



11PM

4AM in London (GMT), 1PM in Tokyo (GMT+9)

Multiscale Visualizations

Moderator: Andreas Bueckle, *Indiana University*

Presenters:

- Tobias Isenberg, *National Institute for Research in Digital Science and Technology (Inria)*
- Ludovic Autin, *Scripps Research Institute*
- Griffin Weber, *Harvard Medical School*



Setting the Stage: Building and Exploring the Human Reference Atlas with Virtual Reality



Andreas “Andi” Bueckle, Ph.D.
Research Lead

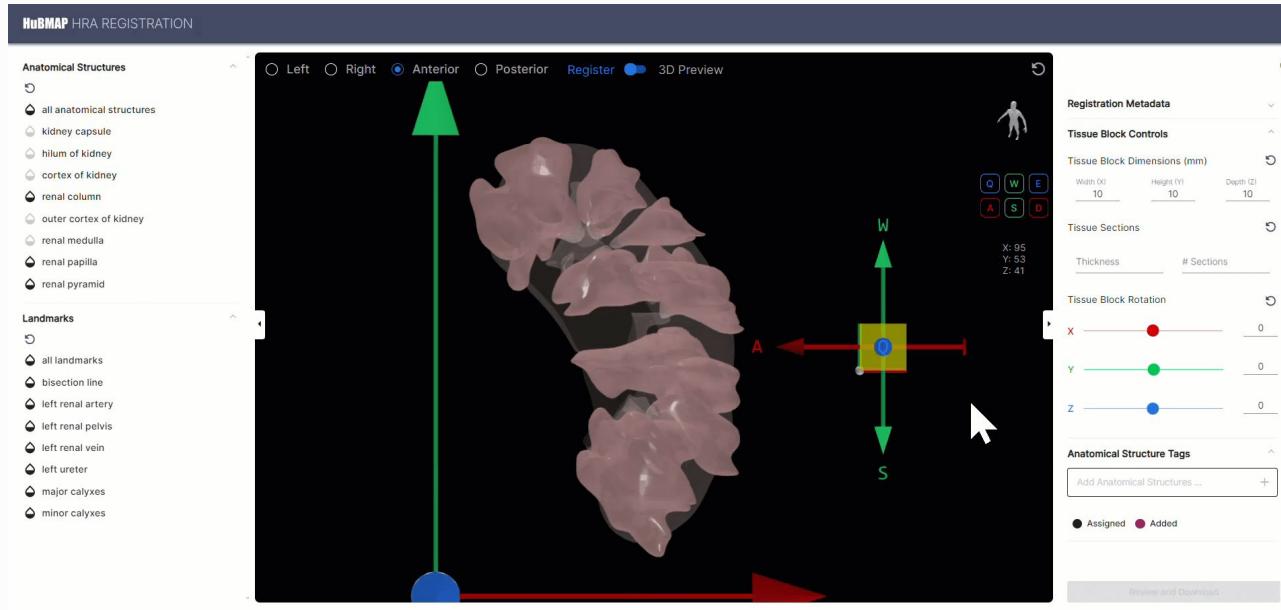
*Cyberinfrastructure for Network Science Center
Department of Intelligent Systems Engineering
Luddy School of Informatics, Computing, and Engineering
Indiana University, Bloomington, IN, USA*

HRA User Interfaces





Registration User Interface (RUI)



<https://apps.humanatlas.io/rui/>



Exploration User Interface (EUI)

HuBMAP HRA EXPLORATION

Sex: Both Age: 1-110 BMI: 13-83

Filter: ✓ AS ✓ CT ✓ B

Anatomical Structures (AS) Tissue Blocks: 730

- ✓ brain 11
- ✓ lymph node 36
- ✓ eye 43
- ✓ fallopian tube 0

Cell Types (CT) Tissue Blocks: 730

- absorptive 67
- absorptive 61
- adipocyte 225
- adipocyte 16

Biomarkers Tissue Blocks: 730

✓ BG ✓ BL ✓ BM ✓ BP ✓ BF

- a smooth muscle actin 0
- A2M 253
- ABC10-43608400015.1 0
- ABCA1 11
- ABCA13 11
- ABCA3 42
- ABCA4 11
- ABCA8 11
- ABCC9 403
- ABCG2 0

Run Spatial Search

body | cell | biomarker

20 Tissue Data Providers
308 Donors
730 Tissue Blocks
939 Tissue Sections
3260 Tissue Datasets

Patient B Cortical biopsy
Entered 4/18/2020, Sett Winfree, KPMP-IUO...

