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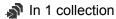


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# Blood sampling, cell isolation, single-cell GEM-generation, globin mRNA blockers and sequencing library preparation protocol



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#### Wim Pierson

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#### **ABSTRACT**

This protocol details blood sampling, cell isolation, single-cell GEM-generation, globin mRNA blockers and sequencing library preparation.

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#### **MATERIALS**

#### Reagents:

- UltraPure 0.5M EDTA, pH 8.0 Thermo Fisher Scientific Catalog #15575-038
- Ethanol ≥70% (v/v) TechniSolv® VWR International Catalog #83801.360
- 1x PBS
- Dulbecco's Phosphate Buffered Saline (without calcium and magnesium) Merck

  MilliporeSigma (Sigma-Aldrich) Catalog #D8537-1L
- Cellaca AOPI Viastain Nexcelom Catalog #CS2-0106-25mL
- 10% 
  MACS BSA Stock Solution Miltenyi Biotec Catalog # 130-091-376
- HISTOPAQUE 1119 Merck MilliporeSigma (Sigma-Aldrich) Catalog #11191-100ML
- MaraPEAK ACK Lysing Buffer Lonza Catalog #BP10-548E
- MicroBeads, mouse Miltenyi Biotec Catalog #130-049-901
- MACS buffer: MACS buffer is a solution containing PBS pH 7.2, 0.5% BSA and 2 mM EDTA. Prepare this by diluting 10% BSA solution 1:20 and 0.5M EDTA 1:250 with 1x PBS.
- PBS-0.04% BSA: PBS-0.04% BSA solution is prepared by diluting 10% BSA solution 1:250 in 1x PBS.
- FastSelect globin mRNA blockers 1X (QIAseq FastSelect Globin Kit, Qiagen GmbH)
- Chromium Next GEM Single Cell 5' Reagent Kits v2 (Dual Index) (10x Genomics)
- Chromium Single Cell V(D)J Amplification Kits, Mouse (10x Genomics)
- Library Construction Kit (10x Genomics)
- Dual Index Kit TT set A, 96 rxns (10x Genomics)

#### **Equipment:**

- Mouse restrainer
- Electric fur clipper (Aesculap, GT415)
- Gauze pads
- Sharps container 1.5L (BD, 305624)
- Vortexer
- Centrifuge with swinging buckets
- Pipettors and tips
- Pipette controller
- Cellaca-MX-AOPI cell counter (Nexcelom)
- QuadroMACS Separator Miltenyi Biotec Catalog #130-090-976

- Chromium Controller (10x Genomics)
- Chromium X (10x Genomics)
- 10x Vortex Adapter (10x Genomics)
- Chromium Next GEM Secondary Holder (10x Genomics)
- 10x Magnetic Separator (10x Genomics)
- Veriti 96-Well Thermal Cycler (Thermo Fisher Scientific)
- NovaSeq6000 platform (PE150) (Illumina)

#### Materials:

- Microlance 3 25G x 5/8" (0.5 x 16mm) needle (BD, 300600)
- Microvette capillary blood system, CB300 K2E, 0.3ML Sarstedt Catalog #16.444
- Microvette capillary blood system, CB300 Z, 0.3 mL Sarstedt Catalog #16.440
- Pluristrainer mini 70 µm filter (Puriselect, 43-10070-40)
- Celltrics 50 µm strainer (Sysmex, 04-004-2327)
- Falcon 50 mL Polypropylene conical Tube (Falcon, 352070)
- Falcon Round-Bottom polystyrene tubes 5 mL (Falcon, 352054)
- LD columns Miltenyi Biotec Catalog #130-042-901
- Cellaca counting plates (Nexcelom, CHM24-A100-004)
- Chromium Next GEM Chip K Single Cell (10x Genomics)

## **Blood sampling**

- 1 Place the animal in the restrainer so that only one of the two the hind legs and tail are free. Stretch out the leg.
- 2 Remove the fur from the lateral side of the hind leg using the electric clipper.
- 3 Locate the lateral saphenous vein (If necessary: swab the skin with a small amount of ethanol to help visualize the vein).

Add 4 10 mL of 1x PBS to the top of the tube to stop the lysis reaction and centrifuge the lysate samp 10m

at 250 x g, Room temperature, 00:10:00 .

Pour off the supernatant and perform a second wash with 4 10 mL of PBS and centrifuge at

10m



350 x g, Room temperature, 00:10:00 .

Pour off the supernatant and resuspend the pellet in equal volume of PBS-0.04% BSA to original whole blood volume.

