Mapping the Human Body at Single-Cell Resolution: Developing the Human Reference Atlas

Rachel Bajema, Indiana University- Bloomington Heidi Schlehlein, Indiana University- Bloomington



Who are medical illustrators?



STAMP

Encapsulated droplets

2. Cell lysis

RNA hybridiza-

Cell 2

Cell 3

4. PCR STAMPs used

as template

m PNIA

5. Sequencing

and analysis

Cells

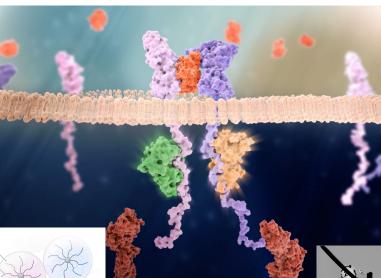
⇒ 🔘 0

1. Microparticle

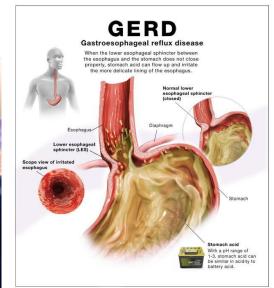
and lysis buffer

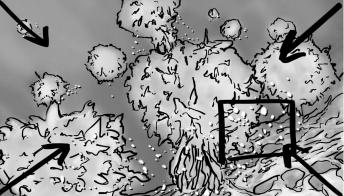
3. Droplets broken

Reverse transcription









What is HuBMAP (The Human BioMolecular Atlas Program)?

- Creating a computable, open-source map of the human body at single-cell resolution
- 18 funded components with over 350 researchers in more than 50 research institutions across the U.S. and Europe
- Initiatives:









Funded by the NIH Common Fund HuBMAP Integration, Visualization and Engagement: Building an atlas tissue maps; tools for visualizing, searching and modeling data; Infrastructure, Engagement and Communications

Indiana University Bloomington > Luddy School of Informatics, Computing, and Engineering > Cyberinfrastructure for Network Science Center (CNS) Rapid Technology Implementation:
Enhancing, large-scale validation, integration of emerging new technologies

3. Tissue Mapping Centers: Collect and analyze a range of normal tissues

 Transformative Technology
Development: Analyze tissue, validation of new methods for mapping the body at high resolution

Tissue Mapping Centers (TMC)

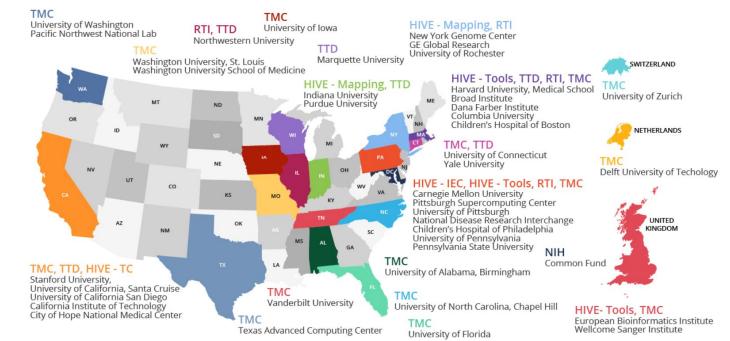
Where data comes from

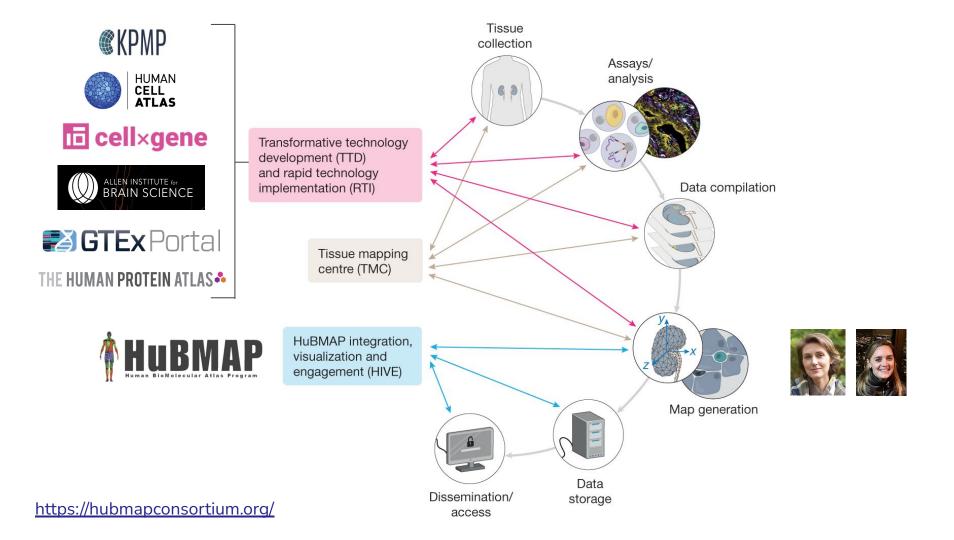
Over 20 different organs from more than 80 donors comprise the samples and datasets (>1000 of each) across 3 main modalities:

