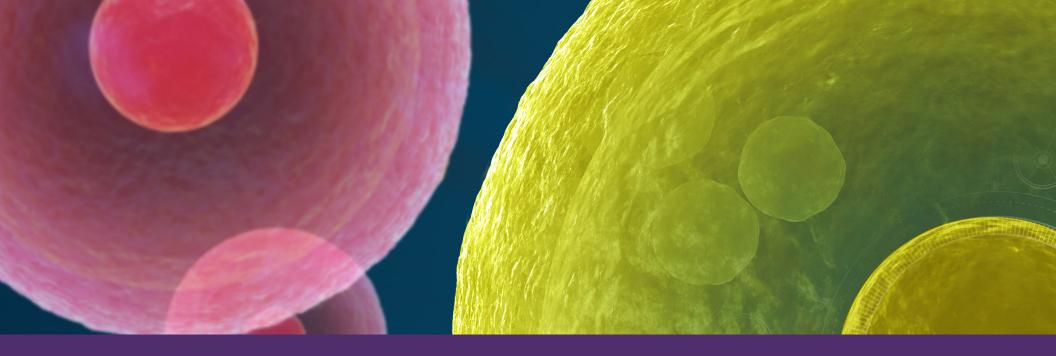
invitrogen



New libraries. New capabilities. New discoveries.





At Thermo Fisher Scientific, we are dedicated to driving innovative technologies to help accelerate your we are your trusted partner for products, services, and support throughout your functional genomics to work better together to deliver great results on your path to discovery.





research. As a leader in life science solutions, screen. Our combined solutions are designed



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Overview of screening solutions

Innovative solutions for every phase of the workflow

With advances in gene editing and RNA interference (RNAi) technologies, it is now possible to efficiently perform genomic screens in a wide variety of model systems. From assay development through target screening and *in vivo* studies, we offer a comprehensive portfolio of products and services for the functional genomics screening workflow (Figure 1). These products empower researchers to expedite discovery of gene targets by performing screens with speed, precision, and accuracy. In addition, our team of experienced scientists can provide custom services to help with your functional genomics screening needs.

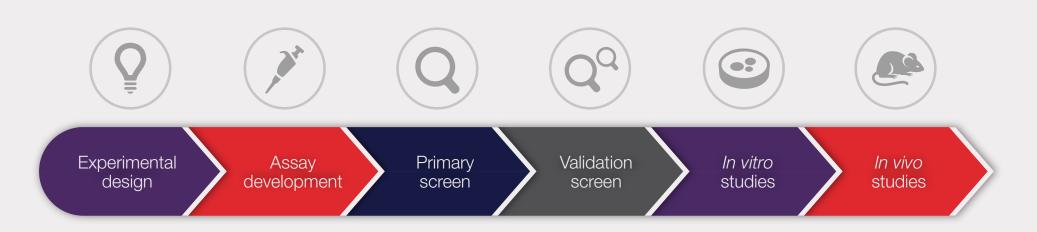


Figure 1. Typical functional genomics screening workflow.

Screening tools available

Accelerate your discovery with the right tool set

Start on the right path to discovery in your gene function studies with advanced tools for experimental design and assay development. Whether you're performing a functional genomics screen in a commonly used cell line or a difficult-to-transfect primary cell type, we offer a large portfolio of workflow

tools to enable superior results from your screening studies. Your choice of tool depends on your model system, the length of knockdown time you require, and other experimental parameters. An overview of our most effective technologies is shown in Table 1 to help you choose the solution that's right for you.

Table 1. Library formats available to choose from.

| | miRNA mimics and inhibitors | <i>Silencer</i> Select siRNA libraries | LentiPool CRISPR libraries | LentiArray CRISPR libraries |
|---------------------------|---|--|--|---|
| Description | Chemically modified, synthetic nucleic acids designed to either mimic mature miRNAs or bind to and inhibit endogenous miRNAs | Ultrapotent siRNAs to knock down gene expression | Pooled libraries of gRNAs in lentiviral expression constructs, providing a potent delivery method for complete gene knockout all in one tube | Arrayed libraries of gRNAs in lentiviral expression constructs, providing a potent delivery method for complete gene knockout |
| RNA type | miRNA oligonucleotide | siRNA oligonucleotide | Full-length gRNA | Full-length gRNA |
| Mechanism of action | Endogenous RNAi-induced silencing complex (RISC) | Endogenous RNAi-induced silencing complex (RISC) | Cas9 endonuclease-mediated indel formation | Cas9 endonuclease-mediated indel formation |
| Effect on gene expression | Transient knockdown | Transient knockdown | Permanent, complete knockout | Permanent, complete knockout |
| Time to effect | 2–3 days | 2–3 days | 2–4 days | 2–4 days |
| Species | Human, mouse, rat | Human, mouse, rat | Human | Human |
| Format | Array or pooled | Array or pooled | Pooled, ready-to-use lentivirus | Arrayed, ready-to-use lentivirus or glycerol stock of bacteria containing lentiviral vector |

Predefined libraries available

Predefined and custom libraries are available to enable you to focus on defined gene sets or perform more expanded screens (Figure 2). Our libraries are available in miRNA, siRNA, gRNA, and lentiviral formats.

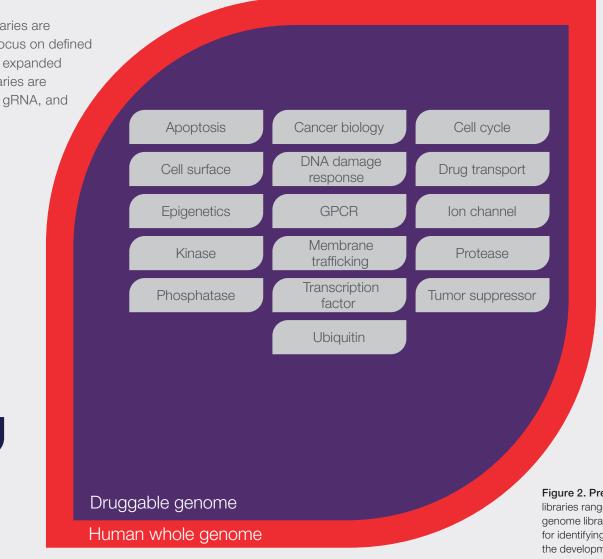


Figure 2. Predefined library options. Available libraries range from focused gene sets to whole-genome libraries. Our druggable genome library is ideal for identifying potential therapeutic targets involved in the development and progression of diseases.



CRISPR libraries

Bring the power of CRISPR-Cas9 technology to high-throughput screening

The CRISPR-Cas9 system is the premier technology for knocking out gene expression and is emerging as the next-generation tool of choice for high-throughput screening. The CRISPR-Cas9 system provides an efficient method for specific, complete, and permanent gene knockout. We are applying the power of the CRISPR-Cas9 system to high-throughput screening applications with our award-winning Invitrogen[™] LentiArray[™] libraries. These arrayed CRISPR libraries are designed to provide you with flexible systems that can be adapted to your needs and help drive new discoveries.



