

# Reduced off-target effects with delivery of active Cas9 protein complexed with sgRNA using gesicle technology

## Guide-it CRISPR/Cas9 Gesicle Production System

- Delivery of active ribonucleoprotein complexes via gesicles results in reduced off-target effects
- Confirmation of reduction in off-target effects is seen in Sanger sequencing results

## Introduction

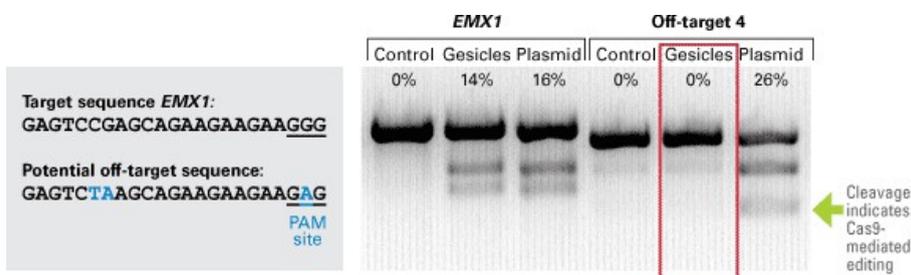
CRISPR/Cas9 gene editing is an RNA-programmable system which has democratized targeted genome modification by virtue of its simplicity and versatility. With this technology, gene editing is mediated by Cas9 nuclease and a single-guide RNA (sgRNA) which directs the Cas9 to a specific genomic locus. However, the utility of any genome modification system—for either basic research or therapeutic development—depends on its specificity (Sternberg and Doudna 2015). Early reports have warned of the frequent off-target effects of the CRISPR/Cas9 system (Fu et al. 2013; Hsu et al. 2013; Mali et al. 2013; Pattanayak et al. 2013), demonstrating the need for developing new methods that reduce these events.

The Guide-it CRISPR/Cas9 Gesicle Production System is a novel methodology that uses cell-derived nanovesicles, called **gesicles**, for co-delivery of active Cas9 protein complexed with a gene-specific sgRNA. Gesicles are loaded with Cas9-sgRNA ribonucleoprotein (RNP) complexes and can be added directly to target cells for CRISPR/Cas9 gene editing. Delivery of active Cas9 protein means no Cas9 coding gene is present in target cells, thus eliminating the problem of persistent and elevated Cas9 levels, which is common to plasmid-based delivery methods.

## Results

### CRISPR/Cas9 gesicles reduce off-target effects compared to plasmid transfection

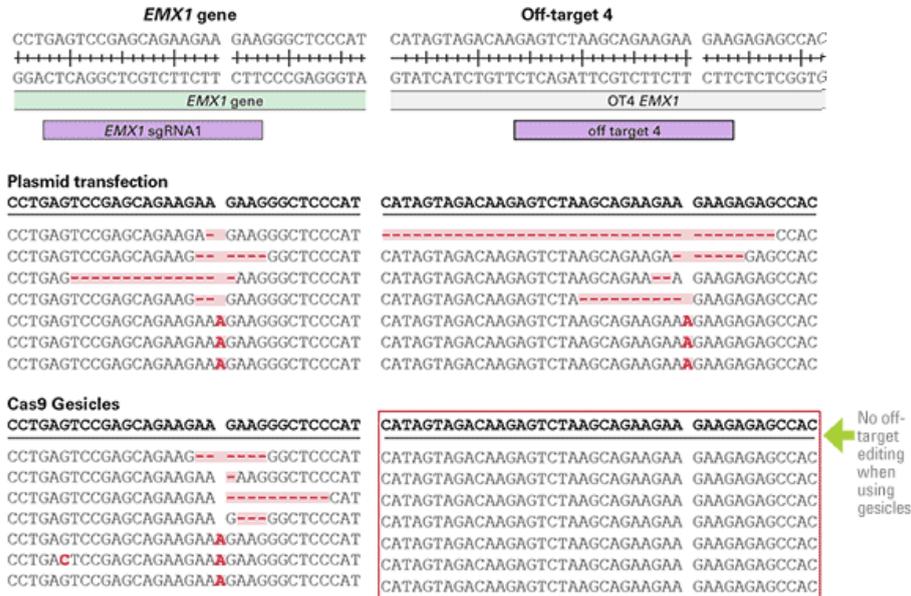
In a side-by-side comparison of Cas9-sgRNA delivery methods, the *EMX1* gene was edited in HEK 293T cells either by treatment with gesicles loaded with Cas9-sgRNA RNP complexes or by transfection with plasmids encoding Cas9 and a gene-specific sgRNA (Figure 1). Following CRISPR/Cas9 modification, the *EMX1* locus and a potential off-target locus (off-target 4) were PCR-amplified from crude cell extracts. The presence of indels was detected using Guide-it Resolvase (a mismatch-specific nuclease included in the Guide-it Mutation Detection Kit), followed by agarose gel electrophoresis. Densitometry (Cong et al. 2013) showed roughly equivalent indel formation at the *EMX1* target locus between the two methods. As expected, plasmid transfection resulted in significant indel formation at the off-target 4 locus, while gesicle delivery resulted in no observable indel formation at this locus (Figure 1).