- mass spectrometry
- microscopy
- sequencing

New Assay types are published as they become available.

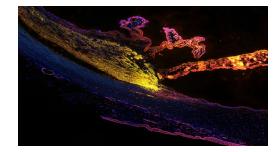
HuBMAP Contributing Sites



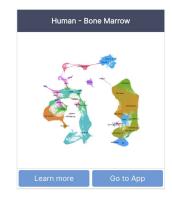


Why another Human Atlas? Why Single-Cell?

- Our goal is multiscale, multidimensional mapping of the human body from the whole body level to the individual cell
- Now possible to classify cells by their expression profiles (level at which RNA or protein is expressed from each gene)
- Large scale data sets can be interpreted by machine learning and visualization tools to find how cells are related to and interact with each other
- Enables the creation of cellular reference maps of the position, function and characteristics of every cell type in the human body







Registration User Interface (RUI)

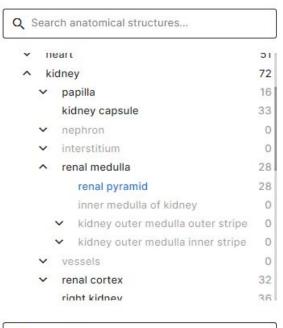
A new way to preserve spatial relationships between different tissue samples from around the world into searchable one 3D space

HuBN	AP CC																					
Blood Vasculatu	Brain Brain	G- C		Fallopian Tube, R	<mark>б</mark> Heart Кі	Ney, L Kidn	1	Knee, R	Large Intestine	Liver Lung		QVary, L	Dvary, R	(r) Pancreas	W. Petvis	Prostate	Skin	Small Intestine	S pleen	ti) Thymus		0
O Lett	O Right	Anterior	O Poster	rior Regi	ster 💽 3	D Preview		THE SEL			P					•				۵ ۲: 43 ۲: 48 ۲: 48	Tissue Block Dimensions (mm) With 00 Height 10 Dieft 20 Tissue Sections Tissue Sections # Sections Tissue Block Rotation # X • Y • Z • Anatomical Structure Tags Add Anatomical Structures • Assigned • Added	20 5 - 0 - 0 - 0 - 0 - 0 - 0 +

Exploration User Interface (EUI)

A way to explore the vast amounts of data entered into the RUI

Anyone can access this data



Q Search cell types...

3D Reference Organ Library

HuB

Glb and fbx files of all 25 organs centrally located

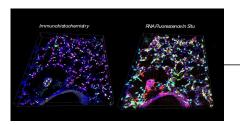
- Free for all to use under Creative Commons license,
- Created for RUI
- Each organ has metadata related to Uberon terms that link to the ASCT+B Table



	aon or us		es via collisi ose releas e				2nd Relea	se, Decemi	ber '21		
		ono				sjoots.					
9	*	•		\mathbf{O}	R	6	6	7	5	F	1
Allen Brain	Blood Vasculatu		Marrow & d/Pelvis	Eye	Fallopian Tube	Heart	Kidney	Knee	Large Intestine	Liver	Lungs
¥	9	-	3	5	_6_	۲	æ	•		*	
ymph Node	Ovary	Pancreas	Placenta	¥ Prostate	Skin	Small Intestine	Spinal	Spleen	Thymus	All Org	ans
NODE						intestine	Cord				_
	¥	$\widehat{\mathbf{V}}$									
Ureter	Urinary Bladder	Uterus									
	Diddaei										
Lun	gs										
Fen						Male					
3D m	odel					3D mod	lel				
		100					2	ł.			
		V.						1			
		-(X			
	1	0							λ		
	(R						/			

https://hubmapconsortium.github.io/ccf/pages/ccf-3d-reference-library.html

Anatomical Structures, Cell Types and Biomarkers (ASCT+B) tables



Data from assays/analysis indicates which anatomical structures, cell types and biomarkers are present in a tissue sample