Features of LentiArray libraries

- Arrayed libraries in flexible formats to allow for a wide variety of assays
- Advanced gRNA designs for maximum knockout efficiency without sacrificing specificity
- Up to 4 high-quality gRNAs per gene target for efficient knockout in a wide variety of cell types
- Complete set of controls to streamline assay development
- 19 predefined libraries and custom options available, allowing you to focus on defined gene sets or perform unbiased whole-genome surveys
- Individual gRNAs and custom arrays available to support rapid assay development and validation of hits

Designed to enable your success

CRISPR libraries are constructed using our proprietary gRNA design algorithm, which incorporates the latest research and our extensive in-house experience. The gRNA designs are selected for maximum knockout efficiency without compromising specificity. For each gene target, we include up to 4 high-quality gRNAs to help ensure that the library will provide high-efficiency knockout of target genes across a wide array of cell types.



LentiArray CRISPR libraries

LentiArray CRISPR libraries are arrayed libraries of gRNAs that are constructed in lentiviral expression vectors, providing a potent delivery method that allows you to perform high-throughput screens in a wide variety of cell types. The libraries are available as either high-titer, ready-to-use lentivirus or as glycerol stocks of bacteria containing lentiviral vector (Figure 3). A full selection of controls and accessory products provides everything you need to successfully develop and perform your screens.



Figure 3. Lentiviral map. All constructs contain sequence-verified gRNA with expression driven by a U6 promoter, as well as a puromycin resistance gene driven by an EF-1a promoter. Puromycin resistance allows you to select for cells that were transduced with gRNA and therefore enrich the population for cells with edits to the target gene, which will ultimately produce stronger phenotypes in your screen.

Case study

Functional characterization studies of the human genome have delivered a tremendous amount of information using genome-based loss-of-function screening in diverse models. RNAi has been the predominant method for loss-of-function screening, but is limited by variable efficiency, frequent incompleteness of protein depletion, and confounding off-target effects. In addition, certain types of mammalian cells are difficult to transfect using lipid reagents or electroporation.

To circumvent these difficulties, lentiviral vectors are commonly used as another delivery method, as they can be easily titrated to manipulate transgene copy number and are stably maintained by integration into genomic DNA during subsequent cell replication. Figure 4 shows how a human kinase loss-of-function screen using our Invitrogen[™] CellSensor[™] NF-κB-*bla* ME-180 cell line easily enables identification of genomic targets associated with the NF-κB pathway. Α TNFα BLA-AM substrate Cell membrane Cytoplasmic esterases Cytoplasm CellSensor cell line 409 nm 409 nm **B**-lactamase TRADD BLA substrate IKK -TNFa +TNFα NF-_KB N β-lactamase С В -TNFa 4 Measurement (green to blue) HPRT ayed gRNA lentiviral particles 4 days post-infection Add TNFa +TNFa CellSensor NF-KB-bla ME-180 cell (stable Cas9-expressing clone)

Figure 4. A subset of kinases screened in CellSensor NF-κB-*bla* ME-180 cells with a ratiometric two-color reporter assay. (A) CellSensor cell lines use β-lactamase reporter technology to provide a rapid and sensitive method of analyzing activation of signal transduction pathways. (B) Overview of the screening workflow. (C) After stimulation with TNFa, the ratio of green to blue fluorescence decreased in unedited cells. Cells infected with lentiviral particles carrying gRNA that effectively disrupted the NF-κB pathway remained green, with a high ratio of green to blue fluorescence. Complete knockout of TNFR1 was observed in nearly 100% of the cells.

TRADD

TNFR1

Learn more at thermofisher.com/lentiarraycrispr

LentiPool CRISPR libraries

Our Invitrogen[™] LentiPool[™] CRISPR libraries are an affordable method to screen a large number of genes, as there is no high-throughput instrumentation required. The key difference between LentiPool and LentiArray CRISPR libraries is the format you receive them in. The pooled CRISPR library includes lentiviruses that encode gRNA to all your target genes, all in one tube, versus an array plate format where you receive one gene per well (Figure 5). All analysis can be performed using a next-generation sequencing (NGS) system. Our LentiPool CRISPR libraries are quality controlled by NGS to confirm gRNA and gene representation.

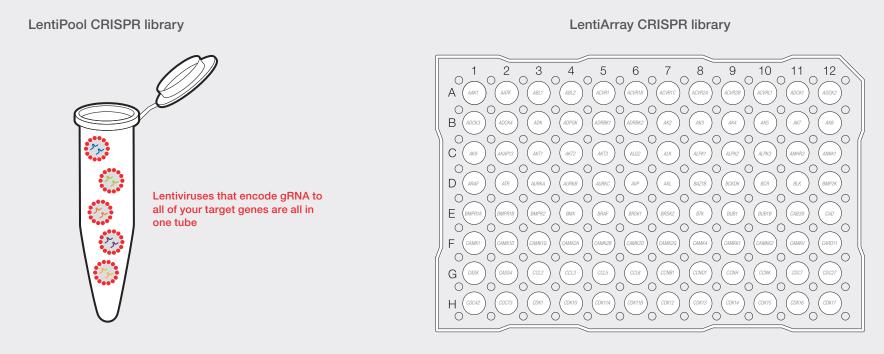


Figure 5. Comparison between LentiPool and LentiArray CRISPR libraries.

Lentivirus is organized to target one gene per well

For more information contact us at GEMServices@thermofisher.com

CRISPR library controls

Optimize your assay design with quality controls

High-quality controls are a cornerstone of successful screens. Our libraries provide a suite of controls to help you quickly develop your assay and define your hit criteria with confidence.

All of the content within the libraries is available as individual, ready-to-transfect gRNA to help you with rapid assay development and hit validation.

Negative controls

Negative controls are nontargeting gRNAs that don't recognize any sequence in the human genome. The negative controls provided are available in multiple package sizes for use in assay development and as on-plate controls when running your screens.

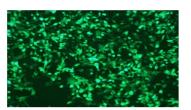
Positive controls

Positive controls are validated gRNA sequences that have demonstrated high-level editing efficiencies of up to 90% in several cell types. These controls are recommended for use in assay development to determine the conditions that provide maximum editing efficiency in your cell models.

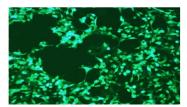
Individual gRNAs against specific genes are available to serve as biological controls for use in assay development and as on-plate positive controls when running your screens.

Delivery controls

Specific to LentiArray CRISPR libraries, we offer a set of delivery controls. These control lentiviruses are available as either negative or positive controls and also express GFP. The inclusion of the GFP marker provides a visual readout to aid in the rapid optimization of viral delivery conditions (Figure 6).



Negative control, scrambled gRNA sequence



Positive control, HPRT1 gene

Figure 6. Delivery controls. HT1080 cells were infected with a negative control lentivirus expressing a scrambled gRNA sequence, or a positive control lentivirus expressing a gRNA specific to the *HPRT1* gene. GFP is expressed from both lentivirus controls, which provides a visual readout of transduction efficiency and helps to determine multiplicity of infection (MOI) for the cell line being used in the screen.

