



Feb 28, 2022

HuBMAP | GE/URMC/SCH Cell DIVE™ Modality Overview

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dx.doi.org/10.17504/protocols.io.b4kgqutw

GE Research



Lisa Lowery

This is an overview of all protocols currently in use for the GE/URMC/SCH Cell DIVE collaboration for the Human BioMolecular Atlas Program (HuBMAP). It includes links to each of the individual protocols that make up this project workflow.

DOI

dx.doi.org/10.17504/protocols.io.b4kgqutw

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Cory Poole 2022. HuBMAP | GE/URMC/SCH Cell DIVE™ Modality Overview.

protocols.io

<https://dx.doi.org/10.17504/protocols.io.b4kgqutw>



protocol ,

Feb 02, 2022

Feb 28, 2022

Feb 02, 2022



Liz McDonough

GE Research

Feb 28, 2022



Lisa Lowery

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Confirm donor acceptance criteria for inclusion.

1

[URMC | Donor Acceptance Criteria for GE/URMC HuBMAP Inclusion](#)
[URMC | COVID+ Donor Acceptance Criteria for GE/URMC HuBMAP Inclusion](#)

2

Prepare paraffin blocks and FFPE sections from tissue samples.

[URMC | Whole Lung and Lobe Processing - Formalin Fixation and Gross Sectioning of Tissue Samples](#)

[URMC | Paraffin Embedding and Sectioning of FFPE Lung Tissue](#)

3

Deparaffinize and rehydrate slides.

[Cell DIVE™ Platform | Slide Clearing and Antigen Retrieval](#)

4

Characterize antibodies (primary/secondary, direct conjugates, and zenon labelled antibodies) and determine any antigen effects from the Cell DIVE dye inactivation process.

[Cell DIVE™ Platform | Antibody Characterization for Multiplexing](#)

[Cell DIVE™ Platform | Antibody Staining & Imaging](#)

5

Prepare direct conjugates for study.

[Cell DIVE™ Platform | Antibody Purification Chemistry](#)

[Cell DIVE™ Platform | Ab Conjugation: Initial Conjugation & Scale up Conjugation](#)

6

Perform Cell DIVE™ multiplexed data acquisition on the final cohort.

Staining is done using the Leica Bond MAX and images are acquired on the IN Cell Analyzer 2200.