Connected to cell ontology: standard set of research-confirmed cell types

• Forces agreement on terminology and interpretation of data

https://hubmapconsortium.github.io/ccf-asct-reporter/

- Text output (.csv) defines organizational hierarchy (relationships) for each organ
 - Organ
 - Anatomical structures
 - Cell types
 - Biomarkers
- Used to define functional tissue units (FTUs) for each organ
- FTUs are related to vascular mapping, a concurrent effort
- FTUs chosen for illustration

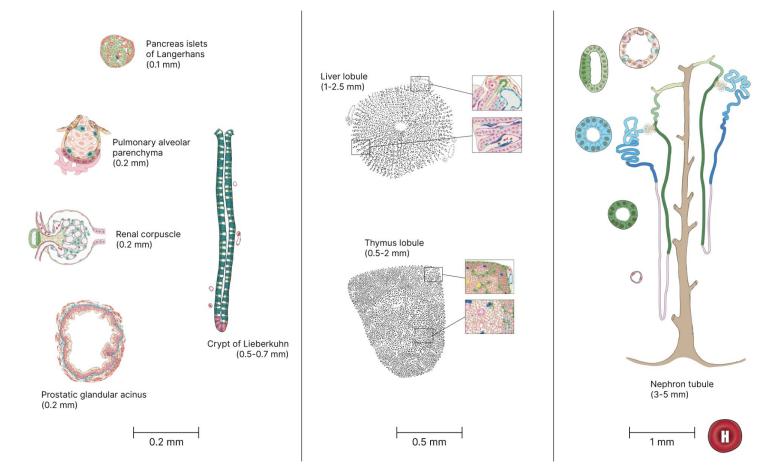
Partial ASCT+B table for Large Intestine

AS/4/LABEL	AS/4/ID	AS/4/NOTES	CT/1	CT/1/LABEL	CT/1/ID
colonic epithelium	UBERON:0000397		absorptive	enterocyte of epithelium of large intestine	CL:0002071
colonic epithelium	UBERON:0000397		goblet	large intestine crypt goblet cell	CL:1000321
epithelium of crypt of Lieberkuhn	UBERON:0011184		epithelial stem cell	intestinal crypt stem cell of large intestine	CL:0009016
epithelium of crypt of Lieberkuhn	UBERON:0011184		Paneth	paneth cell of epithelium of large intestine	CL:0009009
epithelium of crypt of Lieberkuhn	UBERON:0011184		neuroendocrine	neuroendocrine cell	CL:0000165
epithelium of crypt of Lieberkuhn	UBERON:0011184		absorptive	enterocyte of epithelium of large intestine	CL:0002071
pithelium of crypt of Lieberkuhn	UBERON:0011184		goblet	large intestine crypt goblet cell	CL:1000321
pithelium of crypt of Lieberkuhn	UBERON:0011184		transient amplifying cell	transit amplifying cell of large intestine	CL:0009011
pithelium of crypt of Lieberkuhn	UBERON:0011184		tuft cell	intestinal tuft cell	CL:0019032
olonic epithelium	UBERON:0000397	Tissue resident lymphocytes; interepithelial lymphocytes	surface intraepithelial CD8+ alph alpha positive alpha/beta lymp	h CD8-alpha-alpha-positive, alpha-beta intraepithe	li; CL:0000915
olonic epithelium	UBERON:0000397	Tissue resident lymphocytes; interepithelial lymphocytes	surface intraepithelial CD8+ lymphocyte gamma/delta		CL:0008364
colonic epithelium	UBERON:0000397	Tissue resident lymphocytes; interepithelial lymphocytes	surface intraepithelial CD8+ alph beta positive alpha/beta lymph	o CD8-alpha-beta-positive, alpha-beta intraepitheli	a CL:0000796
olonic epithelium	UBERON:0000397	Tissue resident lymphocytes; interepithelial lymphocytes	surface intraepithelial CD4+ lymphocyte	CD4-positive, alpha-beta intraepithelial T cell	CL:0000793
olonic epithelium	UBERON:0000397	Tissue resident lymphocytes; interepithelial lymphocytes	surface intraepithelial CD4-CD8- lymphocyte alpha/beta	CD4-negative, CD8-negative, alpha-beta intraepit	h CL:0000935
rypt of Lieberkuhn of colon	UBERON:0013485		eosinophil	mature eosinophil	CL:0000041
amina propria of mucosa of colon	UBERON:0007177		subepithelial membrane		
amina propria of mucosa of colon	UBERON:0007177		pericryptal fibroblastic sheath		
amina propria of mucosa of colon	UBERON:0007177		capillary endothelium	capillary endothelial cell	CL:0002144
amina propria of mucosa of colon	UBERON:0007177		lymphatic endothelium	endothelial cell of lymphatic vessel	CL:0002138
amina propria of mucosa of colon	UBERON:0007177		pericyte	pericyte cell	CL:0000669
amina propria of mucosa of colon	UBERON:0007177		myofibroblast	myofibroblast cell	CL:0000186
amina propria of mucosa of colon	UBERON:0007177		fibroblast	fibroblast	CL:0000057
amina propria of mucosa of colon	UBERON:0007177		nerve/schwann cell	Schwann cell	CL:0002573
amina propria of mucosa of colon	UBERON:0007177		ganglion		
amina propria of mucosa of colon	UBERON:0007177		neuroendocrine cell	neuroendocrine cell	CL:0000165
ut-associated lymphoid tissue	UBERON:0001962		M cell	M cell of gut	CL:0000682
ut-associated lymphoid tissue	UBERON:0001962		B cell	lymphocyte of large intestine lamina propria	CL:0009018
ut-associated lymphoid tissue	UBERON:0001962		CD4+ T cell	lymphocyte of large intestine lamina propria	CL:0009018
ut-associated lymphoid tissue	UBERON:0001962		regulatory CD4+ T cell	lymphocyte of large intestine lamina propria	CL:0009018
ut-associated lymphoid tissue	UBERON:0001962		CD8+ T cell	macrophage	CL:0009018
gut-associated lymphoid tissue	UBERON:0001962		macrophage	macrophage	CL:0000235
gut-associated lymphoid tissue	UBERON:0001962		NK cell	lymphocyte of large intestine lamina propria	CL:0009018
gut-associated lymphoid tissue	UBERON:0001962		ILF lymphatic endothelium	endothelial cell of lymphatic vessel	CL:0002138