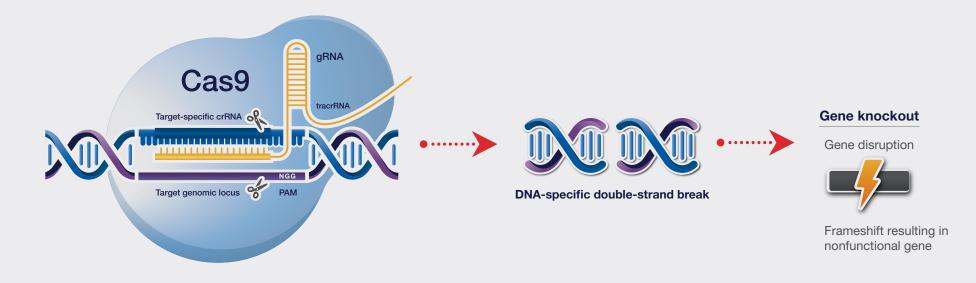


Figure 7. A CRISPR-Cas9 targeted double-strand break. Cleavage occurs on both strands, 3 bp upstream of the NGG in the protospacer adjacent motif (PAM) sequence on the 3' end of the target sequence.

Support resources New to CRISPR genome editing? Learn more at thermofisher.com/genomeedit101

High-efficiency gene knockout with the CRISPR-Cas9 system

The CRISPR-Cas9 system offers specific, complete, and permanent gene knockout, making it a potent tool for new discoveries about gene function. Derived from components of a simple bacterial immune system, this tool enables targeted gene cleavage and gene editing in a variety of eukaryotic cells. Since Cas9 nuclease specificity is determined by a gRNA sequence, editing can be directed to virtually any genomic locus by engineering the gRNA sequence and delivering it along with the Cas9 nuclease to your target cell.

Knockouts are created by the CRISPR-Cas9 system by introducing a double-strand break at a site within the target gene. This double-strand break is then repaired by nonhomologous end joining (NHEJ). The error-prone nature of NHEJ leads to the introduction of insertion or deletion (indel) mutations, resulting in a frameshift and ultimately a nonfunctional gene (Figure 7).

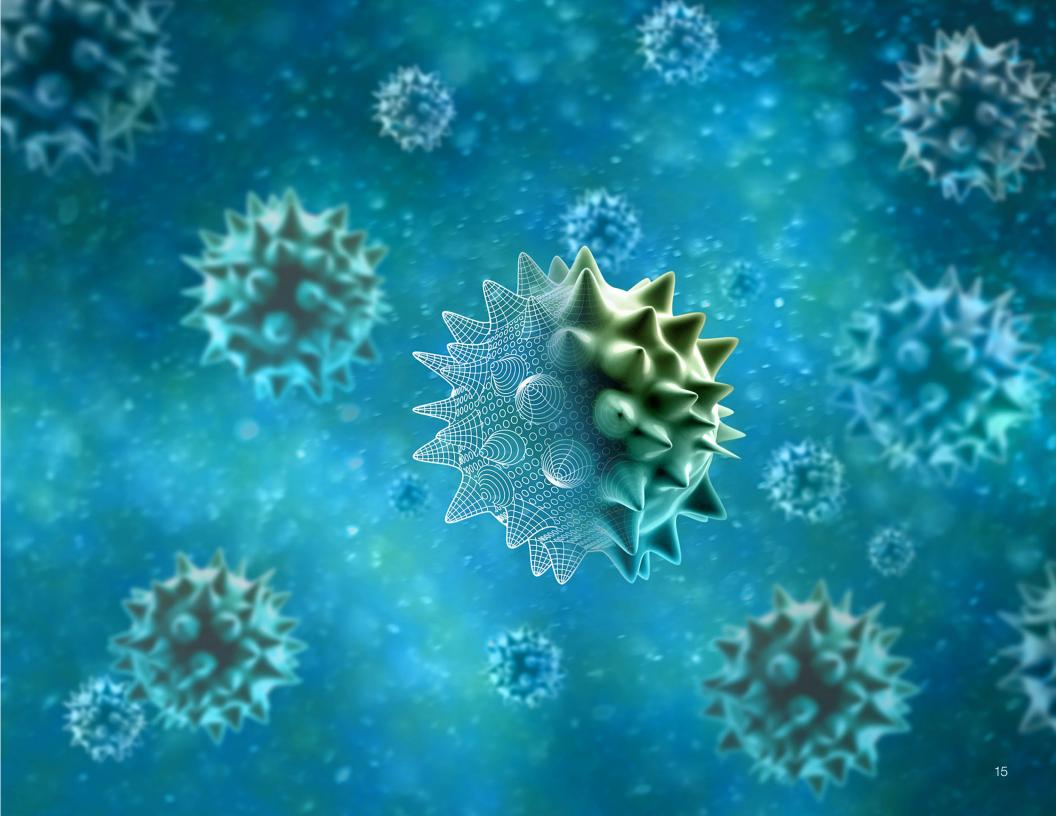


Figure 8. LentiArray Cas9 Lentivirus map. An EF-1a promoter drives expression of a Cas9 gene linked to the blasticidin resistance gene by a self-cleaving P2A peptide. Blasticidin resistance allows for the isolation of stable Cas9-expressing cells.

Invitrogen[™] LentiArray[™] Cas9 Lentivirus provides an efficient method to drive high-level Cas9 nuclease expression in a wide variety of cell types. The LentiArray Cas9 Lentivirus construct includes a human codon–optimized version of Cas9 with 2 nuclear localization signals (NLS, one on the C-terminus and one on the N-terminus) to facilitate efficient delivery into the nucleus (Figure 8). LentiArray Cas9 Lentivirus gives you the ability to create a stable pool of Cas9-expressing cells or isolate stable Cas9-expressing clones. LentiArray Cas9 Lentivirus can be used in screening applications with LentiArray CRISPR libraries. LentiArray Cas9 Lentivirus is provided as high-titer, ready-to-use lentiviral particles.

Cas9 formats available

We offer Cas9 nuclease in multiple formats. For screening applications, we offer Cas9 protein, Cas9 mRNA, and Cas9 lentiviral particles. These gene editing solutions are paired with optimal cell culture reagents, delivery methods, and analysis tools based on your application and cell type.



RNAi libraries

Transient knockdown of multiple gene targets

RNA interference (RNAi) is a proven and highly successful approach for lossof-function studies in virtually all eukaryotic organisms. We offer two types of small RNAs that function in RNAi: short interfering RNA (siRNA) and microRNA (miRNA) mimics and inhibitors. These products are available for *in vitro* and *in vivo* studies and include libraries for high-throughput applications. Your choice of tool depends on your model system, the length of knockdown time you require, and other experimental parameters.

siRNA libraries

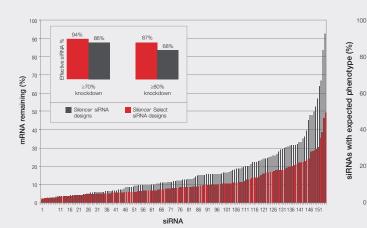
Ultrapotent siRNA for *in vitro* RNAi applications is the best way to effectively knock down gene expression to study protein function in a wide range of cell types. Traditional RNAi methods for gene knockdown in mammalian cells involve the use of synthetic RNA duplexes consisting of two unmodified 21-mer oligonucleotides annealed together to form siRNAs. Invitrogen[™] Silencer[™] Select siRNA products (Table 2) incorporate the latest improvements in siRNA design, off-target effect prediction algorithms, and chemical modifications to enhance strand bias.

