

# Cas9/sgRNA ribonucleoprotein nucleofection using Lonza 4D nucleofector Version 6

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## Abstract

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## Before start

Grow cells to 80-90% confluency. Plate enough cells for 200K cells per nucleofection reaction.

Mg<sup>2+</sup> is required for cleavage of DNA by Cas9. Avoid buffer containing high concentration of EDTA as it can chelate Mg<sup>2+</sup>.

Use SF cell line solution with added supplements as nucleofection solution for HEK293T cells. Don't leave cells in nucleofection solution for a long time as it might be toxic to the cells.

## Protocol

Warm up trypsin, media and 1x PBS

### Step 1.

Prepare cells (part 1)

### Step 2.

Trypsinize cells and spin down at 200 x g for 10 minutes at RT:

-Leave cells in trypsin at 37C for 3-5 minutes.

-Add in warm media to neutralize trypsin.

Note: don't leave cells in trypsin for a long period of time.

Prepare cells (part 1)

### Step 3.

Remove media containing trypsin and resuspend cells in an appropriate amount of warm 1x

PBS. This step is critical as trypsin and FBS commonly contain RNase activity.

#### Prepare cells (part 1)

##### Step 4.

Count cells. Record the cell concentration (cells/uL). In the meantime, put solution containing cells in 37C

#### Form the crRNA : tracrRNA duplex (if needed)

##### Step 5.

Resuspend RNA oligos (cr and tracr) in IDT duplex buffer to final concentrations of 200 uM.

#### Form the crRNA : tracrRNA duplex (if needed)

##### Step 6.

Mix the two oligos in equimolar concentrations to a final duplex concentration of 100 uM. For example, mixing 1.25 uL of 200 uM crRNA and 1.25 uL of 200 uM tracrRNA yields 2.5 uL of 100 uM guide duplex.

#### Form the crRNA : tracrRNA duplex (if needed)

##### Step 7.

Heat at 95C for 5 minutes. Allowing slow cooling to RT by leaving tubes on block before proceeding.

#### Prepare ribonucleoproteins RNPs mix

##### Step 8.

Add 100 pmol of Cas9 to 120 pmol of gRNA very very slowly:

-For example, if Cas9 is at 40 uM and gRNA stock at 200 uM, add 2.5 uL of 40 uM Cas9 to 0.6 uL of 200 uM gRNA.

#### Prepare ribonucleoproteins RNPs mix

##### Step 9.

Incubate at RT for 10-20 minutes to let RNP form.

#### Prepare cells (part 2)

##### Step 10.

For each nucleofection, pipette 200k cells using a P200 or larger into a 1.5 mL tube.

#### Prepare cells (part 2)

##### Step 11.

Spin 200 x g for 10 minutes at RT to pellet cells softly.

#### Prepare cells (part 2)

##### Step 12.

Carefully remove media off of tubes.

## Prepare cells (part 2)

### Step 13.

Prepare a 12-well plate containing 1 mL of media per well. Pre-warm at 37C.

## Nucleofection

### Step 14.

Prepare and label wells on nucleofection cuvettes. To avoid cells staying in nucleofection solution for a long period of time in the subsequent steps, configure Lonza 4D ahead of time using the recommended cell-type program. Use SF cell line program CM-130 for HEK293T cells.



## REAGENTS

Amaya SF Cell Line 4D-Nucleofector Kit S (96 RCT) V4SC-2096 by [Lonza](#)

Lonza Nucleofector 4d AAF-1002X by [Lonza](#)

## Nucleofection

### Step 15.

After centrifugation, cell pellets are soft so carefully remove media from cells.

## Nucleofection

### Step 16.

Resuspend cells in 20  $\mu$ L of nucleofector solution (SF cell line solution with added supplement for HEK293T) using a P200.

## Nucleofection

### Step 17.

Add the entire RNP mix to the 20  $\mu$ L resuspension and mix using a P200.

## Nucleofection

### Step 18.

If using a repair template, add 1 $\mu$ L of 100 $\mu$ M single-stranded donor DNA (100 pmoles) and mix well.

## Nucleofection

### Step 19.

Add nucleofection mixes to the multiwell cuvette, and cap.

## Nucleofection

### Step 20.

Insert cuvette into nucleofector and zap using the configured program.

## Nucleofection

### Step 21.

Allow cells to sit in nucleofection strips for 10 minutes post-nucleofection. This is supposed to increase efficiency.

## Nucleofection

### Step 22.

Add 80 $\mu$ L of pre-warmed media to each well. Pipette mixture out with a P200 into your pre-warmed 12-well plate.

**Step 23.**

Allow cells 24 hours - 48 hours to settle and recover before attempted downstream analysis. Consider including un-zapped controls to test viability.