**Figure 1. Reduced off-target effects with CRISPR/Cas9 gesicles.** HEK 293T cells were either treated with gesicles loaded with Cas9-sgRNA RNP complexes or transfected with plasmids encoding Cas9 and an sgRNA against *EMX1*. After 72 hr, the *EMX1* gene and a potential off-target locus (off-target 4) were amplified from the treated cells by direct PCR. Using the Guide-it Mutation Detection Kit, the amplicons were melted and rehybridized, and mismatched targets were cleaved using Guide-it Resolvase. A control sample that lacked Guide-it Resolvase was included for comparison (Control). The percentage of indel formation was determined by densitometry. No off-target effects were detected following the gesicle treatment.

To confirm the results of the resolvase assay, PCR amplicons of both the *EMX1* locus and off-target 4 were subcloned and sequenced.

Corroborating the previous results (Figure 1, above), plasmid delivery resulted in indel formation at both the *EMX1* and off-target 4 loci (Figure 2). Gesicle delivery resulted in indel formation only at the *EMX1* locus.



**Figure 2. Sanger sequencing confirmed reduced off-target effects with CRISPR/Cas9 gesicles.** *EMX1* and off-target 4 PCR amplicons were subcloned using the Guide-it Indel Identification Kit. Sequencing data for the different clones were aligned with the wild-type sequence (underlined), revealing a range of deletions and insertions (highlighted in red) in both the *EMX1* and off-target 4 sites when cells were treated with plasmid transfection. For the cells treated with gesicles, indels were detected only at the *EMX1* target site, not at the off-target 4 site.

## Conclusions

The [Guide-it CRISPR/Cas9 Gesicle Production System](#) is a novel methodology for the delivery of active Cas9-sgRNA RNP complexes to target cells for CRISPR/Cas9 gene editing. Delivery of these active RNP complexes in this manner prevents both the overexpression and genomic integration of Cas9 inherent to plasmid-based delivery. Therefore, Guide-it CRISPR/Cas9 Gesicles advance genome modification by enabling efficient editing of target loci while also reducing potential off-target effects.

## Methods

### Production of gesicles containing Cas9 protein and sgRNA

The target sgRNA against *EMX1* was cloned into the prelinearized pGuide-it-sgRNA1 vector included in the Guide-it CRISPR/Cas9 Gesicle Production System. This cloned plasmid was added to the provided Guide-it Gesicle Packaging Mix. The Gesicle Packaging Mix contains lyophilized Xfect Transfection Reagent premixed with an optimized formulation of plasmids encoding Cas9 and all the other elements needed for gesicle production.

### Modification of the *EMX1* and off-target 4 genes in HEK 293T target cells

$5.0 \times 10^5$  HEK 293T cells were plated in 24-well plates. 24 hr later, cells were either cotransfected with 500 ng each of plasmids encoding Cas9 and a sgRNA targeting *EMX1* using [Xfect Transfection Reagent](#), or treated with 30  $\mu$ l of Cas9 gesicles (produced as described above).

### Determination of indel formation

72 hr later after plasmid or gesicle treatment, the level of indel formation was determined using the [Guide-it Mutation Detection Kit](#). Crude DNA extracts were prepared from cells. The modified *EMX1* locus and potential off-target site (off-target 4) were amplified using direct PCR. The PCR amplicons were melted and rehybridized, then analyzed using the mismatch-specific nuclease, Guide-it Resolvase. The cleavage reactions were run on an agarose gel, and the percentage of DNA cleavage was determined by densitometry.

## Sanger sequencing for confirmation of indels

*EMX1* and off-target 4 PCR amplicons were subcloned using the [Guide-it Indel Identification Kit](#) and submitted for Sanger sequencing.

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## References

Cong, L. *et al.* Multiplex genome engineering using CRISPR/Cas systems. *Science* **339**, 819–23 (2013).

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<p>The Gesicle Producer 293T Cell Line is a subclone of the transformed human embryonic kidney cell line, HEK 293, which is highly transfectable and supports high levels of protein expression. The cell line also constitutively expresses the simian virus 40 (SV40) large T antigen. Gesicles are produced in these cells via co-overexpression of packaging mix components, which include a nanovesicle-inducing glycoprotein and a protein that is displayed on the cell surface and mediates binding and fusion with the cellular membrane of target cells. Simultaneous overexpression of another protein cargo can result in incorporation of that protein within the gesicles. When combined with a gesicle production system, these cells are capable of producing high gesicle yields.</p>					
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