* Guarantee details at thermofisher.com/sirnaguarantee

With our expanded line of Invitrogen[™] *Silencer[™]* and *Silencer[™]* Select siRNA libraries, you are empowered to expedite your functional genomics screen with precision and accuracy using an array of gene classes. Our premium siRNAs are designed to provide superior results for your loss-of-function studies, helping you to achieve your goals and confidently move to the next phase of your work (Figures 9–11).

We synthesize all siRNAs in-house to exacting quality standards. All *Silencer* and *Silencer* Select siRNA libraries, whether predefined or custom-made, are supplied with full siRNA sequence information and, when available, with siRNA validation data. *Silencer* Select siRNA consistently offers:

- Efficacy—guaranteed that with at least two of the siRNAs you will achieve >70% reduction in target mRNA levels, to help prevent additional siRNA testing, false negatives, and missed discoveries*
- **Specificity**—precise mRNA target recognition helps eliminate false positives, helping to reduce time-consuming and costly off-target validations
- Potency—100-fold more potent than other siRNAs, resulting in lower cost per experiment and low off-target effects



Improved design

Consistent phenotypes

80

60

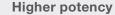
10

20

n = 28

Silencer Select siRNA

n = 43



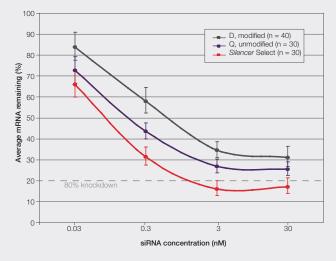


Figure 9. Silencer Select siRNA design algorithm significantly improves effective siRNA prediction accuracy. The Silencer Select siRNA algorithm was used to design 155 siRNAs to 40 different targets. These siRNAs were tested side by side with siRNAs designed using the previous algorithm, at 5 nM in HeLa cells. mRNA knockdown was measured 48 hours posttransfection via qRT-PCR using Applied Biosystems™ TaqMan[®] Gene Expression Assays. Results are expressed as percentage of mRNA remaining compared to cells treated with Invitrogen[™] Silencer[™] Select Negative Control No. 1 siRNA. The inset shows the percentages of siRNAs that elicited ≥70% and ≥80% mRNA knockdown.

Figure 10. Silencer Select siRNAs elicit the expected phenotype at a higher rate than other siRNAs. siRNAs to 7 gene targets with well-understood RNAi-induced phenotypes were individually transfected at 3 nM and phenotypes measured 48 hours later. Each bar represents the percentage of siRNAs that gave the expected, silenced phenotype. siRNAs to BUB1B, AURKB, WEE1, and PLK1 were assessed using a multiparametric cell growth/apoptosis assay in U2OS human osteosarcoma cells. siRNAs to HMGCR, LDLR, and FDFT1 were assessed using an LDL uptake assay in HUH7 human hepatoma cells.

Supplier D, modified siRNA Supplier Q, unmodified siRNA Silencer siRNA

n = 21

n = 34

Figure 11. Silencer Select siRNAs provide up to 100x higher potency compared to other siRNAs. Silencer Select siRNAs and siRNAs from two other suppliers to the same 10 gene targets were individually transfected into HeLa cells in triplicate at the indicated siRNA concentrations. mRNA knockdown levels were tested 48 hours later by gRT-PCR. Average percentage of mRNA remaining is shown for each set of siRNAs.

| Table 2. siRNA | product | selection | guide. |
|----------------|----------------|-----------|--------|
|----------------|----------------|-----------|--------|

| | Silencer siRNA | Stealth RNAi [™] siRNA | Silencer Select siRNA |
|----------------------------|---|--|--|
| | Good knockdown, cost-effective siRNA | Excellent knockdown, low off-target effects | Best knockdown, lowest off-target effects |
| Potency | 20 nM recommended concentration | 20 nM recommended concentration | 5 nM recommended concentration |
| Efficacy (>70% knockdown)* | 2 of 3 siRNAs guaranteed | 2 of 3 siRNAs guaranteed | 2 of 2 siRNAs guaranteed |
| Specificity | Moderate | High | Highest |
| Coverage | Coding RNA | Coding RNA | Coding and noncoding RNA |
| Target species | Human, mouse, rat (use custom tool for other species) | Human, mouse, rat (use custom tool for other species) | Human, mouse, rat (use custom tool for other species) |
| Recommended product | <i>Silencer</i> Pre-designed siRNA, 5 nmol (Cat. No. AM16708) | Stealth RNAi siRNA, tube (Cat. No. 1299001) | <i>Silencer</i> Select Pre-designed siRNA, 5 nmol (Cat. No. 4390771) |

* Guarantee details at thermofisher.com/sirnaguarantee

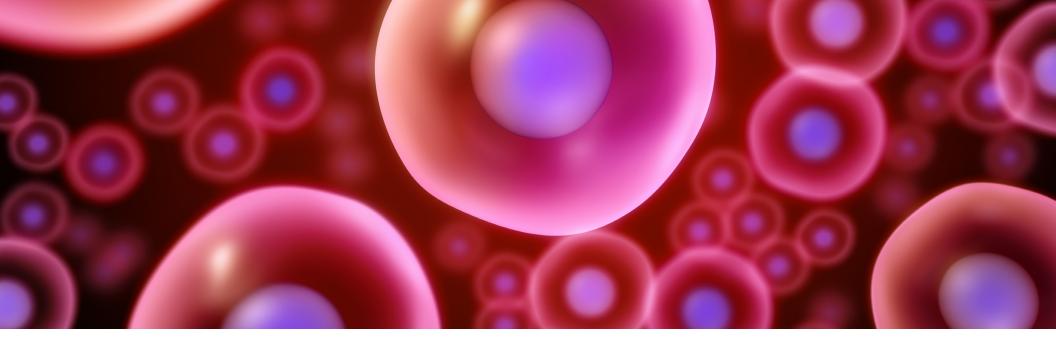
Request more information at **thermofisher.com/sirnaquote** To learn more or place an order, go to **thermofisher.com/sirna**

siRNA controls

Proper controls are essential to help ensure success in every RNAi experiment. The number and types of controls chosen depend on the ultimate research goal. We offer positive and negative controls as well as Invitrogen[®] GeneArt[®] gene synthesis for siRNA-resistant genes that can be used in RNAi rescue experiments.

