WORKFLOW





# rhAmp PCR TECHNOLOGY INNOVATED AT IDT

*BMC Biotechnology* (2011) 11:80

METHODOLOGY ARTICLE

Open Access

## RNase H-dependent PCR (rhPCR): improved specificity and single nucleotide polymorphism detection using blocked cleavable primers

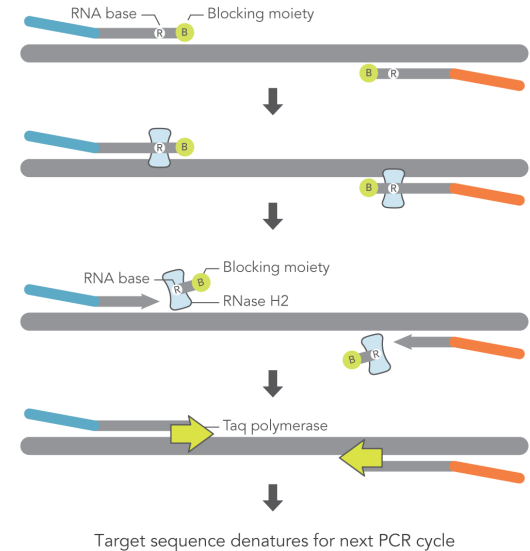
Joseph R Dobosy, Scott D Rose, Kristin R Beltz, Susan M Rupp, Kristy M Powers, Mark A Behlke<sup>\*</sup> and Joseph A Walder

Blocked inactive primers anneal to target

RNase H2 recognizes hybridized internal RNA base

RNase H2 cleaves hybridized primers

DNA polymerase extends newly unblocked primers



# rhAmpSeq DESIGN TOOL

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You are using the IDT portal.

## rhAmpSeq Design Tool

Welcome to the IDT rhAmpSeq design tool. If you encounter any questions while using the tool, contact [applicationsupport@idtdna.com](mailto:applicationsupport@idtdna.com). If you don't see what you're looking for, submit a request to the rhAmpSeq Custom Design Service here.

Create new design

Review design results

Dashboard

Application

Input format

Species

Minimum insert size

Maximum insert size

Panel name

DESIGN

CLEAR AND RESET ↶

[Open tool resources](#)

[Open user guide](#)

Paste/Type input

Upload file

Enter your coordinates separated by tab below.

