

# Optimized crRNA and tracrRNA for Your In Vitro and In Vivo CRISPR Experiments

Ready-to-use crRNA and tracrRNA – no costly cloning necessary  
Increased half-life due to proprietary chemical modifications  
Ideal for screening – available in various quantities

## Background Information

CRISPR/Cas has advanced to a standard technology in modern molecular biology and biotech labs in a very short time since its first publication in 2012.

Briefly summarized: An endonuclease (Cas) is directed to a target DNA

sequence using guide RNA(s). The nuclease initiates a double strand break which leads to a knockout of the gene of interest. Basically every target can be made accessible for knocking out by this method. The easy programmability, the simplicity of the proto-

cols and the stability of the system are clear advantages of CRISPR/Cas. Common experiments involving CRISPR/Cas are performed by cloning the Cas nuclease and the sgRNA into a plasmid for transcription and expression – this is no longer needed!

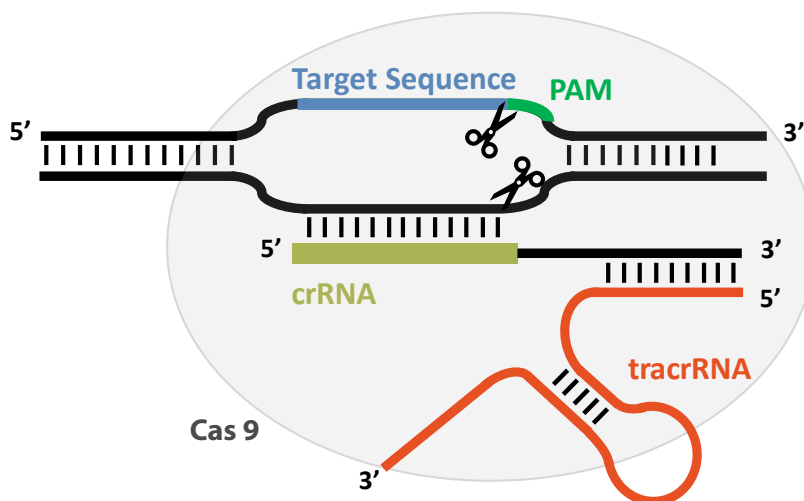
## Microsynth's Optimized Guide RNA's for Your CRISPR/Cas Experiments

To further simplify your lab work, Microsynth offers you two **ready-to-use guide RNAs** (crRNA and tracrRNA) which – together with the Cas endonuclease – can directly be used in your experiment **without any cloning steps**.

The two guide RNAs have been opti-

mized by including **various types of chemical modifications** resulting in an enhanced stability against nucleases. Compared with standard RNA, this **prolongs the half-life** of the guide RNAs in the cell and **increases the success** of your CRISPR/Cas experiment. High turnaround for **multiple**

**target sites** since you only need to design a new crRNA while you use the same tracrRNA. These advantages as well as the availability of various nmol quantities make the Microsynth guide RNA's **ideal for screening**.



**Figure 1:** The crRNA consists of a 19-21 nt sequence (binding to the complementary DNA strand) as well as a tracrRNA fusion domain.