To learn more, go to thermofisher.com/sirnacontrols





miRNA mimics and inhibitors

For artificial regulation of target mRNA translation, Invitrogen[™] *mir*Vana[™] miRNA mimics and inhibitors are chemically modified, synthetic nucleic acids designed to either mimic mature miRNAs or to bind to and inhibit endogenous miRNAs. These products provide a means to functionally study the role of specific miRNAs within cellular systems or to validate the role of miRNAs in regulating target genes. *mir*Vana miRNA mimics and inhibitors are:

- **Versatile**—ready-to-use *in vitro* and *in vivo*, allowing consistency and specificity throughout your entire study (Figures 13 and 14)
- **Highly potent**—lack toxicity and immunostimulation essential to *in vivo* studies
- Available as high-throughput screening libraries—most up-to-date miRBase sequence content
- Current-updated frequently using miRBase database content

Resources

- RNAi Handbook-go to thermofisher.com/rnai
- Genome Editing Selection Guide—go to thermofisher.com/geneedit and select "Genome Editing Resource Center"
- Transfection Protocol Calculator-go to thermofisher.com/transfection
- Publications-go to thermofisher.com/rnaipublications



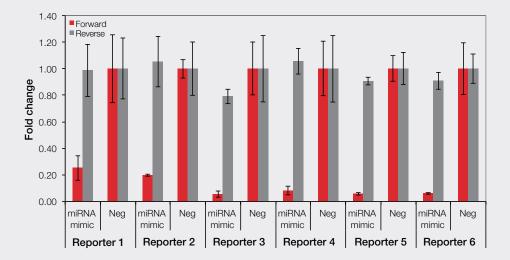


Figure 12. Superior miRNA specificity. The guide strand of *mir*Vana miRNA mimics is highly potent and the star strand is inactivated. The key advantage of *mir*Vana miRNA mimics is inactivation of the star strand. miRNA mimics, like natural microRNAs, have 2 strands. The guide strand is functional and used by argonaute protein to target miRNAs. The star, or passenger strand, is nonfunctional and is normally cleaved and expelled from the silencing complex. Most scientists want to analyze one strand of miRNA at a time while the other strand remains totally inactive. *mir*Vana miRNA mimics achieve this specificity. For this assay we measured activity from both strands of miRNA mimics. One reporter had a target in the forward orientation to measure activity of the guide miRNA strand. Another reporter had the target cloned in reverse orientation to test activity of the star strand of the miRNA mimic. For all 6 sequences, activity of the guide strand is high (resulting in 75–90% lower expression of the reporter), and activity of the star strand is minimal (similar to negative control).

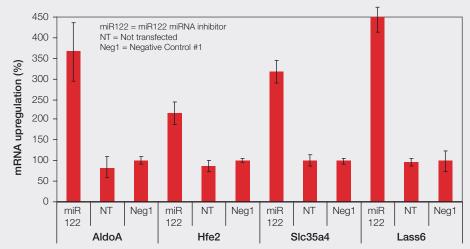


Figure 13. *mirVana miRNA inhibitors effectively suppress miRNA in vivo.* Invitrogen[™] *mirVana*[™] miR122 inhibitor or Negative Control #1 (Neg1) were complexed with Invitrogen[™] Invivofectamine[™] 2.0 Reagent and delivered to BALB/c mouse liver via tail vein injection on three consecutive days at a dose of 7 mg/kg body weight. Expression of four mRNA targets (AldoA, Hfe2, Slc35a4, and Lass6), natural targets of miR122, were measured using Applied Biosystems[™] TaqMan[®] MicroRNA Assays in transfected livers of mice injected with miR122 inhibitor or Negative Control #1 and livers of mice that were not transfected (NT). The results indicate that the inhibitor is efficiently delivered to the liver with Invivofectamine 2.0 Reagent and inactivate miR122, leading to upregulation of genes naturally suppressed by miR122.

miRNA libraries

Complete Invitrogen[™] *mir*Vana[™] libraries containing mimics and inhibitors for human, mouse, and rat miRNA are available.

For information on all our predefined and custom miRNAs libraries, contact us at **RNAiLibraries@lifetech.com**

To learn more or place an order, go to thermofisher.com/mirna

Custom library services

We collaborate with you as partners, from start to finish, to help accelerate your discovery

You're not on this journey alone. Our experienced scientists including technical specialists, project support specialists, and other professionals who appreciate your challenges can help you find answers efficiently and accurately. Whether you're validating an assay, setting up your experiment, purchasing supplies, or verifying compatibility, our team is here to assist you. Especially at a time when you're constantly challenged to do more with less, a problem with an assay is the last thing your lab needs.

Through a consultative approach to all services offered, our RNAi and gene editing team can work with you to design and implement the solutions that fit. We offer custom pooled libraries, custom libraries with genes selected from our pooled libraries, and more. From smaller validation projects and consulting engagements to complete turnkey solutions on a regional or nationwide scale, we can help you achieve your goals.



Questions? Ready to get started?

Contact our dedicated technical support team today at **GEMServices@thermofisher.com**

Transfection technologies

Transfection selection guide

Transfection is the process by which nucleic acids are introduced into eukaryotic cells. Transfection techniques vary widely, including lipid nanoparticle–mediated transfection and physical methods such as electroporation. Our Invitrogen[™] Lipofectamine[™] family of reagents paired with the Invitrogen[™] Neon[™] Transfection System provide complete delivery solutions for your genome editing needs. We have optimized protocols to achieve high cleavage efficiency and ease of delivery. An overview of our most effective transfection products is shown in Table 3.

Table 3. Transfection product selection guide.

| | siRNA, miRNA | mRNA | Protein | Lentivirus production |
|---|--------------|------|---------|-----------------------|
| Lipofectamine CRISPRMAX Cas9 Transfection Reagent | | | V | |
| Lipofectamine 3000 Transfection Reagent* | | | | V |
| Lipofectamine RNAiMAX Transfection Reagent | V | | | |
| Neon Transfection System | V | v | V | |
| Lipofectamine MessengerMAX Transfection Reagent | | v | | |

* Invitrogen[™] Lipofectamine[™] 3000 Transfection Reagent can be used to produce lentivirus.

Choose the solution that's right for you at thermofisher.com/transfection

Lipofectamine CRISPRMAX Cas9 Transfection Reagent

First transfection reagent optimized for CRISPR-Cas9 protein delivery

With Invitrogen[™] Lipofectamine[™] CRISPRMAX[™] Cas9 Transfection Reagent, it is now possible to use a lipid-based reagent to deliver CRISPR-Cas9 protein complexes (Table 4). Lipofectamine CRISPRMAX reagent is the first lipid-based transfection reagent for optimized delivery of CRISPR-Cas9 protein complexes, providing up to 85% cleavage efficiency when combined with Invitrogen[™] GeneArt[™] Platinum[™] Cas9 Nuclease. It is also cost-effective and gentle on cells. Deliver our superior GeneArt Platinum Cas9 Nuclease with a reagent that provides:

- **Demonstrated cleavage efficiency**—tested in over 20 cell types, including iPSCs, mESCs, N2a, CHO, A549, HCT116, HeLa, HEKa, HEK 293, and several others
- Low cell toxicity-fewer cells needed to initiate your experiment
- Cost savings—whether cost per reaction or initial investment
- Easy scalability—an ideal delivery solution for high-throughput experiments

