

# P1 Kidney Cold-Active Protease Single Cell Dissociation

# **Andrew Potter, Steve Potter**

#### **Abstract**

Method used to derive single cell suspension from P1 mouse kidneys on ice, generating a cell suspension with greatly reduced artifact gene expresion changes and suitable for downstream analysis using 10x Chromium or DropSeq.

Citation: Andrew Potter, Steve Potter P1 Kidney Cold-Active Protease Single Cell Dissociation. protocols.io

dx.doi.org/10.17504/protocols.io.nzkdf4w

Published: 23 Mar 2018

# **Guidelines**

# **Storage Conditions of Reagents**

#### Reagent

DPBS (Thermofisher, 14190144) 0.5 M EDTA (Ambion, AM9260G) BSA (Sigma, A8806) Protease from *Bacillus Licheniformis* (Sigma,

Protease from *Bacillus Licheniformis* (Sigma, P5380) DNAse 1 (Applichem, A3778)

#### **Storage Condition**

4°C room temp. 4°C

Store 100  $\mu L$  aliquots (100 mg/mL) in DPBS at -80°C Store 10  $\mu L$  aliquots (250 U/10  $\mu L)$  in DPBS at -80°C

#### **Required Equipment**

**Equipment** Supplier Catalog no. gentleMACS dissociator Miltenyi 130-093-235

# The protocol workflow is as follows:

- A. Isolate Kidney
- B. Initial digestion: triturate on ice
- C. Perform gentleMACS
- D. Continue triturating on ice
- F. Preparing cells for Chromium/DropSeq

## **BEFORE STARTING**

Prepare Bacillus Licheniformis enzyme mix just prior to starting dissociation:

Volume (μl) Reagent			Final concentration
	894	DPBS	1X
	1	0.5 M EDTA	0.5 mM
	5	DNAse 1 (250 U/10 $\mu$ L)	125 U / mL
	100	B. Lich (100 mg/mL)	10 mg/mL

+25 mg tissue / 1 mL enzyme mix

To prepare 0.01% BSA/PBS:

Make stock of 10% BSA in DPBS and store at -20 °C. To make PBS/BSA 0.01% aliquot 50 mL of DPBS in 50 mL conical and pipet in 50  $\mu$ L of 10% BSA stock.

Prepare 10% FBS/PBS with heat-inactivated FBS.

#### **Protocol**

#### Step 1.

Extract & isolate P1 kidneys in ice-cold PBS.

#### Step 2.

Mince kidneys on top of petri dish, on ice, using razor blade.

#### Step 3.

Weigh out 25 mg of tissue for each tube of B. Lich. enzyme mix (2 tubes total).

**■** AMOUNT

25 mg Additional info:

#### Step 4.

Incubate tissue + enzyme on ice for 7 minutes while triturating 15 strokes using 1 mL pipet every 2 minutes set to 700  $\mu$ L - first with tip cut off.

#### Step 5.

Monitor digestion by taking small aliquot and visualizing under scope (every 5 minutes).

#### Step 6.

After 7 minutes, take the digest mix (combine the two tubes) and pipet into Miltenyi C-tube (placed on ice); take C-tube to gentleMACS placed in 4° cold room. Run program brain 03 two times.

- **↓** TEMPERATURE
- 4 °C Additional info:

#### Step 7.

After MACS, briefly quick spin the MACS tube (to 500 G) at 4 °C to ensure contents are in the bottom of the tube.

- **TEMPERATURE**
- 4 °C Additional info:

# Step 8.

Re-suspend and visualize cells using scope by taking small aliquot and using a slide; continue digesting cells in C-tube on ice for 8 additional minutes while triturating every 2 min 15 strokes using a 1 mL pipet.

#### Step 9.

Add 3 mL ice-cold 10% FBS/PBS to digest mix in C-tube to inhibit the protease.

**■** AMOUNT

3 ml Additional info: ice-cold 10% FBS/PBS

# Step 10.

Transfer digest mix to a 15 mL conical. Spin 600 G for 5 minutes at 4 °C; discard supernatant; resuspend cell pellet in 2 mL ice-cold PBS/BSA.

**■** AMOUNT

2 ml Additional info: re-suspend in PBS/BSA

## **▮** TEMPERATURE

4 °C Additional info:

# **Step 11.**

Filter re-suspended cells using 30 uM filter on sterile 50 mL conical on ice - rinse filter with 4 mL ice-cold PBS/BSA. Transfer flow-through to 15 mL conical.

**■** AMOUNT

4 ml Additional info: rinse filter

# **Step 12.**

Spin 15 mL conical tube containing filtered cells 600 G for 5 minutes at 4 °C; discard supernatant and re-suspend pellet in 10 mL ice-cold PBS/BSA.

**■** AMOUNT

10 ml Additional info: PBS/BSA

**↓** TEMPERATURE

4 °C Additional info:

# **Step 13.**

Repeat rinse/spin in ice-cold PBS/BSA.

# Step 14.

Remove supernatant and re-suspend in 1-2 mL ice-cold PBS/BSA.

# **Step 15.**

Examine using hemocytometer and adjust concentration to 100 cells/uL for DropSeq or 1,000 cells/ $\mu$ L for 10X Chromium.