protocols.io

🌐 Imaging Mass Cytometry Modality Overview V.2

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Version 2 ▼

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Human BioMolecular Atlas Program (HuBMAP) Method Development Community
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1   Process donor organs into samples for analysis.
Lymph Node: dx.doi.org/10.17504/protocols.io.bbgnijvehttps://drive.google.com/drive/my-drive
Thymus: dx.doi.org/10.17504/protocols.io.bbgmiju6
Spleen: dx.doi.org/10.17504/protocols.io.bc3kiykw

2   Register organ donor, organs received, and common coordinate region information in the HuBMAP UUID generator at
https://uuid.hubmapconsortium.org

3   Prepare paraffin blocks from tissue samples.
Cassette Processing: dx.doi.org/10.17504/protocols.io.8nzhvf6
Paraffin Embedding: dx.doi.org/10.17504/protocols.io.bam9ic96

4   Package and send paraffin blocks to the Bodenmiller lab: dx.doi.org/10.17504/protocols.io.bf5njq5e

5   Conjugate primary antibodies using the Maxpar Antibody Labeling Kit: dx.doi.org/10.17504/protocols.io.bf5jjq4n

6   Cut and mount paraffin sections for staining and analysis.
Sectioning: dx.doi.org/10.17504/protocols.io.bfz4jp8w

7   Antibody staining of paraffin sections for Imaging Mass Cytometry: dx.doi.org/10.17504/protocols.io.bfz8jp9w

8   Prepare a slide to use for compensation: dx.doi.org/10.17504/protocols.io.bf2djqa6

9   Tune the Image Mass Cytometer for acquisition: dx.doi.org/10.17504/protocols.io.bf2gjqbw

10   Data Acquisition by Image Mass Cytometry: dx.doi.org/10.17504/protocols.io.bf2ijqce

11   Compensation Slide Acquistion: dx.doi.org/10.17504/protocols.io.bf2jjqcn

12   Image Processing and Segmentation: https://github.com/BodenmillerGroup/ImcSegmentationPipeline

13   Load Imaging Mass Cytometry Datasets onto Hipergator:
☐ **IMC_Transfer Data Zurich to HPG.docx**

14   Prepare and upload data to the HuBMAP HIVE Globus landing pad using Hipergator.
☐ **IMC Transfer data from Hipergator to HuBMAP HIVE Globus.docx**