Currently more than 25,000 nodes and edges in the ASCT+B knowledge graph

https://docs.google.com/spreadsheets/d/1d_KWKnQq3HT5nzDmfhlvFG4P_qdviu0vyhGZ6QHqNIk/edit#gid=2043181688

2D Functional Tissue Unit (FTU) Illustrations



HuBMAP CCF Portal: where it all lives

Explore and use

Everything is open source- all recreatable, can be used for any purpose



https://hubmapconsortium.github.io/ccf/index.html



The Human Body Atlas: High-Resolution, Functional Mapping of Voxel, Vector, and Meta Datasets

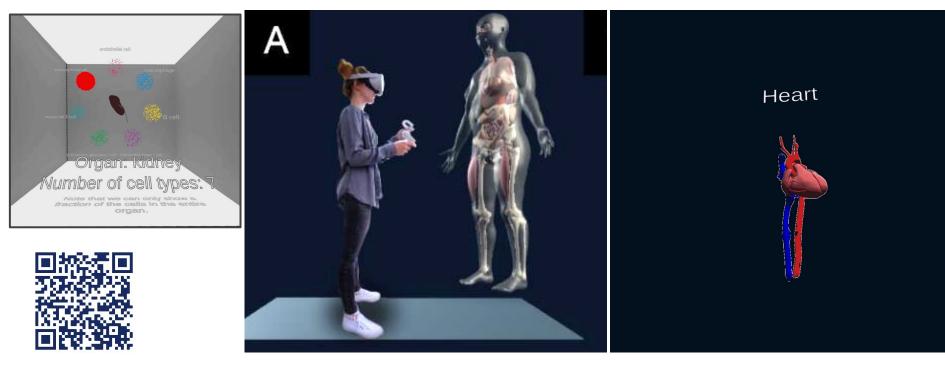
MC-IU team within the HuBMAP HIVE

The ultimate goal of the HIVE Mapping effort is to develop a common coordinate framework (CCF) for the healthy human body. This framework will support cataloging different types of individual cells, understanding the functions of and relationships between those cell types, and modeling their individual and collective function. During the initial three years of HuBMAP, the MC-IU team has built many elements of the CCF. We co-organized the construction of ASCT+B Tables and implemented a CCF Ontology. We collaborated with NIAID at NIH on the design of a 3D Reference Object Library. Lastly, we developed three interactive user interfaces. The CCF ASCT+B Reporter supports the authoring and interactive review of ASCT+B Tables. The CCF Registration User Interface (RUI) supports uniform tissue data registration across organs and labs. The CCF Registration User Interface (RUI) supports of semantically and spatially explicit data —from the whole body to the single cell level. For an introduction to HuBMAP goals, data, and code visit the Visible Human MOOC (VHMOOC).