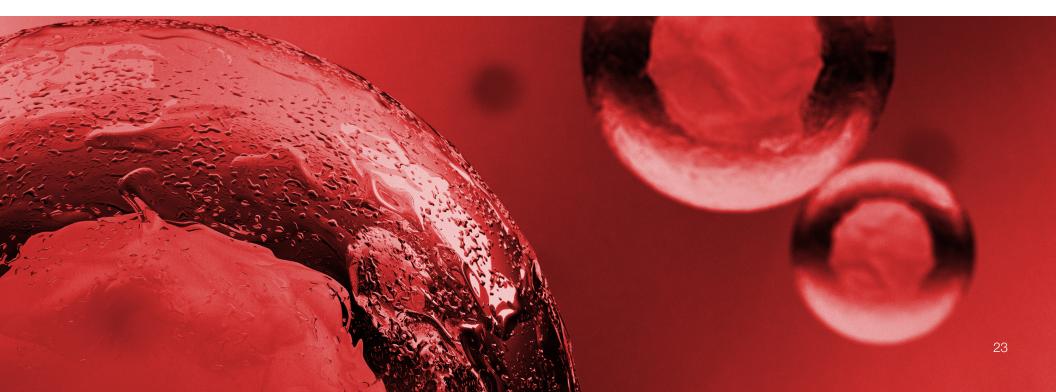


Table 4. Summary of CRISPR-Cas9 delivery methods.

| | CRISPR protein | CRISPR mRNA | CRISPR plasmid | CRISPR libraries |
|--|--|--|--|---|
| | Cas9 protein | IVT gRNA (from GeneArt CRISPR T7 Strings DNA Fragment) + + Cas9 mRNA | GenArt CRISPR vector with OFP reporter | Lentiviral gRNA |
| | GeneArt Platinum | GeneArt CRISPR | GeneArt CRISPR | CRISPR LentiArray |
| | Cas9 Nuclease | Nuclease mRNA | Nuclease Vector | Libraries |
| gRNA design | Use the Ger | - | ol for optimal design and minimal off-t r.com/crisprdesign | arget effects. |
| gRNA synthesis | GeneArt Precision gRNA Synthesis Kit | GeneArt Precision gRNA Synthesis Kit | DNA oligos cloned into plasmid | |
| Reporter-based enrichment | GeneArt Genomic Cleavage Selection Kit (sold separately) | GeneArt Genomic Cleavage Selection Kit (sold separately) | All-in-one expression plasmid (included) | |
| No promoter constraint | v | V | Cytomegalovirus (CMV) promoter | |
| Ready to use | v | ✓ | Required cloning step | Ready-to-use lentiviral particles |
| No random integration concern | v | v | No, could be a concern | Stable expression of Cas9; lentivirus may randomly integrate |
| Controlled dosage | V | ✓ | | |
| Fast turnover | 4 | ✓ | | |
| Microinjection-ready | v | ✓ | Larger payload | |
| Multiplexing and screening capable | v | v | Larger payload | High-throughput screening |
| Ready-to-act, stable ribonucleoprotein (RNP) complex | v | | | |
| Modification options | Knockout and knock-in | Knockout and knock-in | Knockout and knock-in | Loss-of-function screening |
| Delivery method* | Lipofectamine CRISPRMAX reagent | Lipofectamine MessengerMAX reagent | Lipofectamine 3000 reagent | |

* For the most efficient transfection of primary cells, stem cells, and difficult-to-transfect cells, use the Neon Transfection System.

Learn more or see what cell types we've tested at thermofisher.com/crisprtransfection

Lipofectamine RNAiMAX Transfection Reagent

Invitrogen[™] Lipofectamine[™] RNAiMAX[™] Transfection Reagent offers an advanced, efficient solution for siRNA or miRNA delivery. It can also be used for cotransfection of gRNA and Cas9 mRNA. No other transfection reagent for RNAi experiments provides such easy and efficient delivery into a broad spectrum of cell types including primary cells, stem cells, and other hard-to-transfect cell types (Figure 14).

Superior transfection efficiency at low siRNA concentrations

When it comes to achieving effective gene knockdown, Lipofectamine RNAiMAX reagent outperforms other siRNA transfection reagents. High knockdown levels of target genes can be achieved with as little as 1 nM siRNA.

Low cytotoxicity profile for easy optimization

Lipofectamine RNAiMAX reagent enables maximal knockdown and excellent cell viability across a 10-fold concentration range of the reagent. This makes Lipofectamine RNAiMAX reagent easy to optimize for the lowest siRNA concentration that can be used while reducing cytotoxicity in your experimental system. Transfection-mediated cytotoxicity can mask the true phenotype of a target gene being studied, so minimizing the amount of reagent during transfection is a critical factor for successful RNAi experiments.

Support resources

View transfection protocols at thermofisher.com/transfectionprotocols

Download your copy of our transfection and genome engineering handbook at **thermofisher.com/transfectionhandbook**

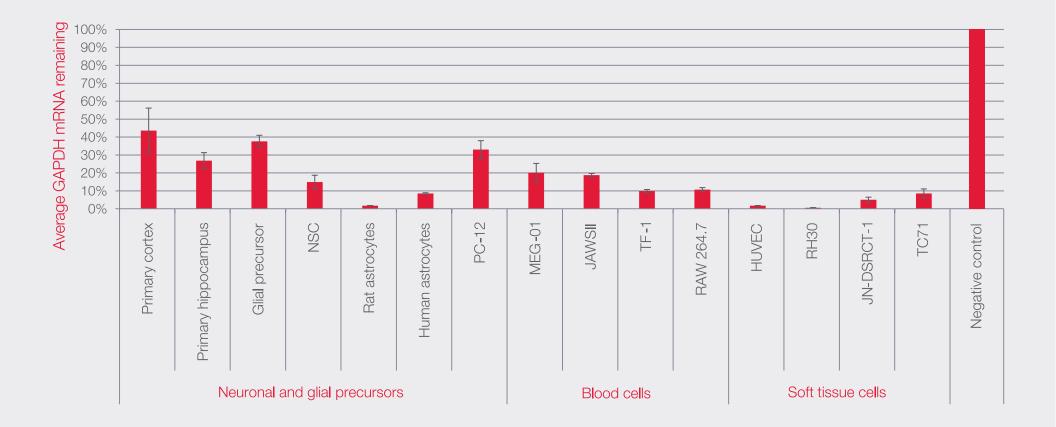


Figure 14. Invitrogen *Silencer* **Select siRNA transfection with Lipofectamine RNAiMAX reagent.** Cells were transfected with Lipofectamine RNAiMAX reagent complexed with *Silencer* Select siRNA at 30 nM/well. Knockdown of GAPDH mRNA was assessed by qPCR. The cell density per well and amount of Lipofectamine RNAiMAX reagent used per well were as follows: rat primary cortex cells: 10⁴ cells/well, 0.6 µL; rat primary hippocampus cells: 10⁴ cells/well, 0.6 µL; rat glial precursor cells: 8 x 10³ cells/well, 0.4 µL; neuroblastoma-spinal cord (NSC) cells: 1.2 x 10⁴ cells/well, 0.3 µL; rat astrocyte cells: 1.2 x 10⁴ cells/well, 0.3 µL; PC-12 cells (derived from a pheochromocytoma of the rat adrenal medulla): 8 x 10³ cells/well, 0.3 µL; megakaryoblastic leukemia cells (MEG-01): 2 x 10⁴ cells/well, 0.3 µL; murine dendritic cells (JAWSII): 1.2 x 10⁴ cells/well, 0.3 µL; human erythroleukemia cells (TF-1): 10⁴ cells/well, 0.3 µL; RAW 264.7 macrophages: 10⁴ cells/well, 0.3 µL; human umbilical vein endothelial cells (HUVEC): 8 x 10³ cells/well, 0.3 µL; rhabdomyosarcoma cells (RH30): 8 x 10³ cells/well, 0.3 µL; human desmoplastic small round cell tumor cells (JN-DSRCT-1): 8 x 10³ cells/well, 0.3 µL; human Ewing's sarcoma cells (TC71): 1.6 x 10⁴ cells/well, 0.3 µL.