chr7 12345 12365

Proprietary



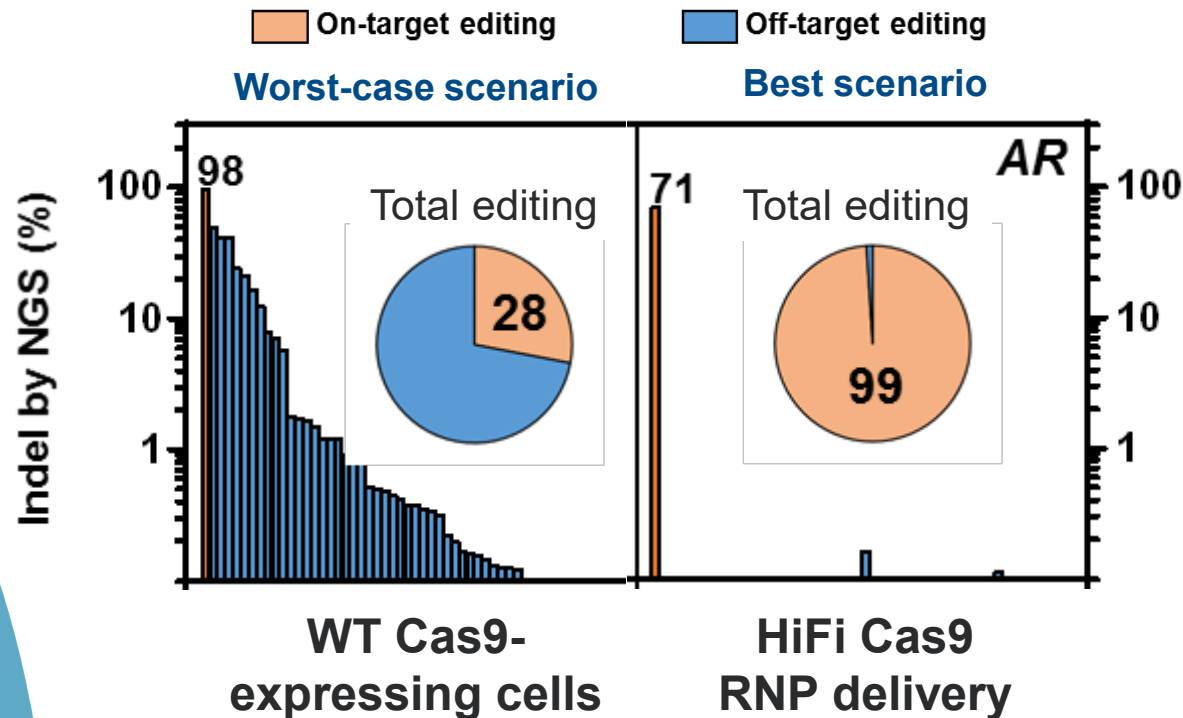
# rhAmpSeq DESIGN TOOL

## Supported Reference Genomes

Species	Reference genome
<i>Homo sapiens</i> (human)	GRCh38
<i>Homo sapiens</i> (human)	GRCh37
<i>Mus musculus</i> (mouse)	GRCm38
<i>Rattus norvegicus</i> (rat)	Rnor6.0
<i>Bos taurus</i> (cow)	ARS UCD1.2
<i>Bos taurus</i> (cow)	UMD3.1
<i>Brassica napus</i> (canola)	Bra napus v2.0
<i>Caenorhabditis elegans</i> (nematode)	WBcel235
<i>Danio rerio</i> (zebrafish)	GRCz11
<i>Gallus gallus</i> (chicken)	GRCg6a
<i>Glycine max</i> (soybean)	Glycine max v2.1
<i>Gossypium hirsutum</i> (cotton)	ASM98774v1
<i>Oryza sativa</i> (rice)	IRGSP 1.0
<i>Solanum lycopersicum</i> (tomato)	SL3.0
<i>Solanum tuberosum</i> (potato)	SolTub 3.0
<i>Sus scrofa</i> (pig)	Sscrofa11.1
<i>Triticum aestivum</i> (wheat)	IWGSC v1.0
<i>Zea mays</i> (maize)	B73 RefGen v4

# rhAmpSeq TECHNOLOGY AND Alt-R HiFi Cas9

Verification and validation Alt-R *S.p.* HiFi Cas9 Nuclease V3



## rhAmpSeq system savings:

- 8 x 40 assay designs
- ~1500 individual PCRs reduced to <96
  - Master mix
  - gDNA
- Library quantification
- Full-time equivalent hours
  - Months to days

【基礎研究から臨床応用まで利用可能 : GMPグレード製品】

## Aldevron CRISPRヌクレアーゼ



遺伝子編集を利用して遺伝子疾患や遺伝性疾患の調査・治療を行うには、このヌクレアーゼが臨床への近道です。なぜなら研究グレードと同等で、完全なcGMPで製造されており、規制当局への提出に必要な品質文書でサポートされているからです。

CRISPR with confidence. 詳しくはこちらから。

### Aldevron SpCas9 Nuclease and SpyFi® Nuclease

ADD TO ORDER

Recombinant *Streptococcus pyogenes* Cas9 nuclease, purified from an *E. coli* strain expressing the SpCas9 nuclease. Contains N- and C-terminal SV40 nuclear localization sequences (sNLS) and no affinity purification tags. Provided in solution at 10 mg/mL. SpyFi® Cas9 nuclease has similar on-target potency to wild-type SpCas9, but with reduced off-target effects, allowing for more precise genome editing.

Quantity	Product	Catalog #	Price
<input type="text" value="0"/>	sNLS-SpCas9-sNLS Nuclease, 0.25 MG	10017687	¥44,760 JPY
<input type="text" value="0"/>	sNLS-SpCas9-sNLS Nuclease, 5 MG	10017688	¥744,600 JPY
<input type="text" value="0"/>	SpyFi Cas9 Nuclease, 0.25 MG	10017689	¥69,700 JPY
<input type="text" value="0"/>	SpyFi Cas9 Nuclease, 5 MG	10017690	¥946,500 JPY