Visual resource

Organ VR Gallery



https://osf.io/z9gm3/ For test build: contact Andi Bueckle at abueckle@iu.edu

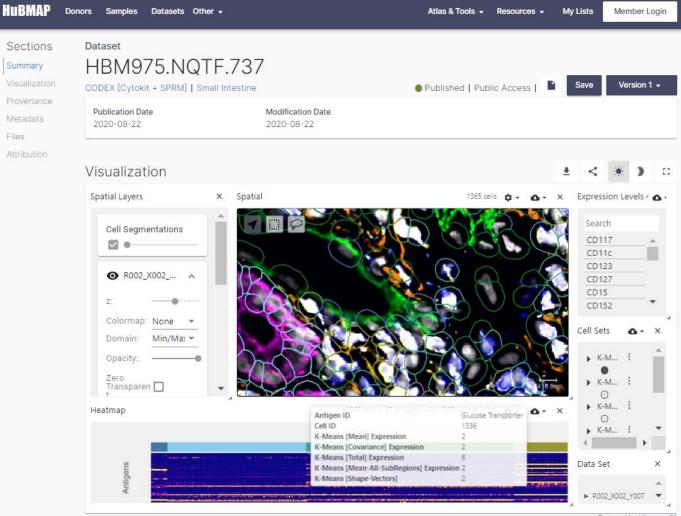


vilessle

Principal Investigator: Nils Gehlenborg

http://vitessce.io/

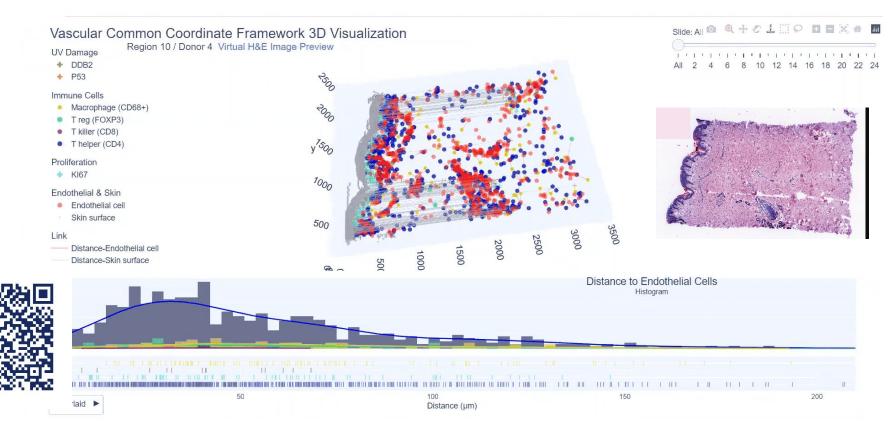




Powered by Vitessce 🗹

Visual resource

Vascular Common Coordinate Framework





- All 3D/2D files are open source, available for download
- Take the free <u>Visible Human MOOC tutorial</u> to learn more
- Explore the <u>HuBMAP Data Portal</u> and <u>CCF Portals</u>, including the visualization tools
- Contribute to and the <u>ASCT+B tables</u> for research on anatomical structures, cell types and biomarkers, and their hierarchical relationships
- Register tissues blocks in the RUI
- Become an SMEs/external reviewers of 3D models and FTUs
 - Expert in an human anatomy organ system
 - Work with medical artist to edit and create 3D organs and FTU illustrations

Contact us

Facebook @cnscenter Instagram @cns_at_iu Twitter @_hubmap



Rachel Bajema - <u>rbajema@iu.edu</u>



Heidi Schlehlein - <u>hschleh@iu.edu</u>

This project has been funded in part with Federal funds from the National Institute of Allergy and Infectious Diseases (NIAID), National Institutes of Health, Department of Health and Human Services under BCBB Support Services Contract **HHSN316201300006W/HHSN27200002** to MSC, Inc.