

# Human Kidney / Tumour Tissue Disaggregation for Single Cell RNA Sequencing (10x Genomics platform)

Kevin Loudon, John Ferdinand, Alexandra Riding, Menna Clatworthy

## Abstract

**Citation:** Kevin Loudon, John Ferdinand, Alexandra Riding, Menna Clatworthy Human Kidney / Tumour Tissue Disaggregation for Single Cell RNA Sequencing (10x Genomics platform). **protocols.io**

dx.doi.org/10.17504/protocols.io.qtrdwm6

**Published:** 09 Jul 2018

## Protocol

### Tissue Preparation

#### Step 1.

Take dissected tissue (renal cortex, medulla or tumour) and weigh tissue (typical biopsy size used 0.5 - 1 gram)

### Tissue Preparation

#### Step 2.

Pour approximately 2-3 mLs of "Digest Mix" onto sample in 10cm<sup>3</sup> petridish

#### 📌 NOTES

#### PREPARATION OF DIGEST MIX

##### Ingredients

- (1) RPMI alone
- (2) Liberase TM (Sigma Aldrich)
- (3) DNase (Sigma Aldrich)

##### For 50mLs of RPMI add:

- > 625 microlitres of Liberase (Stock solution 2.5mg/mL)
- > 250 microlitres of DNase (Stock solution 0.05mg/mL)

### Tissue Preparation

#### Step 3.

Using a razor blade mince into small pieces approximately 2mm<sup>3</sup>.

### Tissue Preparation

#### Step 4.

Transfer tissue into a gentleMACS C tube and add further 3-4 mLs of Digest mix.

#### 📌 NOTES

GentleMACS C tube by Miltenyi Biotec (Cat.130-096-334)

## Tissue Preparation

### Step 5.

Place in shaking incubator at 37°C for 30 minutes.

## Tissue Preparation

### Step 6.

Homogenise sample in GentleMACS tube using program "Spleen 4" and "Lung 2" on GentleMACS dissociator.

## Tissue Preparation

### Step 7.

Pass through a 100µm cell strainer with of a 2.5ml syringe plunger and wash through with cold running buffer.

#### NOTES

#### PREPARATION OF RUNNING BUFFER

#### Ingredients (for 1 litre)

- (1) 1L PBS
- (2) 5ml BSA (from reagent diluent kit)
- (2) 4ml 0.5M EDTA

## Tissue Preparation

### Step 8.

Centrifuge in a bench top centrifuge at 2000 RPM for 10 minutes and CAREFULLY remove the supernatant.

## Tissue Preparation

### Step 9.

If sample is contaminated with red blood cells an additional red cell lysis step can be taken.

## Tissue Preparation

### Step 10.

To ensure optimal yield for 10X Genomics single cell platform, a live cell enrichment step is required - this was performed using Miltenyi 'Dead Cell Removal Kit' (Please see manufacturers instructions for further details).

#### REAGENTS

Dead Cell Removal Kit 130-090-101  
by Miltenyi Biotec

#### NOTES

## **LIVE CELL ENRICHMENT (Miltenyi - Dead Cell Removal Kit)**

### **Ingredients**

- (1) Dead Cell removal Kit - Miltenyi (Order No. 130-090-101)
- (2) MACS Column (LS or MS)

### **In brief for MACS colum LS**

- (1) Use LS column for  $10^8$  dead cells or  $10^9$  total cells.
- (2) Remove supernatant completely following previous steps
- (3) Resuspend pellet in 100  $\mu$ L of '*Dead Cell Removal MicroBeads*' per approximately  $10^7$  total cells.
- (4) Incubate 15 minutes at room temperature (20-25 °C).
- (5) Rinse column with 1x binding buffer as per manufacturers instructions.
- (6) Apply cell suspension in 1-10mLs of binding buffer and collect the effluent as the **NEGATIVE** cell population (i.e the live cells).
- (7) Wash cells with PBS for 5 minutes at 1500rpm.

---

## Tissue Preparation

### **Step 11.**

Count the cells and resuspend the live cell suspension in appropriate volume of PBS for the 10X application.

---