



# Genome editing in the choanoflagellate Salpingoeca rosetta

### David Booth<sup>1</sup>

<sup>1</sup>University of California, San Francisco



King Lab Protist Research to Optimize Tools in Genetics (PROT-G)







### **ABSTRACT**

This protocol details the preparation and execution of CRISPR/Cas9 genome editing in S. rosetta. The protocol builds on a method to transfect macromolecules into S. rosetta for delivering a purified Cas9 ribonucleoprotein from Streptomyces pyogenes (SpCas9 RNP) into S. rosetta. Upon cleaving the S. rosetta genome at locations specified by the guide RNA (gRNA) of the SpCas9 RNP, S. rosetta can use DNA oligonucleotides as templates to repair the double-stranded break. Those repair templates can encode foreign sequences and mutations for editing the S. rosetta genome, so long as DNA oligonucleotides have >30 bases of sequence that is homologous to both sides of the Cas9 cleavage site.

#### **GUIDELINES**

Perform cell culturing and transfection procedure inside of a biosafety cabinet to maintain sterility.

#### STEPS MATERIALS

NAME ~	CATALOG #	VENDOR ~
crRNA	Custom Order	Integrated DNA Technologies
Duplex Buffer	11-01-03-01	Integrated DNA Technologies
tracrRNA	1072534	Integrated DNA Technologies
DNA Oligonucleotide	Custom Order	Integrated DNA Technologies
EnGen Cas9 NLS, S. pyogenes - 400 pmol	M0646T	New England Biolabs
Salpingoeca rosetta cultured with Echinicola pacifica (SrEpac)	PRA-390	ATCC
Falcon 525cm² Rectangular Straight Neck Cell Culture Multi-Flask, 3-layer with Vented Cap	353143	Corning
SF Cell Line 96-well Nucleofector Kit	V4SC-2096	Lonza
Papain from papaya latex	P3125-100MG	Millipore Sigma

BEFORE STARTING

Please consult the attached file of media recipes for artificial seawater, high nutrient media, and low nutrient media.

MediaRecipes.pdf

**Culture Cells** 

Seed a large culture of S. rosetta.

1.1 Two days prior to transfection, inoculate 120 ml of high nutrient media with a culture of *S. rosetta* feeding on *E. pacifica* to a final concentration of *S. rosetta* of [M]8000 cells/ml.

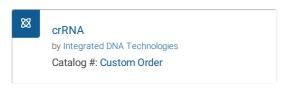


1.2 Grow the culture for **48:00:00** in a 3-layer flask at **22 °C with 60% humidity**.



## Prepare Gene Editing Cargo

- 2 Prepare a guide RNA (gRNA) that binds to SpCas9 and targets DNA by annealing CRISPR RNA (crRNA) with the trans-activating CRISPR RNA (tracrRNA).
- 2.1 Resuspend crRNA in duplex buffer (30 mM HEPES-KOH, pH 7.5; 100 mM potassium acetate) to a final concentration of [M]200 Micromolar (µM).





2.2 Resuspend tracrRNA in duplex buffer to a final concentration of [M]200 Micromolar (µM).



2.3

Mix equal volumes of crRNA (  $\odot$  go to step #2.1 ) and tracrRNA (  $\odot$  go to step #2.2 ) to have a final concentration of [M]100 Micromolar ( $\mu$ M) gRNA, which is the annealed complex of crRNA and tracrRNA.

- 2.4 Incubate the gRNA solution at § 95 °C in an aluminum block for © 00:05:00.
- 2.5 Place the aluminum block was placed at & Room temperature to slowly cool the qRNA to & 25 °C.
- 2.6 Store the gRNA at § -20 °C.
- 3 Prepare DNA oligonucleotides that serve as repair templates after SpCas9 cleavage.
- 3.1 Dissolve oligonucleotides to a final concentration of [M]250 Micromolar (µM) in 10 mM HEPES-KOH, pH 7.5.



- 3.2 Incubate the dissolved oligonucleotides at § 55 °C for © 01:00:00.
- 3.3 Store oligonucleotides at § -20 °C.
- 3.4 Before starting nucleofections, ensure that the oligonucleotides are fully dissolved by incubating them at § 55 °C for © 01:00:00, which concurs with the assembly of the SpCas9/gRNA complex.
- 4 Assemble SpCas9 with the gRNA to form the SpCas9 RNP.

4.1 For one transfection, place **2 μl** of [M]**20 Micromolar (μM) SpCas9** in the bottom of a 0.2 ml PCR tube.



4.2

Add  $\square 2 \mu l$  of [M] 100 Micromolar ( $\mu M$ ) gRNA (  $\odot$  go to step #2 ) by slowly pipetting up and down with SpCas9 to gently mix the gRNA together. This solution is called the "SpCas9 ribonucleoprotein (RNP)."

