Alt-R™ CRISPR-Cas12a (Cpf1) System

Delivery of ribonucleoprotein complexes into Jurkat T cells using the Neon™ Transfection System

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## Revision History

<table>
<thead>
<tr>
<th>Version</th>
<th>Date released</th>
<th>Description of changes</th>
</tr>
</thead>
<tbody>
<tr>
<td>2</td>
<td>June 2021</td>
<td>Updated to include new Cas12a variants (A.s. Cas12a (Cpf1), A.s. Cas 12a (Cpf1) Ultra, L.b. Cas12a Ultra)</td>
</tr>
<tr>
<td>1.1</td>
<td>May 2018</td>
<td>Added note about use of updated Cas12a enzymes (V3)</td>
</tr>
<tr>
<td>1</td>
<td>May 2017</td>
<td>Original protocol</td>
</tr>
</tbody>
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Alt-R CRISPR-Cas12a (Cpf1) System: Delivery of RNP into Jurkat cells (Neon Transfection System)
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INTRODUCTION

This protocol describes the optimized delivery of a Cas12a (Cpf1) ribonucleoprotein (RNP) complex, containing Alt-R™ CRISPR-Cas12a (Cpf1) crRNA and any of the available Cas12a variants (A.s. Cas12a (Cpf1), A.s. Cas 12a (Cpf1) Ultra, L.b. Cas12a Ultra), into Jurkat T cells using the Neon™ Transfection System (Thermo Fisher) with 10 µL electroporation cuvettes [1]. Application of this protocol can be found in a DECODED newsletter article [2]. Other cell lines may require optimization of electroporation parameters and RNP concentration to ensure desired editing efficiency for your experimental needs. To investigate on-target mutations and estimate editing efficiency, we recommend using the Alt-R Genome Editing Detection Kit [3].

The CRISPR-Cas12a (Cpf1) system is distinct from the more commonly used CRISPR-Cas9 system. For example, Cas12a (Cpf1) nuclease does not require a tracrRNA. This nuclease recognizes a T-rich, protospacer-adjacent motif (PAM: TTTV, where V is an A, C, or G base) and creates a staggered double-stranded DNA cut with a 5’ overhang. For additional information, visit www.idtdna.com/CRISPR-Cpf1.

IMPORTANT

1. Use low-passage, healthy cells. A critical factor affecting the success of electroporation is the health of the cells. It is important to:
   - Use the lowest passage number cells available
   - Subculture cells for at least 2–3 days before the electroporation procedure
   - Replace the media the day before electroporation
   - Determine the optimal confluency for your cell type

2. Wash cells before electroporation. FBS and trypsin may contain RNase activity that can quickly degrade CRISPR RNA components. Therefore, it is crucial to wash cells with PBS to remove FBS-containing media or trypsin. Alternatively, for adherent cells, use “enzyme-free” dissociation media instead of trypsin.

3. Alt-R Cas12a (Cpf1) Electroporation Enhancer, a non-targeting, single-stranded carrier DNA, is required for successful electroporation. See the Performance section at www.idtdna.com/CRISPR-Cpf1 for supporting data.

4. We recommend the use of appropriate controls in your experiment such as HPRT-specific, positive control crRNA and a non-targeting negative control. For suggested sequences for studies in human, mouse, or rat cells, see page 5.

   To detect on-target editing of the suggested HPRT control crRNA, Alt-R HPRT PCR Primer Mix (available for human, mouse, or rat) may be used with the Alt-R Genome Editing Detection Kit (T7 endonuclease I assay).
CONSUMABLES AND EQUIPMENT

