



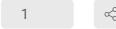


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## S Human Fallopian Tube and Ovary Dissociation for Single Cell RNA-Seq

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This protocol provides a procedure for human fallopian tube and ovary dissociation into single cell suspension prior to single cell RNA-sequencing. It involves removing the epithelial fraction and then further digesting the remaining solid tissue fraction to gain maximum yield from all tissue compartments.

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Single Cell, Ovary, Fallopian Tube

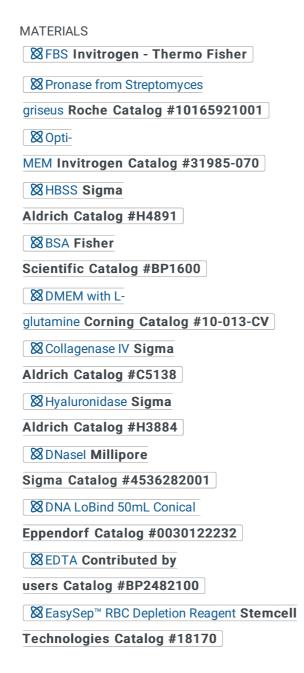
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Apr 30, 2020

Feb 14, 2022

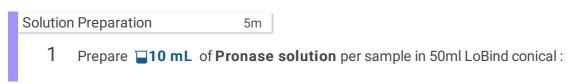
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Prepare DMEM/10%FBS stock
Prepare 0.5M EDTA stock
Prepare 1mL PBS/0.05% BSA per sample

Note: Our samples are retrieved from the OR immediately upon removal from patient and transported to pathology. The pathologist grossly examines the sample and provides 100-200mg transverse tissue segments from the isthmus, ampulla and fimbriae of the fallopian tube and a longitudinal segment from the center of the ovary. Tissue segments are transported to the lab in DMEM/10% FBS.



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Reagent	Quantity
Opti-MEM	10 ml
Pronase (~7 U/mg)	18 mg

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Karst AM, Drapkin R (2012). Primary culture and immortalization of human fallopian tube secretory epithelial cells.. Nature protocols. https://doi.org/10.1038/nprot.2012.097

Prepare **10 mL** of **digestion buffer** per sample in a 50 mL LoBind conical:

Reagent	Quantity
HBSS	10 ml
Collagenase IV (>120 U/mg)	15 mg
Hyaluronidase (750-3000 U/mg)	10 mg
DNasel (20,000 U/ml)	10 μΙ

2.1 Note: Keep all solutions at § 37 °C throughout protocol.

Dissociation 1h 30m

- 3 Rinse gross blood off samples with DMEM/10% FBS.
  - 3.1 Fallopian tube isthmus: bivalve
  - 3.2 Fallopian tube ampulla: bivalve
  - 3.3 Fallopian tube fimbriae: leave whole

3.4 Ovary: mince  $\square 100 \text{ mg}$  tissue into 1-2 mm pieces.

4 Place each sample into 10 mL of Pronase solution. Place in § 37 °C orbital shaker at 200 rpm for © 00:30:00.

## 5 /

Using a 10 ml pipette, triturate tissue to remove loose *epithelial fraction*. Pass all of the pronase solution through 70  $\mu$ m filter into new 50 ml LoBind conical and rinse filter with  $\blacksquare 10$  mL DMEM/10%FBS. Recover solid fraction from original conical and/or off of filter.

NOTE: You now have an *epithelial fraction* and a *solid fraction*. You will continue to process the *epithelial fraction* and *solid fraction* separately in the next steps. Both contain potentially unique cell types.

- 6 Place remaining *solid fraction* in **digestion buffer**. Place on § 37 °C orbital shaker at 200 rpm for © 00:30:00 to © 00:45:00
- 7 Spin down *epithelial fraction* at 400 rcf for **© 00:04:00**. Aspirate and discard supernatant. Resuspend in **3 mL** DMEM/FBS and wait for solid tissue fraction to complete digestion.
- 8 Pass *solid fraction* solution through a 70 μm filter into conical containing *epithelial fraction* and rinse filter with **10 mL** DMEM/FBS.
- 9 Spin down at 400 rcf for **© 00:04:00** . Aspirate and discard supernatant. Re-suspend in **1 mL** PBS/0.05% BSA

## RBC Clean-up 10m

- 10 Move solution to Eppendorf tube and add 12 μL [M]0.5 Molarity (M) EDTA
- Vortex the EasySep RBC depletion reagent bottle for © 00:00:30 to re-suspend

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- 12 Add  $\mathbf{25} \, \mu \mathbf{L}$  reagent and gentle vortex for  $\mathbf{00:00:03}$  until solution appears homogeneous
- 13 Place on magnet and incubate © 00:05:00

5m

- 14 Leaving on magnet, pipet off clear solution
- 15 Re-suspend in pre-warmed § 37 °C DMEM/10%FBS.
- 16 Assess viabilty with trypan blue if proceeding immediately to single cell library creation or method of choice if optimizing protocol.