Patient A Cortical biopsy
Biopsy from Nephrology biobank-salvaged fro...

Cover Nephrectomy
Biopsy from Nephrology biobank-salvaged fro...

Female, Age 38
Entered 7/3/2023, Andrea Radtke, NIH

Male
Entered 5/26/2023, Anna Martinez Casals, Isl...

Male, Age 19, Patient APAP1
Entered 8/24/2023, Jordan Portman, UEdinbu...

Female, Age 48, Patient APAP10
Entered 8/24/2023, Jordan Portman, UEdinbu...

Female, Age 23, Patient APAP2
Entered 8/24/2023, Jordan Portman, UEdinbu...

Male, Age 66, Patient APAP3
Entered 8/24/2023, Jordan Portman, UEdinbu...

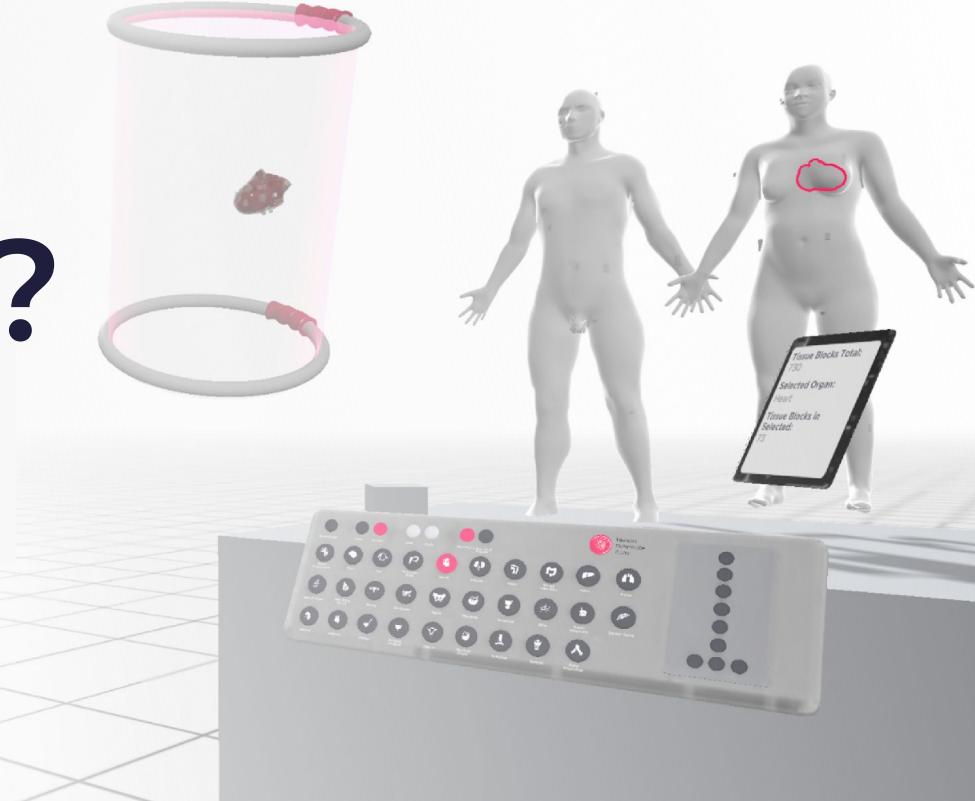
Male, Age 38, Patient APAP4
Entered 8/24/2023, Jordan Portman, UEdinbu...