**Instruments, kits, and reagents**

<table>
<thead>
<tr>
<th>Consumables and equipment</th>
<th>Ordering information</th>
</tr>
</thead>
<tbody>
<tr>
<td>Neon Transfection System</td>
<td>Thermo Fisher Scientific (Cat # MPK5000)</td>
</tr>
<tr>
<td>Neon Transfection System 10 µL Kit</td>
<td>Thermo Fisher Scientific (Cat # MPK1096)</td>
</tr>
<tr>
<td>RPMI-1640 Medium (RPMI)</td>
<td>ATCC (Cat # 30-2001)</td>
</tr>
<tr>
<td>Fetal bovine serum (FBS)</td>
<td>General laboratory supplier</td>
</tr>
<tr>
<td>1X Phosphate buffered saline (PBS)</td>
<td>General laboratory supplier</td>
</tr>
<tr>
<td>Alt-R CRISPR-Cpf1 crRNA</td>
<td>IDT custom crRNA (<a href="www.idtdna.com/CRISPR-Cpf1">www.idtdna.com/CRISPR-Cpf1</a>)</td>
</tr>
<tr>
<td>Alt-R Cas12a (Cpf1)</td>
<td>IDT (Cat # 1076300)</td>
</tr>
<tr>
<td>• variant A.s. Cas12a (Cpf1)</td>
<td>1081068, 1081069</td>
</tr>
<tr>
<td>• variant A.s. Cas 12a (Cpf1) Ultra</td>
<td>10001272, 10001273, 10007804</td>
</tr>
<tr>
<td>• variant L.b. Cas12a Ultra</td>
<td>10007922, 10007923, 10007924</td>
</tr>
<tr>
<td>Nuclease-Free IDTE, pH 7.5 (1X TE solution)</td>
<td>IDT (Cat # 11-01-02-02)</td>
</tr>
<tr>
<td>Alt-R Cas12a (Cpf1) Electroporation Enhancer</td>
<td>IDT (Cat # 11-01-02-02)</td>
</tr>
</tbody>
</table>

**Improved enzymes**

The Alt-R Cas12a (Cpf1) enzyme has recently been further optimized. The latest version (V3) can be directly substituted into this protocol in place of the prior Alt-R Cpf1 enzyme.

To order control crRNAs, enter the appropriate sequence into the Cas12a crRNA ordering tool (accessible at [www.idtdna.com/CRISPR-Cpf1](www.idtdna.com/CRISPR-Cpf1)). These sequences are available online for copying and pasting into the ordering tool.

**Recommended control crRNAs**

<table>
<thead>
<tr>
<th>Sequence</th>
</tr>
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<tbody>
<tr>
<td>GGTAAAAGATGGTTAAATGAT</td>
</tr>
<tr>
<td>GGTAAAAGATGGTTAAATGA</td>
</tr>
<tr>
<td>ATGCTTAAGAGGTATTTGTTA</td>
</tr>
<tr>
<td>CGTTAATCGCGTATAATACGG</td>
</tr>
<tr>
<td>CATATTGCGCGTATAGTCGCG</td>
</tr>
<tr>
<td>GGCCTGCATAGTGCAGCG</td>
</tr>
</tbody>
</table>

**Prepare cell cultures for electroporation**

1. Do not use freshly thawed cells for electroporation: passage your cells 1X after thawing, and verify that they grow well and look healthy.
2. Use cells with the lowest passage number possible.
3. Change the cell culture media on the cells 1 day before electroporation.
4. Split cells, if necessary, to obtain optimal confluency for electroporation.

**Note:** For Jurkat cells, optimal cell density is between $1 \times 10^5$ and $1 \times 10^6$ cells/mL at the time of transfection [4].
**Prepare the crRNA**

1. At first use, resuspend Alt-R CRISPR-Cas12a crRNA in IDTE Buffer to create a 100 µM stock solution. For assistance, use the IDT Resuspension Calculator at [www.idtdna.com/SciTools](http://www.idtdna.com/SciTools).

   **Store resuspended RNAs at −20°C.**

2. Dilute crRNA (from Step 1) to a 75 µM working dilution. You will need 1 µL of working dilution for each electroporation in **Form the RNP complex, Step 1**.

**Form the RNP complex**

1. For each well undergoing electroporation, combine the crRNA and Cas12a (Cpf1) Nuclease components, gently swirling the pipet tip while pipetting:

<table>
<thead>
<tr>
<th>Component</th>
<th>Amount</th>
</tr>
</thead>
<tbody>
<tr>
<td>Alt-R CRISPR-Cas12a crRNA (from Prepare the crRNA, Step 2)</td>
<td>1.0 µL</td>
</tr>
<tr>
<td>Alt-R Cas12a (Cpf1) Nuclease</td>
<td>1.0 µL</td>
</tr>
<tr>
<td><strong>Total volume</strong></td>
<td><strong>2.0 µL</strong></td>
</tr>
</tbody>
</table>

   * This 2 µL volume is for 1 electroporation reaction; scale up as necessary for your experiment—we recommend making 1.2X volume needed to correct for pipetting errors.