20 Count cells using the Cellaca or Luna cell counter.

## **Anti-Ter119 Microbeads selection**

45m

21

Note

Important! Use all reagents at 🖁 Room temperature .

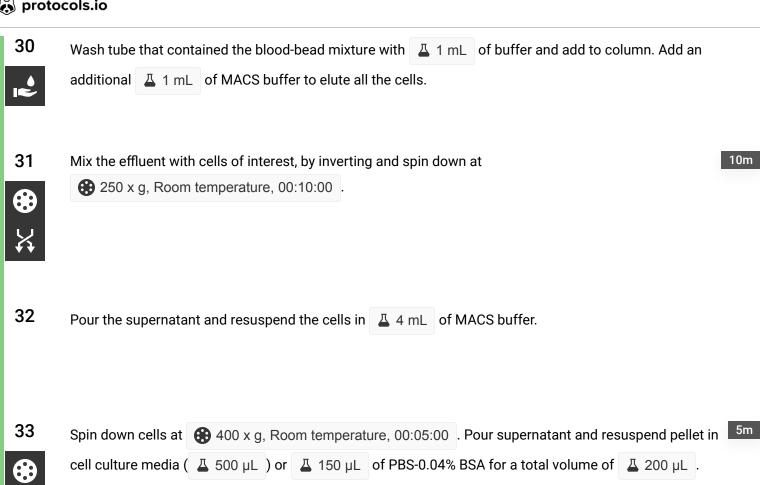
Transfer the blood from the capillary tubes over a 70  $\mu$ m pluristrainer placed on a 5 mL round-bottom tube to remove any clots.

Rinse the original blood tubes twice with  $\Delta 100 \, \mu$ L of MACS buffer and rinse the filter with  $\Delta 3 \, \text{mL}$  MACS buffer.

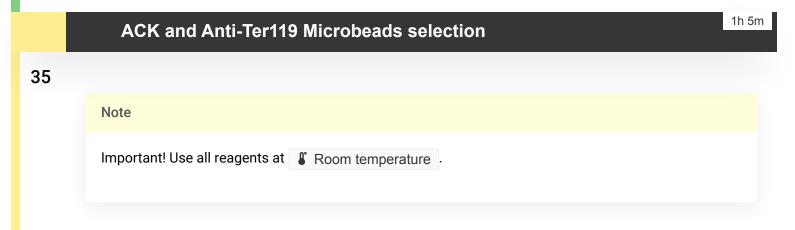
23 5m Centrifugate the samples at 400 x g, Room temperature, 00:05:00 . 24 Pipet off the supernatant ( $\sim 4.3 \, \text{mL}$ ) with a P1000 pipettor. 25 15m Briefly vortex the Anti-Ter119 microbeads and add 🚨 200 µL of beads. Mix well and incubate for (5) 00:15:00 Room temperature . 26 10m Add 🚨 3.5 mL of MACS buffer, mix the solution by inverting and centrifugate at 300 x g, Room temperature, 00:10:00 . 27 Prepare the LD columns by putting them in the magnetic field of a QuadroMACS Separator and rinse columns with A 2 mL MACS buffer; discard flowthrough and place a new 5 mL round-bottom tube under the columns. 28 From the centrifugated blood, remove supernatant using a P1000 pipettor and resuspend cells in Δ 1000 μL of MACS buffer.

Apply the cell suspension onto the column and collect the unlabeled cells which pass through.

29



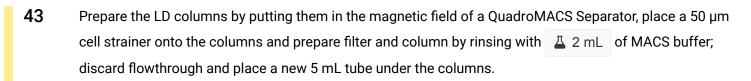
34 Count cells using the Cellaca or Luna cell counter.



Pipet the blood out of the capillary tubes and transfer the entire volume of blood, ~150 μL to a 50 mL tube.