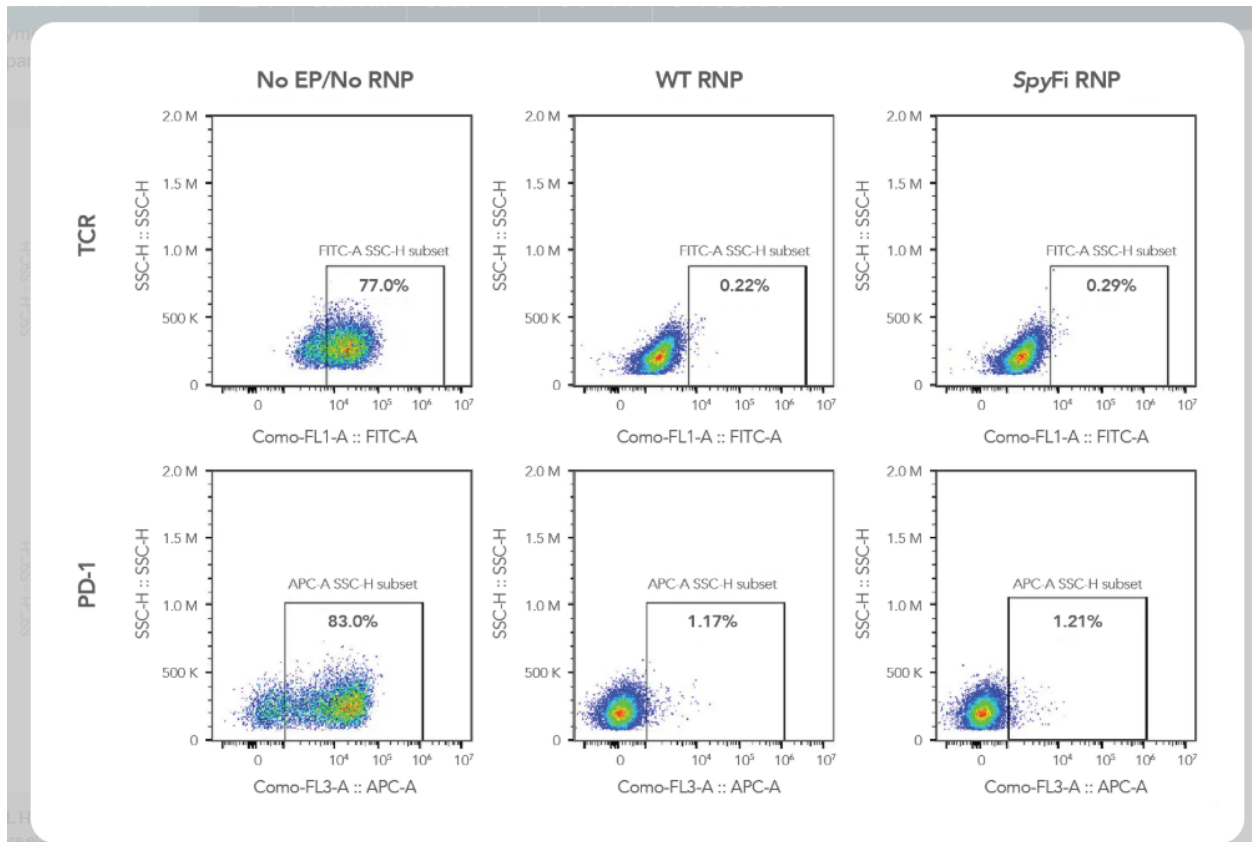
### Aldevron Eureka-V™ Nuclease

ADD TO ORDER

Recombinant Class 2, Type V CRISPR-Cas nuclease, purified from an *E. coli* strain expressing the recombinant nuclease. Contains N- and C-terminal SV40 nuclear localization sequences (sNLS) and no affinity purification tags. Provided in solution at 10 mg/mL. Requires acknowledgment of Limited Use Label License (LULL).