To learn more or place an order, go to thermofisher.com/rnaimax

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High-content analysis

CRISPR analysis steps using HCA

- Process development
- Rapid phenotyping
- Off-target screening

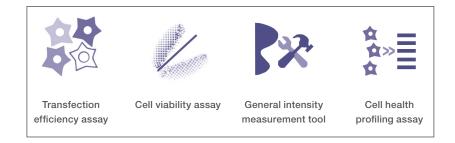
When screening a CRISPR library involves large cell numbers and complex phenotypic responses, HCA is an obvious choice. HCA provides automation throughout the process, and the imaging format allows for quantitative morphological analysis.

Process optimization

Cell analysis using fluorescent protein reporters of transfection or dsDNA cleavage is routine in automated workflows generally run at high throughput in HCA. Cell health assays prior to editing can be a stringent QC step to avoid wasting time and reagents, and after editing they are critical metrics for process optimization.

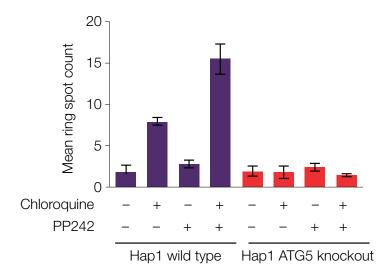


Thermo Scientific[™] CellInsight[™] CX7 High Content Analysis Platform



Cell phenotyping

The CRISPR-Cas9 system is routinely used for knockout, knock-in, and modulation of gene expression. HCA provides automated analysis with quantitative rigor to directly analyze changes in protein expression levels, compartmentalization, or cell morphology (Figure 15).



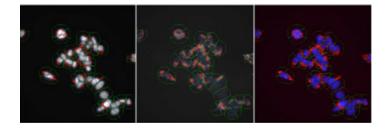


Figure 15. Automated quantitation of autophagy using HCS. The CellInsight CX7 High Content Analysis (HCA) Platform was used to identify and count LC3B granules in CRISPR-edited cells. Nuclei (blue overlay), cell perimeter (green outline), and LC3B granules (red overlay) are shown.

Off-target analysis

HCA allows you to track your targeted protein and also monitor its impact on cell health and behavior. Invitrogen[™] reagents and the CellInsight CX5 High Content Analysis (HCA) Platform makes analysis of multiple cell health parameters routine (Figure 16).

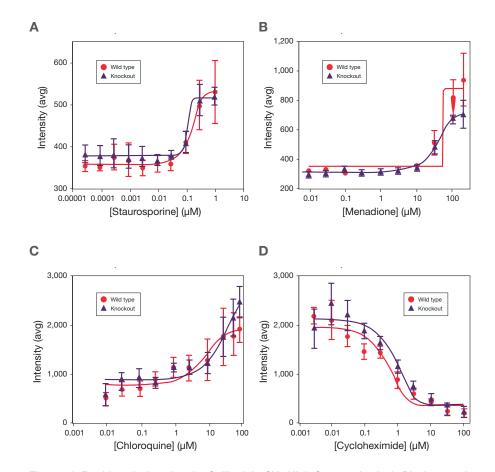


Figure 16. Rapid analysis using the CellInsight CX5 High Content Analysis Platform and Invitrogen[™] reagents. Wild type and CRISPR-edited cells were compared for (A) apoptosis, using CellEvent[™] Caspase-3/7 Green Detection Reagent, (B) oxidative stress, using CellROX[™] reagents, (C) protein degradation, using LysoTracker[™] reagents, and (D) protein synthesis, using the Click-iT[™] Plus OPP kit.

Find out more at thermofisher.com/detectcrispr

Ordering information

| | Product and Cat. No. | | | No. |
|--|---------------------------------------|--|--|--|
| Product | <i>Silencer</i> siRNA (unmodified) | <i>Silencer</i> Select siRNA (chemically modified) | LentiArray CRISPR (ready to use) | LentiArray CRISPR (glycerol stock) |
| Screening libraries | | | | |
| Human Whole Genome Library | NA | NA | A31949 | A32185 |
| Human Druggable Genome Library | A30077 | NA | A31948 | A32184 |
| Human Apoptosis Library | A30095 | A30095 | A31940 | A32176 |
| Human Cancer Biology Library | A30143 | A30142 | A31933 | A32169 |
| Human Cell Cycle Regulation Library | A30137 | A30136 | A31936 | A32172 |
| Human Cell Surface Library | A30145 | A30144 | A31943 | A32179 |
| Human DNA Damage Response Library | A30090 | A30089 | A31946 | A32182 |
| Human Drug Transporter Library | A30135 | A30133 | A31941 | A32177 |
| Human Epigenetics Library | A30086 | A30085 | A31934 | A32170 |
| Human GPCR Library | A30082 | NA | A31947 | A32183 |
| Human Ion Channel Library | A30084 | NA | A31942 | A32178 |
| Human Kinase Library | A30079 | NA | A31931 | A32167 |
| Human Membrane Trafficking Library | A30139 | A30138 | A31937 | A32173 |
| Human Nuclear Hormone Receptor Library | A30083 | NA | A31939 | A32175 |
| Human Phosphatase Library | A30080 | NA | A31932 | A32168 |
| Human Protease Library | A30081 | NA | A31944 | A32180 |
| Human Transcription Factor Library | NA | A30087 | A31938 | A32174 |
| Human Tumor Suppressor Library | A30147 | A30146 | A31945 | A32181 |
| Human Ubiquitin Library | A30141 | A30140 | A31935 | A32171 |

| Product | Quantity | Cat. No. |
|--|----------|----------|
| CRISPR library controls | | |
| LentiArray CRISPR Positive Control Lentivirus, human HPRT | 100 µL | A32056 |
| LentiArray CRISPR Positive Control Lentivirus, human HPRT | 1 mL | A32829 |
| LentiArray CRISPR Positive Control Lentivirus, human HPRT with GFP | 100 µL | A32060 |
| LentiArray CRISPR Positive Control Lentivirus, human HPRT with GFP | 1 mL | A32830 |
| LentiArray CRISPR Negative Control Lentivirus, human non-targeting | 100 µL | A32062 |
| LentiArray CRISPR Negative Control Lentivirus, human non-targeting | 1 mL | A32327 |
| LentiArray gRNA | Custom | A32042 |
| LentiArray Custom CRISPR Plate | Custom | A32045 |