36 10m Add \( \begin{aligned} \Lambda & 2 mL \) of ACK buffer (1x) and incubate the sample on a rocker for (5) 00:10:00 Room temperature . 37 5m Add A HL of PBS to the top of the tube to stop the lysis reaction and centrifuge the sample at 400 x g, Room temperature, 00:05:00 . 38 10m Pour off the supernatant and add 4 mL of MACS buffer to wash the cells, centrifuge at 250 x g, Room temperature, 00:10:00 . 39 Resuspend the pellet in  $\perp$  540 µL of MACS buffer and add  $\perp$  50 µL of Anti-Ter119 MicroBeads. 40 15m Mix well and incubate for 00:15:00 Room temperature . 41 10m After incubation, add 4 3.5 mL of MACS buffer and centrifuge at 300 x g, Room temperature, 00:10:00 . 42 After centrifugation, pour off the supernatant and resuspend cells in \$\textstyle 500 \mu L \text{ MACS buffer.}\$

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- From the centrifugated blood, take off supernatant using a P1000 pipettor and resuspend cells in  $\perp$  1000  $\mu$ L of MACS buffer.
- 45 Apply the cell suspension onto the column and collect the unlabeled cells which pass through.
- Wash tube with 4 1 mL of buffer and add to column. Add an additional 4 1 mL of MACS buffer to elute all the cells. Collect the total effluent which contains the white blood cells.
- Mix the suspension by inverting and spin the total effluent down at 250 x g, Room temperature, 00:10:00.
- Pour the supernatant and resuspend the cells in 4 mL of MACS buffer.
- Spin down cells at 400 x g, Room temperature, 00:05:00. Pour supernatant and resuspend pellet in cell media ( \$\frac{1}{4}\$ 500 \(mu\text{L}\)) or \$\frac{1}{4}\$ 150 \(mu\text{L}\) of PBS-0.04% BSA for a total volume of \$\frac{1}{4}\$ 200 \(mu\text{L}\).

10m

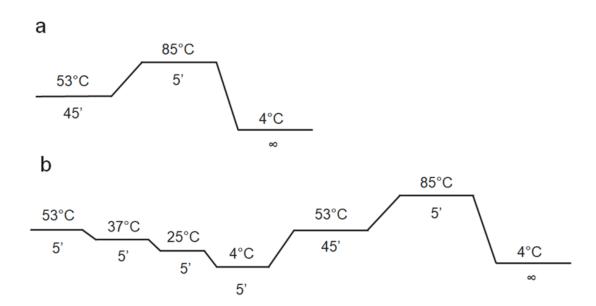
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50 Count cells using the Cellaca or Luna cell counter.

# 45m Histopaque-1119 selection 51 Note Important! Use all reagents at \$\mathbb{\mathbb{L}}\ Room temperature . Prepare 5 mL tube with 4 3 mL Histopaque-1119 gradient fresh on the day of the experiment. 52 Pipet the blood out of the capillary tubes and slowly add it to the top of the gradient, and centrifuge the tu 40m for \$\infty\$ 400 x g, 00:30:00 | with low deceleration for \$\infty\$ 00:10:00 Room temperature |. 53 Remove, slowly, the white ring of cells at the top of the gradient. 54 5m Spin down cells at 400 x g, Room temperature, 00:05:00. 55 Resuspend cells in $\perp$ 200 $\mu$ L of PBS-0.04% BSA. 56 Count cells using the Cellaca or Luna cell counter.

## Library preparation and sequencing

- Evaluate cell viability and cell counts with acridine orange/propidium iodine (AO/PI) onto the Luna FX7 cell counter (Logos Biosystem).
- Adjust the volume of the cell suspension to 1000 cells/µL in PBS-0.04% BSA.
- For each sample, load 20,000 cells (supplementary table 1) into the Chromium Controller (10X Genomics) and partition into single 10X barcoded droplet according to the Chromium single cell 3' Gel Bead Kit v2 manufacturer's instructions.
  - For a selection of samples that underwent anti-Ter 119 Microbeads isolation protocol, use a modified master mix recipe: Δ 1 μL of 0.1X FastSelect globin mRNA blockers 1X (QIAseq FastSelect Globin Kit, Qiagen GmbH) was introduced at the expense of Δ 1 μL of water according to manufacturer's instructions.
  - **59.2** For a selection of other samples, use a modified GEM-RT incubation on a Veriti 96- well thermal cycler (Thermofisher), as shown below:



- For the single-cell gene expression libraries, use the Chromium Single Cell 3' Reagent kit v2 (10X Genomics) according to manufacturer's instructions.
  - 60.1 cDNAs recovered are amplified on Veriti 96-Well Thermal Cycler, the cDNAs products are amplified, cleaned up, quality controlled and quantified.
- Add Illumina P5, P7, Read2 primers and Sample Index to the cDNAs products to generate sequencing libraries.
- Perform quality control and quantify using a quantitative PCR (KAPA Biosystems Library Quantification Kit for Illumina platforms).
- 63 Load the sequencing libraries on a NextSeq2000 P3 flow cell with a sequencing depth of 50,000 reads per cell.

