

Validation of Single-cell Properties and Conservation of Cell Proportions



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Early Access Program, Brochure - v1.Dec2



Capturing Rich Cell Diversity in Cell Heterogenous Samples

6-month Brain Organoids (Shallow Sequencing, 10k reads/cell)



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Blood -> PBMC -> 9 libraries (testing: fresh/frozen, operator, blood draw, library prep): 15,151 cells (total)



Pseudo-bulk expression correlations

Cell proportions



Service: 3' scRNA-seq

PBMC isolation (optional) Single-cell library construction Library sequencing Data processing: cell calling, expression matrix

Data analysis: cell annotations/proportions (optional)

Samples accepted for 3' scRNA-seg analysis:

PBMCs (fresh or cryopreserved): 1 M cells per sample Peripheral blood (EDTA-treated collection tubes*): 1-5 mL * Free tests available with other collection tubes Lymphoprep™ option, SepMate™ option Other options: Bone marrow mononuclear cells (BMMNC)

RBC-depleted blood (free tests available)

Lead time:

1-2 weeks from date of sample arrival at UST

Deliverables:

Fasta files

QC/single-cell report (common metrics) Cell matrix (raw or filtered) in .csv or .h5 formats Further data processing, figures for paper (optional)

Contact us to learn more about the Early Access Program or schedule your service: service@universalsequencing.com

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