4.3 Incubate the *Sp*Cas9 RNP at & **Room temperature** for **© 01:00:00** (roughly the time to complete the preparation of *S. rosetta* for priming, see below).

### Prepare transfection rReagents

5 Prepare SF Buffer (Lonza) for transfections.



5.1

Add all of buffer B (smaller volume that may also be called supplement 1) to buffer A (larger volume).

- 5.2 Store on ice until ready for use. The combined buffer can also be stored at 4°C for up to 3 months.
  - The combined buffer can be stored at 4°C for up to 3 months.
  - Because the Lonza kits can be so expensive, we recommend aliquoting large volumes of the SF components (900 µl aliquots for buffer A and 200 µl aliquots for buffer B) to prevent SF buffer from spoiling after buffers A and B have been combined.
- 6 Prepare the priming buffer.

6.1

Dilute papain to a final concentration of 100  $\mu$ M in dilution buffer (50 mM HEPES-KOH pH 7.5, 200 mM sodium chloride, 20% [v/v] glycerol, and 10 mM cysteine) from a stock solution of 1 mM papain (Millipore Sigma, St. Louis, MO; Cat. No. P3125-100MG]), and incubate at room temperature just before priming cells for transfection.



The dilution buffer [50 mM HEPES-KOH pH 7.5, 200 mM sodium chloride, 20% (v/v) glycerol and 10 mM cysteine] should be sterile filtered through a 0.22 µm filter.

The dilution buffer may also be prepared ahead of time and stored in a -80°C freezer just before its use.

6.2 Make a solution of the remaining components of the priming buffer (40 mM HEPES-KOH, pH 7.5; 34 mM lithium citrate; 50 mM L-cysteine; 15% [wt/vol] PEG 8000). DO NOT combine the papain and priming buffer unti just before adding the priming buffer to cells.



The priming buffer without papain should be sterile filtered through a  $0.22\,\mu m$  filter.

The priming buffer without papain can also be made ahead of time and stored at -80°C until it is used. Be sure that the priming before is warmed to room temperature prior to use.

## Wash Cells

- 7 Prepare S. rosetta for transfection by washing away feeder bacteria.
- 7.1 The 120 ml culture of *S. rosetta* feeding on *E. pacifica* ( 5 go to step #1 )was homogenized by vigorous shaking and then split into 40 ml aliquots in 50 ml conical tubes.
- 7.2

Vigorously shake the aliquots and centrifuge the cells for **⊙ 00:05:00** at **⊚ 2000 x g** and **♂ 22 °C** in a swinging bucket rotor.

Use a serological pipette to gently remove from the cell pellet all but 2 ml of the supernatant, which remains cloudy with *E. pacifica* bacteria. With a fine tip transfer pipette, gently remove the remaining liquid near the pellet.

- 7.4 The three cell pellets were resuspended in a total volume of 50 ml artificial seawater, combined into one conical tube, and vigorously shaken to homogenize the cells.
- For a second time, the resuspended cells were centrifuged for for © 00:05:00 at @2000 x g and & 22 °C in a swinging
- 7.6 The supernatant was removed as before ( 🕁 go to step #7.3 ).

7.5

bucket rotor.

- 7.7 The pellet was resuspended in **50 ml** of artificial seawater, and the cells were homogenized by vigorous shaking.
- 7.8
  - The cells were centrifuged for a third time for  $\bigcirc 00:05:00$  at  $②2200 \times g$  and  $②22^{\circ}C$ .
- 7.9 Remove the supernatant as before ( go to step #7.3 ).
- 7.10 Resuspend the cell pellet in 400 µl of artificial seawater. This resuspension is called the "washed cells."
  - Prepare  $\blacksquare 100 \, \mu l$  aliquots of [M]50000000 cells/ml.
- 8.1 Dilute □2 μl of "washed cells" ( ♦ go to step #7.10 ) into □196 μl of artificial seawater.
- 8.2 Fix the diluted cells with  $2 \mu$  of 37.5% formaldehyde and homogenize by vortexing.