Ordering information (continued)

| Product | Quantity | Cat. No. |
|---|----------|-----------|
| Cas9 formats | | |
| LentiArray Cas9 Lentivirus | 100 µL | A32064 |
| LentiArray Cas9 Lentivirus | 1 mL | A32069 |
| GeneArt Platinum Cas9 Nuclease (1 µg/µL) | 10 µg | B25642 |
| GeneArt Platinum Cas9 Nuclease (1 µg/µL) | 25 µg | B25640 |
| GeneArt Platinum Cas9 Nuclease (3 µg/µL) | 75 µg | B25641 |
| GeneArt CRISPR Nuclease mRNA | 15 µg | A29378 |
| Related products | | |
| GeneArt Genomic Cleavage Detection Kit | 20 rxns | A24372 |
| THE RNA Storage Solution | 50 mL | AM7001 |
| Lipofectamine CRISPRMAX Cas9 Transfection Reagent | 10 rxns | CMAX00001 |
| Lipofectamine CRISPRMAX Cas9 Transfection Reagent | 25 rxns | CMAX00003 |
| Lipofectamine CRISPRMAX Cas9 Transfection Reagent | 75 rxns | CMAX00008 |
| Lipofectamine CRISPRMAX Cas9 Transfection Reagent | 150 rxns | CMAX00015 |
| Lipofectamine RNAiMAX Transfection Reagent | 1.5 mL | 13778150 |
| Lipofectamine MessengerMAX Transfection Reagent | 1.5 mL | LMRNA015 |
| Neon Transfection System Starter Pack | 1 pack | MPK5000S |

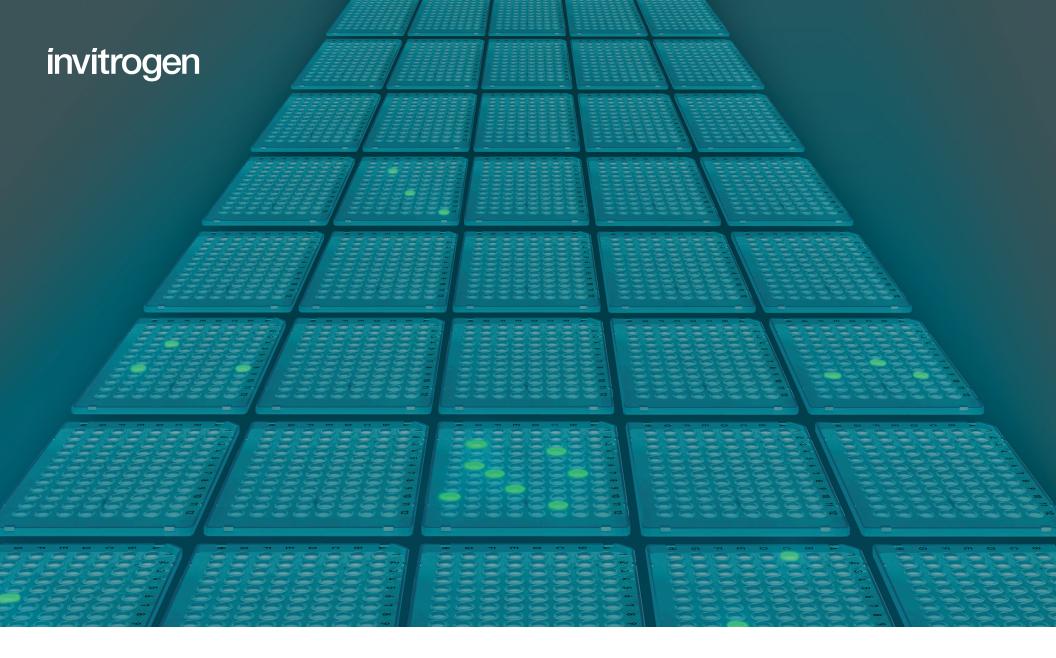
| Product | Quantity | Cat. No. | | | |
|---|--|----------|--|--|--|
| mirVana miRNA mimics and inhibitors*-next-generation miRNA chemistries for lowest off-target ef | mirVana miRNA mimics and inhibitors*—next-generation miRNA chemistries for lowest off-target effects and highest potency | | | | |
| mirVana Predesigned miRNA Mimic | 5 nmol | 4464066 | | | |
| mirVana Predesigned miRNA Mimic, in vivo use | 250 nmol | 4464070 | | | |
| mirVana Predesigned miRNA Inhibitor | 5 nmol | 4464084 | | | |
| mirVana Predesigned miRNA Inhibitor, in vivo use | 250 nmol | 4464088 | | | |
| mirVana miRNA Mimic, Negative Control #1 | 5 nmol | 4464058 | | | |
| mirVana miRNA Inhibitor, Negative Control #1 | 5 nmol | 4464076 | | | |
| mirVana miRNA Mimic, miR-1 Positive Control | 5 nmol | 4464062 | | | |
| mirVana miRNA Inhibitor, let-7c Positive Control | 5 nmol | 4464080 | | | |
| mirVana Custom miRNA Mimic | 5 nmol | 4464068 | | | |
| mirVana Custom miRNA Mimic, in vivo use | 250 nmol | 4464071 | | | |
| mirVana Custom miRNA Inhibitor | 5 nmol | 4464086 | | | |
| mirVana Custom miRNA Inhibitor, in vivo use | 250 nmol | 4464089 | | | |
| Pre-miR precursors and anti-miR inhibitors—trusted, affordable miRNA | | | | | |
| Predesigned | | | | | |
| Anti-miR miRNA Inhibitor | 5 nmol | AM17000 | | | |
| Pre-miR miRNA Precursor | 5 nmol | AM17100 | | | |

Ordering information (continued)

| Product | Cat. No. |
|---|----------|
| Positive controls | |
| Silencer Select GAPDH Positive Control siRNA | 4390850 |
| Silencer GAPDH siRNA (human, mouse, rat) | AM4624 |
| Silencer GAPDH siRNA (human) | AM4633 |
| Silencer Select MALAT1 Positive Control siRNA | 4455877 |
| Silencer Cy®3-labeled GAPDH siRNA (human, mouse, rat) | AM4649 |
| Silencer GFP (eGFP) siRNA | AM4626 |
| Silencer Firefly Luciferase (GL2 + GL3) siRNA | AM4629 |
| mirVana miRNA Mimic, miR-1 Positive Control | 4464062 |
| mirVana miRNA Inhibitor, let-7c Positive Control | 4464080 |
| Negative controls | |
| Silencer Select Negative Control No. 1 siRNA | 4390843 |
| Silencer Select Negative Control No. 2 siRNA | 4390846 |
| Silencer Negative Control No. 1 siRNA | AM4611 |
| Silencer Negative Control No. 2 siRNA | AM4613 |
| mirVana miRNA Inhibitor, Negative Control #1 | 4464077 |
| mirVana miRNA Mimic, Negative Control #1 | 4464058 |

* Covers human, mouse, rat, and other species.





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