<https://apps.humanatlas.io/eui/>

HRA Organ Gallery in VR

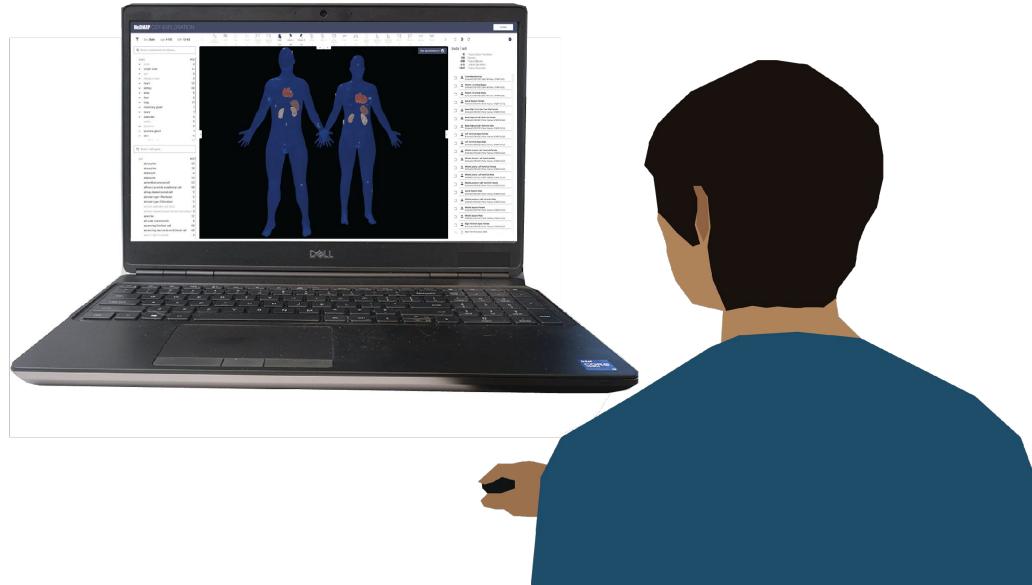


What if...?





What if?



HRA Exploration User Interface: <https://apps.humanatlas.io/eui/>



HRA Organ Gallery in VR



Scan to
download

 Meta

Identified **3 major use cases**:

- Quality Assurance/Quality Control for registered tissue blocks
- Onboarding to the HRA
- Telling Embodied Data Stories

Paper:

<https://doi.org/10.3389/fbinf.2023.1162723>

HRA Organ Gallery in VR:

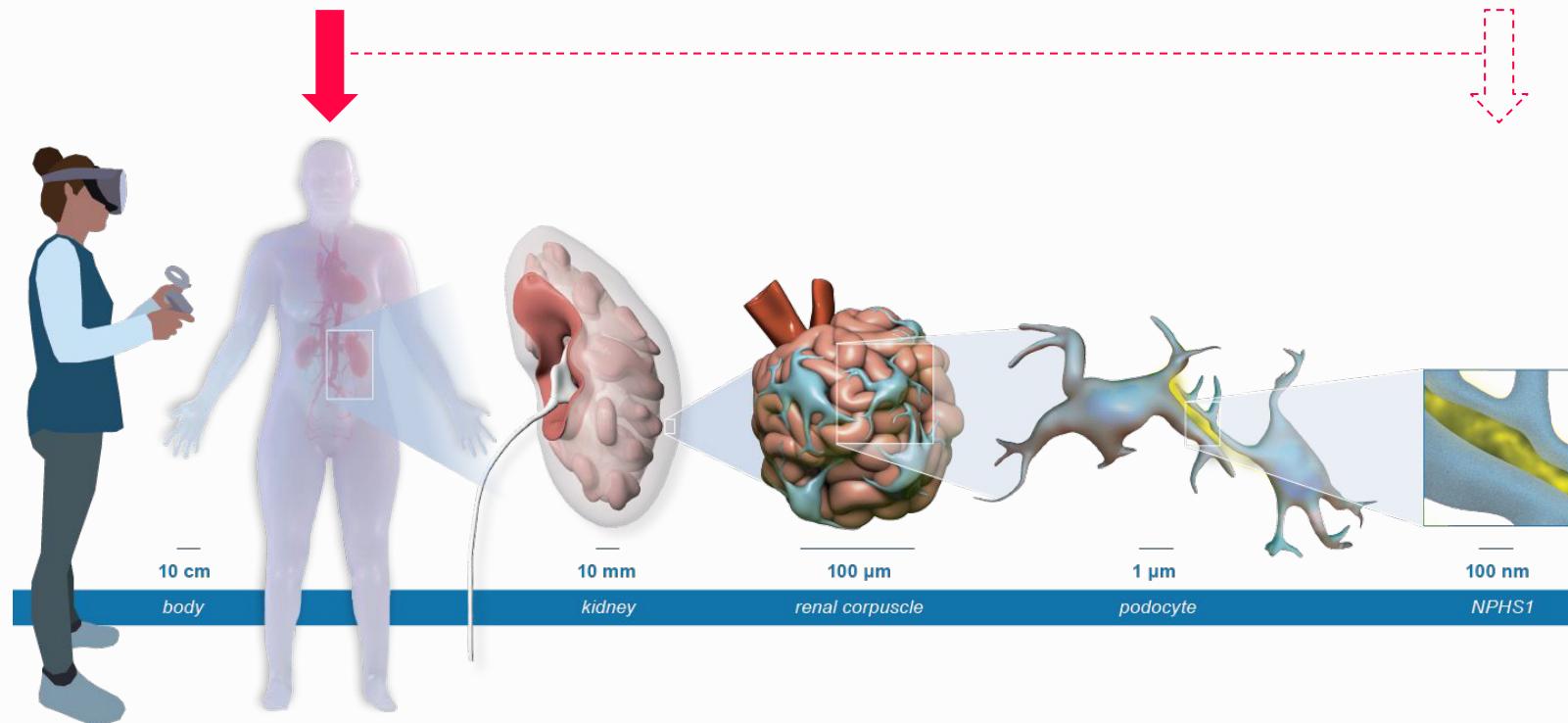
<https://www.meta.com/experiences/5696814507101529>