Pipet the fixed cells into a fixed chamber slide and determine the cell concentration. 8.3 Remember that concentration of diluted and fixed cells is a 100-fold dilution from the "washed cells." Be sure to factor that dilution into your concentration. Cells can be counted on a hemacytometer (Neubauer with brightlines) or with an automated cell counter. We recommend a Luna-FL automated cell counter. LUNA-FL **Dual Fluorescence Cell Counter** Logos Biosystems L20001 👄 After determining the cell concentration, dilute the "washed cells" to final concentration of [M]50000000 cells/ml and split into 100 µl aliquots. One aliquot provides enough cells for 12 nucleofections. **Prime Cells** Prime cells for nucleofection by degrading the glycocalyx that surrounds S. rosetta. 9.1 Spin the □100 µl aliquots of washed cells ( ♦ go to step #8.4 ) at ⊕800 x g and § 22 °C for ♦ 00:05:00 . Gently remove the supernatant from the cell pellet with a gel-loading pipette tip. 9.2 X 9.3 Combine the priming buffer components ( ogo to step #6) to make a final priming buffer (40 mM HEPES-KOH, pH 7.5; 34 mM lithium citrate; 50 mM l-cysteine; 15% [wt/vol] PEG 8000; and 1 µM papain) 9.4 Resuspend each cell pellet in 100 µl of priming buffer. 9.5

Incubate cells for © 00:35:00 at § Room temperature .

9.6 Add 10 µl of [M] 50 mg/ml Bovine Serum Albumin to each aliquot of primed cells for quenching proteolysis from the priming buffer. 9.7 Centrifuge cells at (3) 1250 x g and (3) 22 °C for (5) 00:05:00 . Gently remove the supernatant from the cell pellet with a gel-loading pipette tip. 9.8 9.9 Resuspended each cell pell in 25 µl of SF Buffer ( 5 go to step #5 ). This suspension of cells is called the "primed cells." 9.10 Store the "primed cells" on ice while preparing nucleofection reactions. Transfect Cells Deliver gene editing cargo via nucleofection. 10 10.1 Add 116 µl of ice-cold SF Buffer ( 5 go to step #5 ) to the SpCas9 RNP ( 5 go to step #4.3 ), which has a total volume of **4 μl** . For reactions that use two different gRNAs, assemble each SpCas9 RNP separately then combine each SpCas9 RNP at this step. After the SpCas9 RNPs have been combined, add 16 µl of ice-cold SF Buffer 10.2 Add 2 ul of the repair oligonucleotide template to the PCR tube with SpCas9 RNP and SF Buffer ( one to step #10.1). 10.3 Add 2 4 of "primed cells" (from 5 go to step #9.10) to the PCR tube with SpCas9 RNP, SF Buffer, and the repair template ( 🜣 go to step #10.2 ). This solution, which is called the "nucleofection mix," should have a total volume of **⊒24 μl** . Transfer the entire nucleofection mix into one well of a 96-well nucleofection plate. 10.4 At this point, prepare for the recovery step, by transferring the recovery buffer into a convenient vessel and setting the pipette to  $\square 100 \mu I$ .

- 10.5 Pulse the nucleofection plate with the CM156 pulse.
  - 4D-Nucleofector Core Unit
    Control system for performing nucleofection
    Lonza AAF-1002B 🖘
  - 96-well Shuttle Device
    Add-on for Nucelofector 4d device to perform plate-based nucleofections
    Lonza AAM-1001S 🖘

## **Rest and Recover Cells**

- 11 Allow membranes to reseal by resting cells in recovery buffer before growing cells again in media.
- 11.1 Immediately after transfection, add **100 μl** of ice-cold recovery buffer (10 mM HEPES-KOH, pH 7.5; 0.9 M sorbitol; 8% [wt/vol] PEG 8000) to each nucleofection transfection and gently mixed by firmly tapping the side of the plate.
- 11.2 Allow cells to rest in recovery buffer for © 00:05:00.
- Gently mix the well in the nucleofection plate by pipetting up and down before transferring the entire volume in nucleofection well (the nucleofection mix plus the recovery buffer) into to **2 ml** of low nutrient media in one well of a 6 well plate.
- 11.4 Incubate at § 22 °C and 60% humidity for © 00:30:00
- 12 Add E. pacifica food and grow transfected cells.
- 12.1 Add **10 μl** of [M]**10 mg/ml** of *E. pacifica* to the wells in the 6 well plate.
  - Prepare the *E. pacifica* solution by resuspending a frozen, 10 mg pellet of *E. pacifica* in 1 ml of artificial seawater.

12.2	
	(

Incubate the 6 well plate at § 22 °C and 60% humidity for © 24:00:00 before using in downstream experiments.

## (Optional) Select for Cycloheximide Resistance

13

Add 10 µl of [M] 1 mg/ml of cycloheximide to the 2 ml culture of transfected cells after allowing the cells to fully recover.



Cycloheximide is toxic. Handle carefully and properly dispose.

14

Incubate the cells in cycloheximide for **96:00:00** prior to genotyping and clonal isolation.

This is an open access protocol distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited