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© CRISPR/Cas9-based knock-out in human primary T cells (24-well setup)



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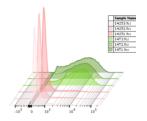
Protocol status: Working

We use this protocol and it's working

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Abstract

This protocol has been optimized for **Thermo Fisher's Neon Electroporation System** and is based on the report Reprogramming human T cell function and specificity with non-viral genome targeting. We have used TrueCut™ Cas9 Protein v2 and reached very comparable KO efficiencies to those previously shown by others.

Guidelines

To reduce variation per electroporation sample and to account for practical issues of electroporation (e.g. sparks), we have also found that a 24-well plate setup where we seeded cells from three independent electroporation reactions (within 10-ul tip) into a single well with 500 ul to 1 ml culture media in it had been the most feasible approach. This setup is especially helpful during initial optimization experiments where the goal is to screen many conditions. Pooling three independent reactions into a single sample provides a virtual buffer against variation across electroporation samples and yields enough treated cells for common assaying approaches, such as Western Blotting. We always recommend optimizing or screening using 10-ul tip setup and then scaling the reactions up to 100-ul once the optimal settings are known.

Materials

MATERIALS

X TrueCut™ Cas9 Protein v2 Thermo Fisher Scientific Catalog #A36497

X TrueGuide™ tracrRNA **Thermo Fisher Scientific Catalog** #A35506

X Neon™ Transfection System 10 μL Kit Thermo Fisher Scientific Catalog #MPK1096



Cas9 RNP preparation

- 1 RNase Zap everything before you start
- 2 Resuspend the crRNA (2 nmol) in 100 ul (making 20 uM) and tracRNA (20 nmol) in 1 ml RNA storage buffer (making 20 uM). Aliquot and keep at -80°C.
- Need 0.375ul crRNA and 0.375 ul tracRNA per well (this way, we'll use 7.5pmol sgRNA per 200K cells as Neon protocol suggests).
- 4 Mix 1.5 ul crRNA and 1.5 ul tracRNA (for 4 rxns) in a PCR tube.
- 5 Keep the RNA mix at 95°C for 5 minutes
 - **(2)** 00:05:00
- 6 Then keep the RNA mix at at 37°C for 25 mins.
 - **(:)** 00:25:00
- For Cas9, Neon suggests 1250 ng Cas9 protein per 200K cells. Our Cas9 is at 5 mg/ml concentration. So, we will need 0.25 ul Cas9/200K cells.
- After incubation of the sgRNA, slowly add 1ul Cas9 (for 4 reactions) in the PCR tube, mix and incubate at 37°C for 15 mins.
 - 00:15:00

Cell preparation and electroporation

- Need 200K cells for one electroporation event. The cells should be in 9 ul T buffer (9 ul T buffer reaction), so that the total volume (9 ul cells + 1 ul RNP mix) will be 10ul for the Neon 10 tip.
- 10 Debead and count the activated cells.
- 11 We need 200,000*3*24 (14.4 million cells).
- To be safe, assume 4 reactions well so we will need 200,000*4*24 (19 million) cells.



- 13 19 million cells are actually good for 96 wells.
- 14 So, we need 96*9 = 864 ul T buffer
- 15 After debeading and counting the cells, spin them down at 200 x g for 7 mins.
- 16 Aspirate the media as much as possible.
- 17 Resuspend the pellet in 864 ul T buffer.
- 18 Add 36 ul of cell mix to each PCR tube and mix well.
- 19 The Cas9 RNP and cells are ready for electroporation
- 20 Electroporation at 1600 V 10 ms 3 pulses

Equipment NAME new equipment Thermo Fisher Scientific Neon $^{\mathsf{M}}$ Transfection System $^{\mathsf{BRAND}}$ SKU MPK5000S

21 Seed the electroporated cells on the prepared 24-well-plate with warm T cell media.

Profiling



22 The time to profile CRISPR/Cas9 treated cell depends on the particular assay of interest but in general, cells can be profiled via flow cytometer for surface proteins and via Western Blot for internal proteins 3 days after electroporation.