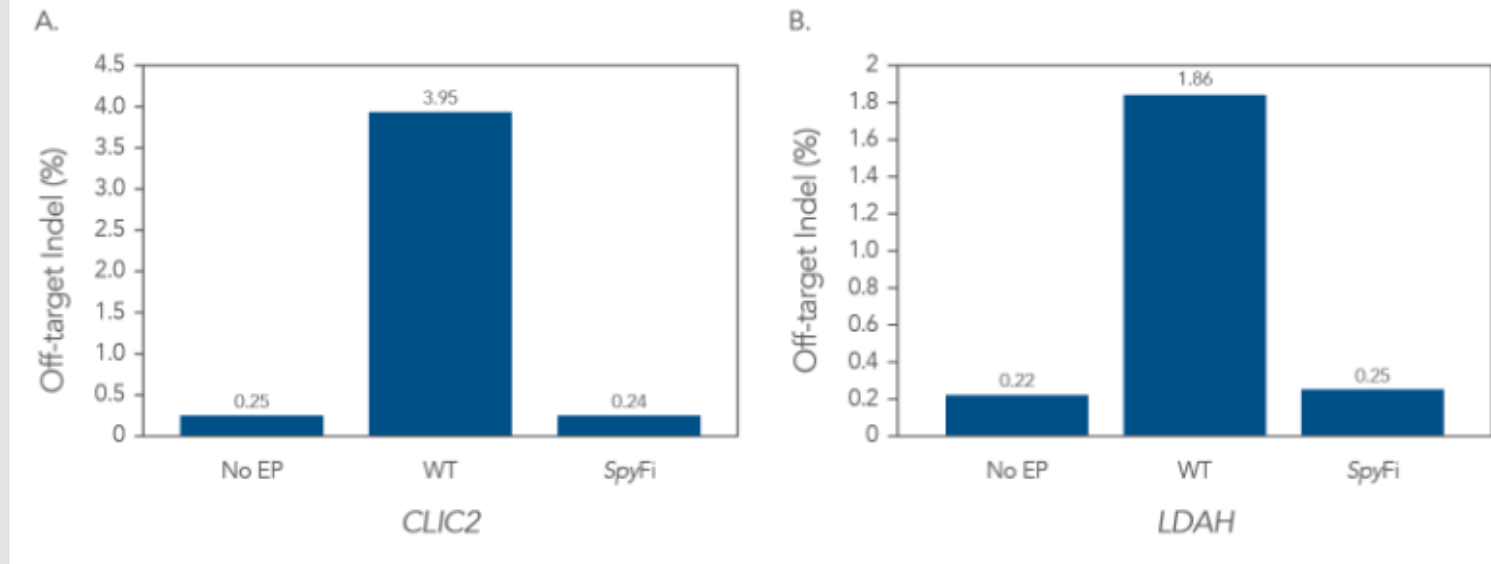
Quantity	Product	Catalog #	Price
<input type="text" value="0"/>	Eureka-V Nuclease, 1MG	10017800	¥211,500 JPY
<input type="text" value="0"/>	Eureka-V Nuclease, 5MG	10017811	¥964,700 JPY

# High-efficiency multiplexed T cell editing



**High multiplexed knockout efficiency with wild type (WT) and SpyFi RNP.** To knockout both *TCR* and *PD-1*, activated human T cells were electroporated with an equimolar mixture of RNP targeting *TRAC*, *TRBC*, and *PD-1*. Expression was measured by flow cytometry 5 days post electroporation. Expression was substantially reduced following electroporation with WT or SpyFi RNPs compared to no electroporated cells,  $n = 1$ .

## Reduced off-target effects



**Figure 2. Fewer off-target events with *SpyFi* vs WT RNP.** SpCas9 or *SpyFi* Cas9 RNP targeting the *TRAC* locus was transfected into activated human T cells using Maxcyte electroporation (EP) technology. The frequency of off-targeting editing for each RNP was determined by measuring InDel percentages at four off-target binding sites previously identified by iGUIDE [1]. DNA flanking off-target binding sites in *CLIC2*, *LDAH*, *ANKS1B*, and *ADCY10* were amplified by PCR, gel-purified, and sequenced by next generation sequencing (NGS). (A) At the *CLIC2* locus *SpyFi* RNP reduced off target editing by 94% compared to WT RNP. (B) Similarly, at the *LDAH* locus there was an 87% reduction in off-target editing. At both these loci *SpyFi* had off-target effects comparable to no electroporation controls. At two other sites (*ANKS1B* and *ADCY10*), off-target editing was generally less frequent but was diminished by *SpyFi* (33% less at *ANKS1B* locus, 15% less at *ADCY10*, data not shown),  $n=1$ .

# CRISPR cGMP gRNA Manufacturing Overview



For customers looking to accelerate their CRISPR gene editing program from discovery to clinical trials, our Engineering Run and full cGMP Compliant gRNA Manufacturing Services offer a streamlined and regulated solution. Backed by comprehensive documentation, our guide RNA CRISPR services simplify regulatory filings, offering a straightforward path to clinical success.