Exploring Spatial Data in VR



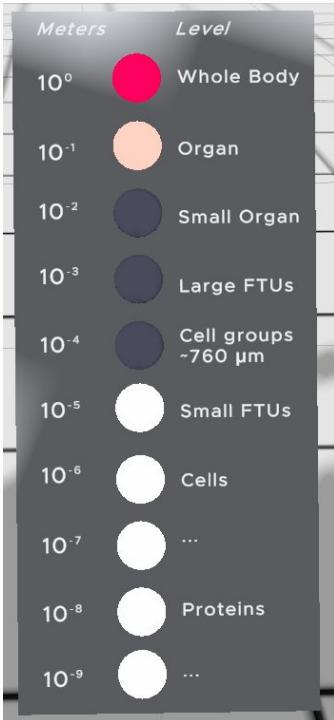


Common Coordinate Framework (CCF)





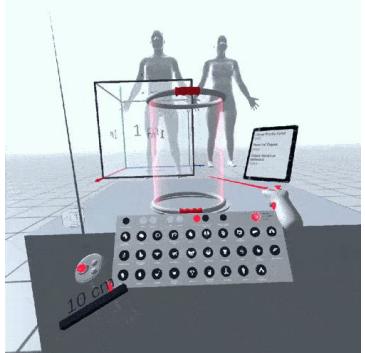
Common Coordinate Framework (CCF)



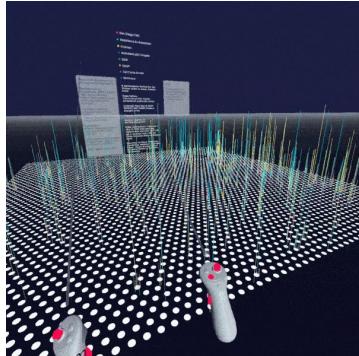
- Start at highest floor
- Take an elevator across levels, from Whole Body to Proteins
- Currently open for business:
 - Whole Body
 - Small Organ
 - Large FTUs
 - Cell Groups (~760 μm)



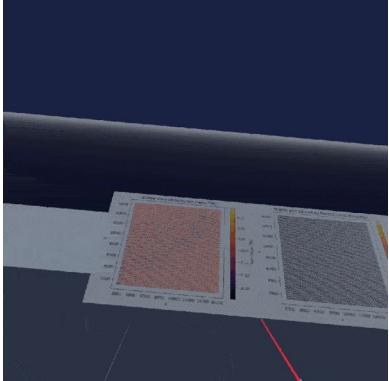
HRA Organ Gallery in VR



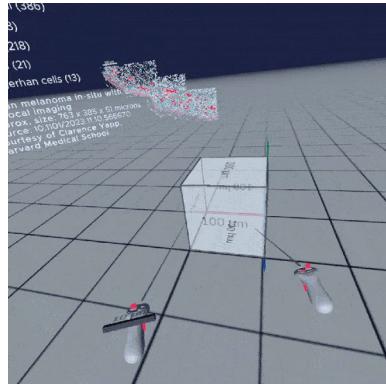
Level 0: Whole Body



Level -3: 3D stepped relief map for
senescence hallmarks in Visium slide



Also Level -3: auxiliary 2D
scatter graphs



Level -4: Cell groups
~760 microns

Multiscale exploration of the HRA

- **Level 0: Whole Body**
- **Level -2: Small Organ**
- **Level -3: Large FTUs**
- **Level -4: Cell Groups ~760 Microns**

