



Version 2

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# Genome editing in the choanoflagellate *Salpingoeca rosetta* V.2

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

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## 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* (*Sp*Cas9 RNP) into *S. rosetta*. Upon cleaving the *S. rosetta* genome at locations specified by the guide RNA (gRNA) of the *Sp*Cas9 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.

## DOI

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## PROTOCOL INTEGER ID

53152

## GUIDELINES

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

## MATERIALS TEXT

### STEP MATERIALS

[DNA Oligonucleotide Integrated DNA](#)

**Technologies Catalog #Custom Order** Step 3.1

[tracrRNA Integrated DNA](#)

**Technologies Catalog #1072534** Step 2.2

[Falcon 525cm<sup>2</sup> Rectangular Straight Neck Cell Culture Multi-Flask, 3-layer with Vented](#)

**Cap Corning Catalog #353143** Step 1.2

[EnGen Cas9 NLS, \*S. pyogenes\* - 400 pmol New England](#)

**Biolabs Catalog #M0646T** Step 4.1

[SF Cell Line 96-well Nucleofector](#)

**Kit Lonza Catalog #V4SC-2096** Step 5

[Salpingoeca rosetta cultured with Echinicola pacifica](#)

**(SrEpac) ATCC Catalog #PRA-390** Step 1.1

[crRNA Integrated DNA](#)

**Technologies Catalog #Custom Order** Step 2.1

[Duplex Buffer Integrated DNA](#)

**Technologies Catalog #11-01-03-01** Step 2.1

[Papain from papaya latex Millipore](#)

**Sigma Catalog #P3125-100MG** Step 6.1

## BEFORE STARTING

Please consult the attached file of media recipes for artificial seawater, high nutrient media, and low nutrient media. [MediaRecipes.pdf](#)

## Culture Cells

### 1 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 [8000 cells/ml](#).

[Salpingoeca rosetta cultured with Echinicola pacifica](#)

**(SrEpac) ATCC Catalog #PRA-390**

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

[Falcon 525cm<sup>2</sup> Rectangular Straight Neck Cell Culture Multi-Flask, 3-layer with Vented Cap](#) **Corning Catalog #353143**

## 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 **100 Micromolar (μM)** .

[crRNA Integrated DNA](#)  
**Technologies Catalog #Custom Order**

[Duplex Buffer Integrated DNA](#)  
**Technologies Catalog #11-01-03-01**

- 2.2 Resuspend tracrRNA in duplex buffer to a final concentration of **100 Micromolar (μM)** .

[tracrRNA Integrated DNA](#)  
**Technologies Catalog #1072534**

- 2.3 

Mix equal volumes of crRNA ( [go to step #2.1](#) ) and tracrRNA ( [go to step #2.2](#) ) to have a final concentration of **100 Micromolar (μM)** 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 gRNA 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.

[DNA Oligonucleotide Integrated DNA](#)

[Technologies Catalog #Custom Order](#)

- 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.

[EnGen Cas9 NLS, S. pyogenes - 400 pmol New England](#)

[Biolabs Catalog #M0646T](#)

- 4.2 

Add **2 μl** of **[M]100 Micromolar (μM) gRNA** ([go to step #2.6](#)) 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 *SpCas9* RNP at **Room temperature** for **01:00:00** (roughly the time to complete the preparation of *S. rosetta* for priming, see below).

#### Prepare transfection Reagents

##### 5 Prepare SF Buffer (Lonza) for transfections.

[SF Cell Line 96-well Nucleofector](#)

[Kit Lonza Catalog #V4SC-2096](#)

- 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 µM 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.

 [Papain from papaya latex Millipore](#)

[Sigma Catalog #P3125-100MG](#)

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 until just before adding the priming buffer to cells.

The priming buffer without papain should be sterile filtered through a 0.22 µm 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 buffer is warmed to room temperature prior to use.

## Wash Cells

## 7 Prepare *S. rosetta* for transfection by washing away feeder bacteria.

7.1 Homogenized the **120 mL** culture of *S. rosetta* feeding on *E. pacifica* ( [go to step #1.2](#) ) by vigorously 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.

7.3 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.

7.5 

For a second time, the resuspended cells were centrifuged for **00:05:00** at **2000 x g** and **22 °C** in a swinging bucket rotor.

7.6 The supernatant was removed as before ( [go to step #7.3](#) ).

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 **00:05:00** at **2200 x g** and **22 °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."

## 8 Prepare **100 µl** aliquots of **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 µl** of 37.5% formaldehyde and homogenize by vortexing.

8.3 Pipet the fixed cells into a fixed chamber slide and determine the cell concentration.

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 [↗](#)

8.4 After determining the cell concentration, dilute the "washed cells" to final concentration of **50000000 cells/ml** and split into **100 µl** aliquots.

One aliquot provides enough cells for 12 nucleofections.

## Prime Cells

### 9 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**.

9.2 Gently remove the supernatant from the cell pellet with a gel-loading pipette tip.

9.3 

Combine the priming buffer components ( [↗ go to step #6.2](#) ) 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 **50 mg/ml Bovine Serum Albumin** to each aliquot of primed cells for quenching proteolysis from the priming buffer.

9.7 Centrifuge cells at **1250 x g** and **22 °C** for **00:05:00**.

9.8 Gently remove the supernatant from the cell pellet with a gel-loading pipette tip.

9.9 Resuspend each cell pellet in **25 µl** of SF Buffer ( [go to step #5.2](#) ). This suspension of cells is called the "primed cells."

9.10 Store the "primed cells" on ice while preparing nucleofection reactions.

## Transfect Cells

### 10 Deliver gene editing cargo via nucleofection.

10.1 Add **16 µl** of ice-cold SF Buffer ( [go to step #5.2](#) ) to the *SpCas9* RNP ( [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 µl** of the repair oligonucleotide template ( [go to step #3.4](#) ) to the PCR tube with *SpCas9* RNP and SF Buffer ( [go to step #10.1](#) ).

10.3 Add **2 µl** of "primed cells" (from [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** .

**10.4** Transfer the entire nucleofection mix into one well of a 96-well nucleofection plate.

At this point, prepare for the recovery step, by transferring the recovery buffer into a convenient vessel and setting the pipette to **100 µl** .

**10.5** Pulse the nucleofection plate with the CM156 pulse.

4D-Nucleofector Core Unit  
Control system for performing nucleofection  
Lonza      AAF-1002B      [link](#)

96-well Shuttle Device  
Add-on for Nucleofector 4d device to perform plate-based nucleofections  
Lonza      AAM-1001S      [link](#)

## 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** .

**11.3** 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 **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 **1 µg/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.