2. Incubate the mixture at room temperature for 10–20 min.

**Prepare Neon Transfection System**

1. Turn on the Neon system.
2. Enter electroporation settings, or choose a setting from the optimization protocol.
   
   **Note:** In internal experiments, the optimum settings for Jurkat cells were found to be 1600 V, 10 ms pulse width, 3 pulses [2].

3. Set up the Neon Pipette Station by filling the Neon Tube with Electrolytic Buffer (included in the Neon Transfection System Kit) and insert into the station.

**Perform electroporation of cells**

   
   a. At first use, resuspend the Alt-R Cas12a (Cpf1) Electroporation Enhancer to 100 µM in IDTE to create a stock solution.
   
   b. For each set of experiments, dilute stock to 10.8 µM (working solution). You will need 2 µL of working solution for each electroporation in Step 11.

2. Prepare culture plates to receive cells following electroporation.
   
   a. Set 1: For each electroporation sample, fill a well with 190 µL of culture media (RPMI, 10% FBS) to resuspend cells in Step 16.
   
   b. Set 2: For each electroporation sample, fill 3 wells with 150 µL of culture media (RPMI, 10% FBS) for growth in Step E17.
   
   c. Store all plates in a tissue culture incubator (37°C, 5% CO₂).

3. Pipette cells up and down to dissociate cell clumps.
4. Remove as much supernatant as possible without disturbing the pellet.

5. Wash cells in 5 mL of 1X PBS, and then centrifuge at 600 rpm (Beckman GH-3.8 rotor: 58 x g) for 10 min at room temperature.

Note: FBS in the growth media may contain RNase activity. Therefore, it is crucial to wash the cells with PBS after spinning down.

6. Remove as much supernatant as possible without disturbing the pellet.

7. Resuspend cells by adding 8 µL of Resuspension Buffer R per 5 x 10⁵ cells.

8. For each electroporation, add the following to a 200 µL PCR tube:

<table>
<thead>
<tr>
<th>Component</th>
<th>Amount (µL)*</th>
</tr>
</thead>
<tbody>
<tr>
<td>crRNA:Cas12a RNP complex (from Form the RNP complex, Step 2)</td>
<td>2</td>
</tr>
<tr>
<td>Cell suspension (from Step 10)</td>
<td>8</td>
</tr>
<tr>
<td>10.8 µM Alt-R Cas12a (Cpf1) Electroporation Enhancer (from Step 1b)</td>
<td>2</td>
</tr>
<tr>
<td><strong>Total volume</strong></td>
<td><strong>12</strong></td>
</tr>
</tbody>
</table>

* The final concentration for each electroporation is 6.25 µM crRNA, 5.25 µM Cas12a nuclease, and 1.8 µM Cas12a (Cpf1) electroporation enhancer.

9. Insert a Neon Tip into the Neon Pipette.

10. Pipette 10 µL of cell/RNP complex mixture (from Step 11) into the Neon Tip, avoiding air bubbles.

11. Insert the Neon Pipette and Tip into the Pipette Station. Verify the presence of Electrolytic Buffer in the Neon Tube.

12. Press Start.

13. After electroporation, transfer cells to wells containing 190 µL of pre-warmed culture media (RPMI, 10% FBS) (from Step 2a) and slowly resuspend.

14. Transfer 50 µL of resuspended cells in triplicate to the wells containing 150 µL of pre-warmed culture media (RPMI, 10% FBS) (from Step 2b).

15. Incubate cells in a tissue culture incubator (37°C, 5% CO₂) for 72 hr.

To investigate on-target mutations with the Alt-R Genome Editing Detection Kit (T7EI mismatch assay), follow the protocol [3].
REFERENCES


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