Talk to Andreas “Andi”
Bueckle if you would like a
demo. Video:

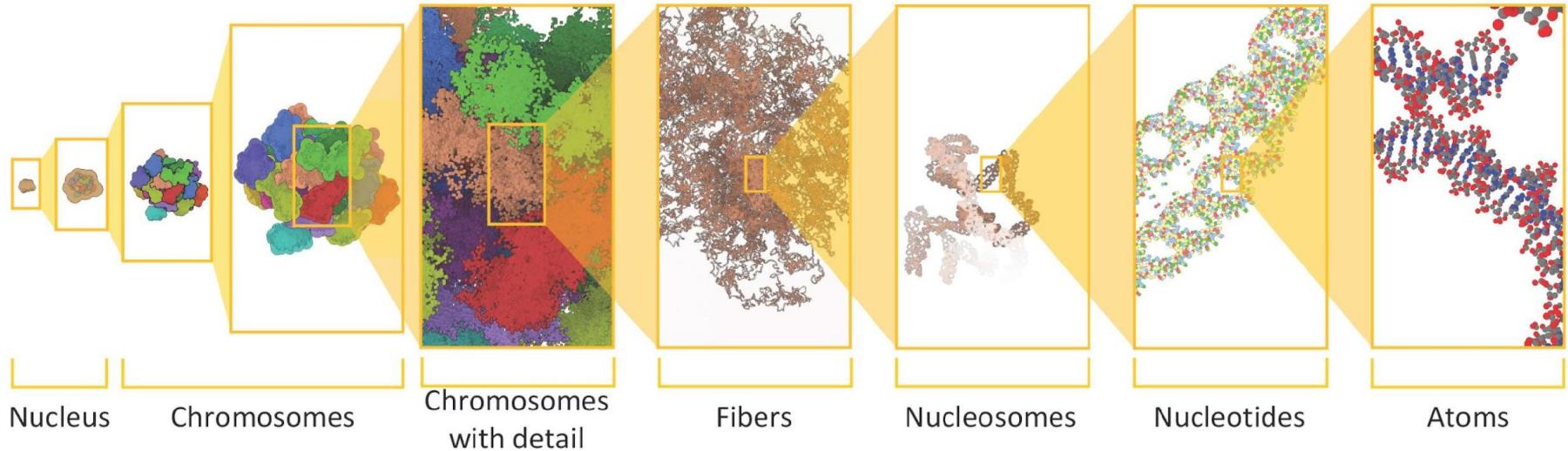
<https://www.youtube.com/watch?v=Wy0BCOFWClk>



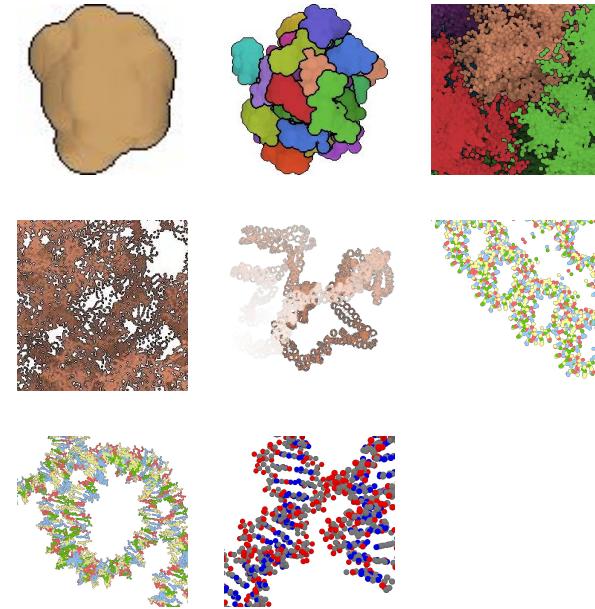
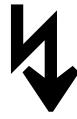
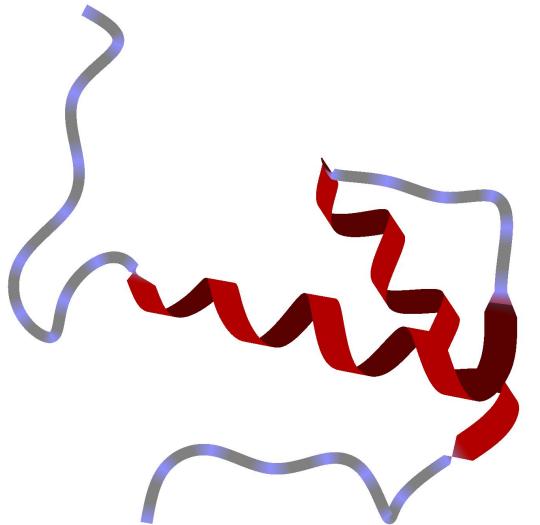
Tobias Isenberg