**Lab to life changing advances. 私たちがお手伝いします。**

	Engineering Run	cGMP
Availability	Delivery times after placing the order	Delivery times after placing the order
Cleanroom	Certification not required	ISO 8 Clean Room - Certified
Changeover	In-place with cleaning validation	In-place with cleaning validation
Materials	QA release on raw materials	QA release on raw materials
Batch Records	Draft based on customer's specifications	Based on customer specification
Release Testing	Qualified methods	Validated methods

# ゲノム編集でのHIBITの利用

---

## 1. HiBiT とは何か

HiBiT は NanoLuc ルシフェラーゼを分割した 11 アミノ酸 (33 Nucleotide seq) の超小型ペプチドタグ

ゲノム編集で 内在性遺伝子の末端にノックインすると、細胞内で LgBiT と結合して強い発光シグナルを出す

## 2. ゲノム編集における最大のメリット

\*\*HDR ノックインが「成功したかどうか」を最短・最も確実に判断できる

『内在性タンパク質のノックイン成功を、迅速・高感度・定量的に評価できる点』

## 従来のノックイン確認方法の課題

方法	課題
PCR / シーケンス	時間がかかる、労力大、スループット低
Western blot	抗体依存、感度・特異性にばらつき
蛍光タグ (GFP等)	タグが大きく、HDR効率が下がる／機能影響

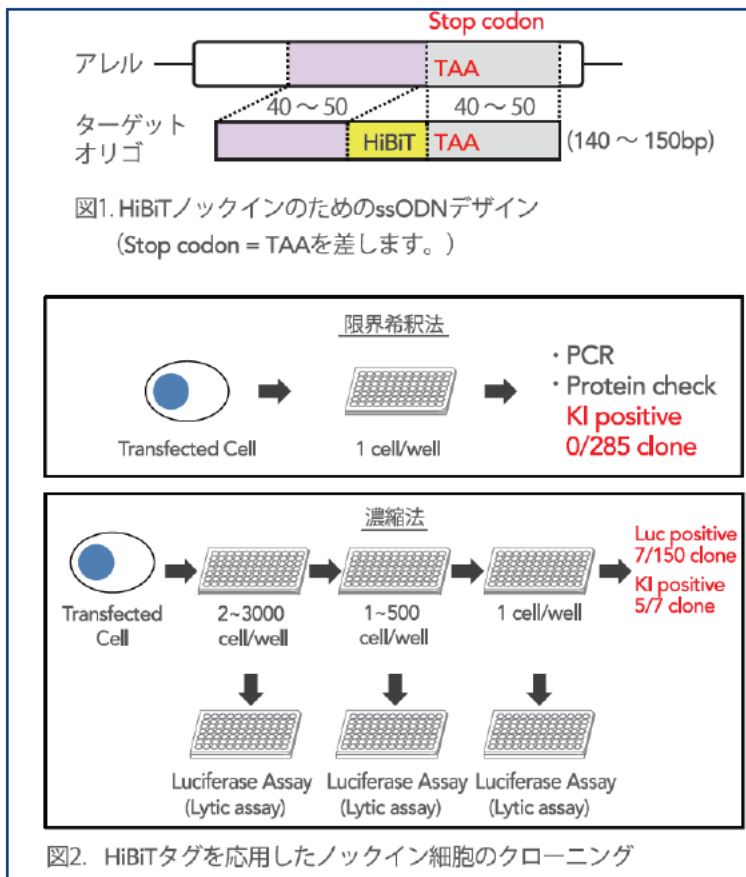
## HiBiT を使うと何がかわるか

ポイント	HiBiT の強み
タグサイズ	11 aa (極小) → HDR成功率が高い
検出	超高感度発光
定量性	発光=タンパク質量 (半定量~定量)
操作	細胞溶解→試薬添加のみ
スピード	編集後 24-48 時間で評価可能

「HDR がどれくらい起きたか」を“見える数字”で即判断できる  
タンパク質の定量が可能

**\* 注意点: タグ挿入位置の設計(最重要)、フレーム設計(HiBiT はここに非常に厳しい)など慎重に確認する事が重要なポイントとなります。万が一塩基にズレが生じた場合、フレームシフトが起きて正しいタンパク発現が起きずに、発光での検出も出来なくなってしまいます。**

# 例:細胞のクローニング作業を簡便化



# まとめ

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# ALT-R™ CRISPR CAS9 SYSTEM の初期費用の目安