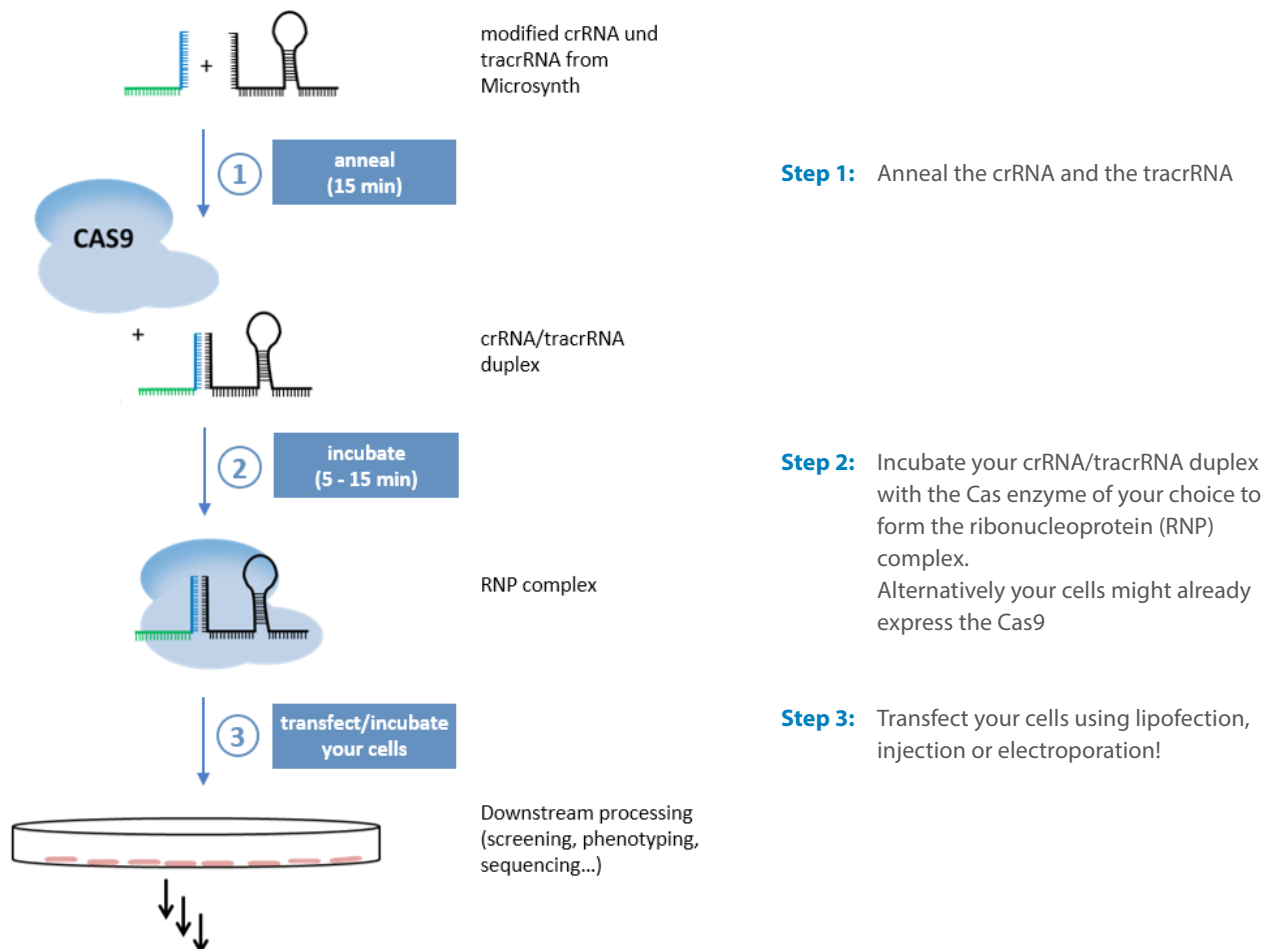
When complexed with tracrRNA and Cas9 protein, a double strand break will be created at a locus complementary to the 19-21 nt guide RNA sequence, 3-4 bp upstream from a PAM sequence (5'-NGG-3').

## Our Product

crRNA	tracrRNA
Chemically modified for increased half-life	Chemically modified for increased half-life
2 nmol minimal yield guaranteed	2 nmol or 5 nmol aliquots
Larger amounts on request	Larger amounts on request
19 - 21 nt target sequence + 3 PAM sequence	Universally usable for all your experiments
PAGE purified for highest purity and efficacy	PAGE purified for highest purity and efficacy
100% quality control	100% quality control

## Experimental Procedure

Simple 3 Step protocol to your knock out or homology-directed repair (HDR) clone.



### How to Order?

Using your gene maps and online available software, you choose your target site of your gene of interest.

Please send your 19-21 bp target sequence from 5' to 3' plus the 3 nt PAM sequence to [oligo.support@microsynth.ch](mailto:oligo.support@microsynth.ch)

### Need More Information?

Call us at +41 71 722 83 33 or

Email us at [oligo.support@microsynth.ch](mailto:oligo.support@microsynth.ch)