Inria

ScaleTrotter

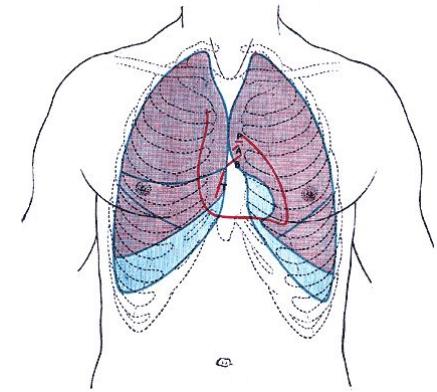
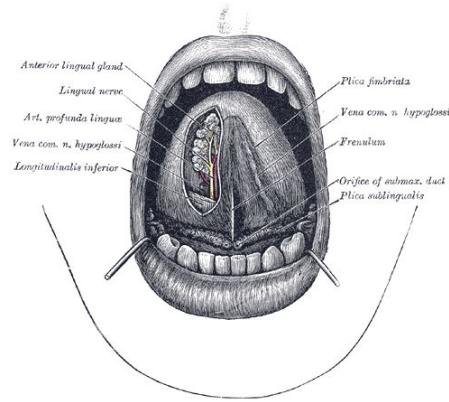
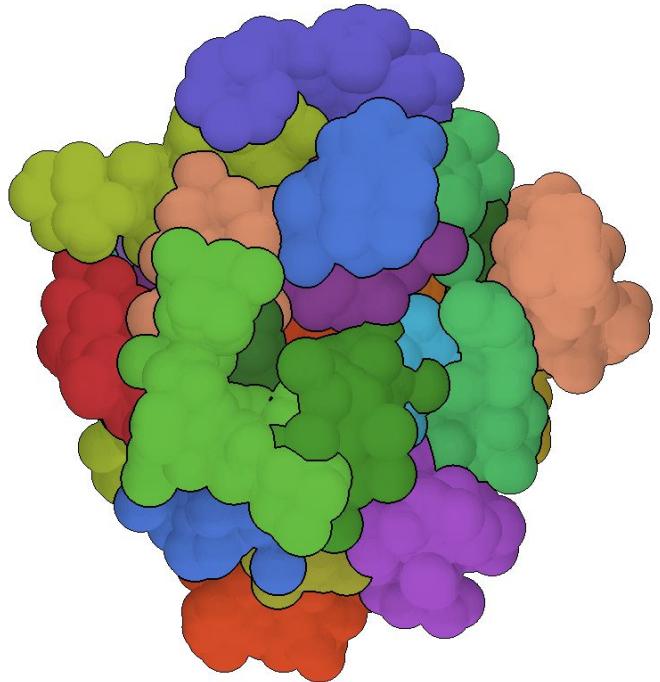


“Simple,” straightforward transitions



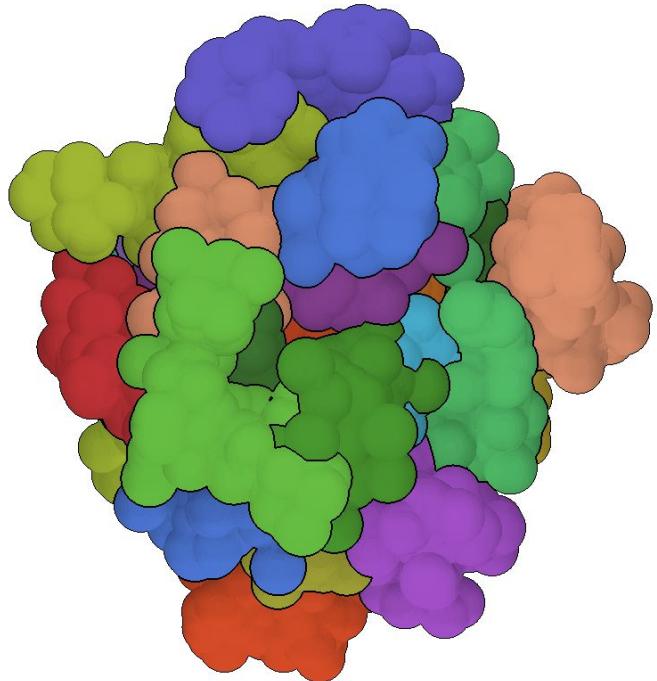
[van der Zwan et al., 2011]