## gRNA:2パートシステム

例①: ヒト/マウス/ラット対象、ターゲット遺伝子用とコントロール用、及びCas9タンパク質を購入

Alt-R® CRISPR crRNA, 2 nmol (ターゲット遺伝子) 10,300円

– 2 nmol Alt-R® CRISPR Control Kit 22,120円

• tracrRNA、crRNA (HPRT)、crRNA (Negative)、プライマーセット、バッファー

– Alt-R® S.p. Cas9 Nuclease V3, 100 µg 24,860円

初回費用定価合計  
¥57,280~

例②: ターゲット遺伝子用のみ購入、あるいはヒト/マウス/ラット以外の実験用に購入

– Alt-R® CRISPR crRNA, 2 nmol 10,300円

– 5 nmol Alt-R® CRISPR tracrRNA 12,160円

– Alt-R® S.p. Cas9 Nuclease V3, 100 µg 24,860円

初回費用定価合計  
¥47,320~

※ Alt-R® S.p. HiFi Cas9 Nuclease V3, 100 µg 30,200円

※ Alt-R® Genome Editing Detection Kit, 25 rxn 19,040円

※ 2パートシステムの場合、残ったtracrRNAは、新たに購入する別のcrRNA(10,300円)と一緒に使用できます。

★初回ご注文時は、該当品目に限り30%オフでのトライアルキャンペーンを実施しております

令和8年3月19日

第90回 共同研テクニカルセミナー  
ゲノム編集

令和8年 4月21日 火 14:00~15:30\*

\*質疑応答含む

【ハイブリッド形式】共同研棟7階セミナー・会議室 (D71-09) /zoom (事前登録制)

第1部 14:00~14:30

「ゲノム編集：遺伝子改変マウスからヒトiPS細胞まで」

講師：吉村 康秀助教 (大阪大学医学系研究科附属共同研ゲノム編集センター)

第2部 14:30~15:30

「明日から始めるゲノム編集」-CRISPRシステムを用いた  
強力でより正確なゲノム編集- 基礎~応用まで

講師：山田 聖子氏 (IDT株式会社)

ゲノム編集は遺伝子技術の中で、最も注目を浴びており、かつ様々な応用法について研究開発されている技術の一つです。


例えば医療分野では革新的治療法への応用に向けて日々急速に進歩しています。また、農業・工業分野においても食やエネルギー問題に対する問題解決に向けて研究が盛んに行われています。

世界最大級の研究用カスタム修飾合成品のサプライヤーであるIDT社では、以前からクローニングフリーの化学合成gRNAとリコンビナントCas9タンパク質とのRNA-タンパク複合体 (RNP) を細胞へ導入するAlt-R CRISPR Systemをご提供しています。明日からすぐに始められるAlt-R CRISPR Systemのご説明と、オンターゲット・オフターゲット切断の同定ツールの他、効率的なHDR実験のための製品・ツール情報、ゲノム編集後の解析法についてご紹介いたします。ゲノム編集の基礎から応用まで網羅したセミナーです。

↓ご登録はこちら

◆セミナートピック

- ・IDT Alt-R CRISPR システム概要 (Cas9/ Cas12a)
- ・蛋白質工学を用いた変異体の開発HiFi Cas9, Cas12a Ultra
- ・オンターゲット切断、オンターゲットの解析ツール
- ・相組み換え(HDR)によるノックイン実験の、効率最適化のためのHDRデザインツールとAlt-R HDR Donor template(ssDNA&dsDNA)について
- ・今後の製品開発における展望



事前登録制(締め切り4月21日(火)正午) 登録URL: <https://forms.office.com/r/HfSk9rEci>

◆参加ご希望の方に、別途セミナー参加用のInvitationを配信させていただきます

\*本セミナーは大学・研究機関・研究セミナー対象です\*

◆問い合わせ先◆大阪大学 医学系研究科附属共同研ゲノム編集センター (担当: 寺尾、内線3890)

◆大阪大学大学院医学系研究科附属共同研ゲノム編集センターゲノム編集サポート事業

◆ポイント

- ・IDTではゲノム編集関連製品をワークフローで揃える事が出来る
- ・設計サポート(有償)
- ・ゲノム編集分野では、高いシェアを誇り安心して使用できる
- ・製品開発に力を入れている

ご購入の際は、是非ゲノム編集サポート事業をご利用ください

◆共同研へのご質問・お問い合わせ  
order@ctrlab.med.osaka u.ac.jp

〒565-0871 大阪府吹田市山田丘 2-2  
大阪大学大学院医学系研究科附属共同研  
ゲノム編集センター



# THANK YOU



記載内容については予告なく変更される可能性があります。最新情報は弊社HPでご確認いただくか、下記までお問合せください。

カスタマーケア: [japan-custcare@idtdna.com](mailto:japan-custcare@idtdna.com)  
03-4510-4061