Visual embedding transition



[Gray's Anatomy, 1918]

Visual embedding transition



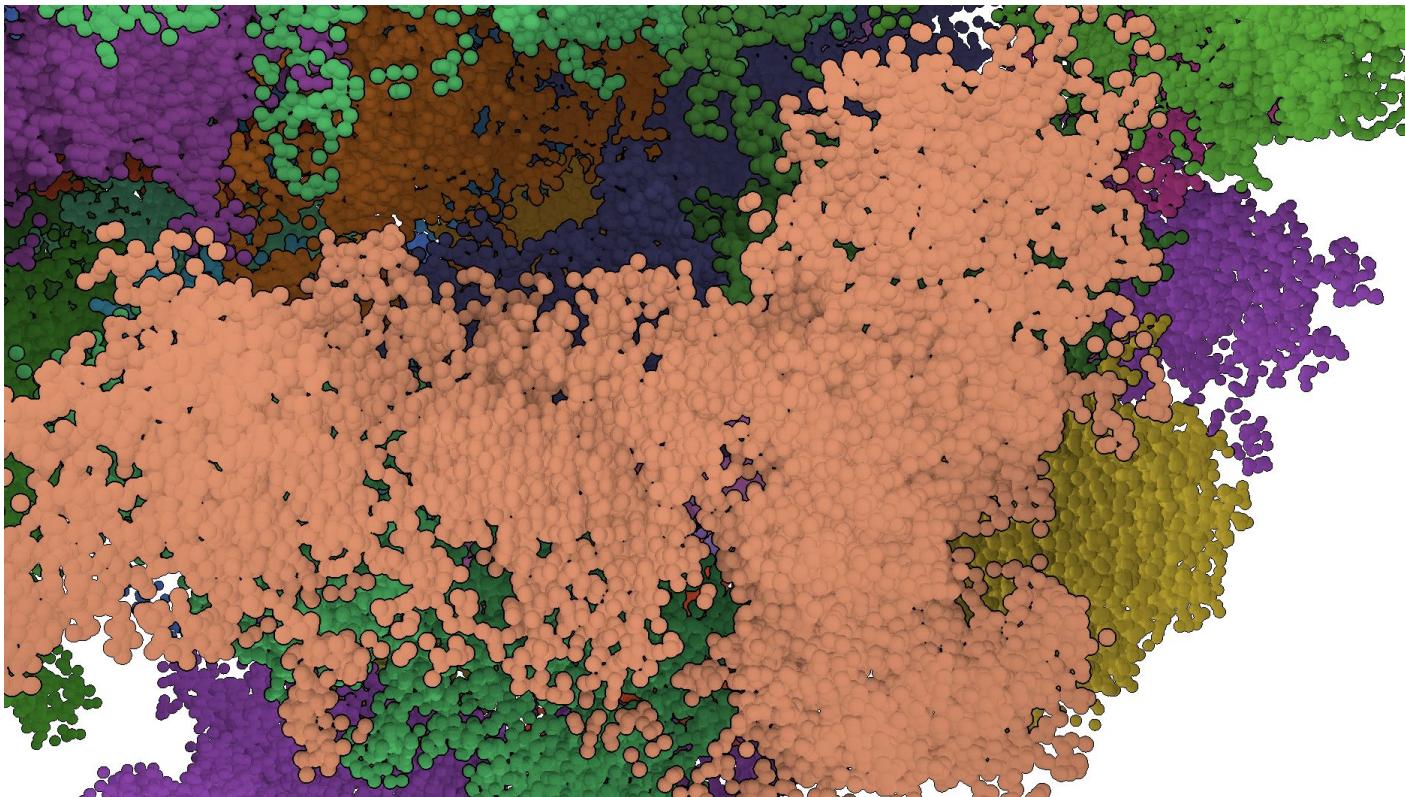
coarse scale in 3D

coarse scale flattened to canvas

detailed scale in 3D on top of canvas

canvas disappears

Visual embedding transition



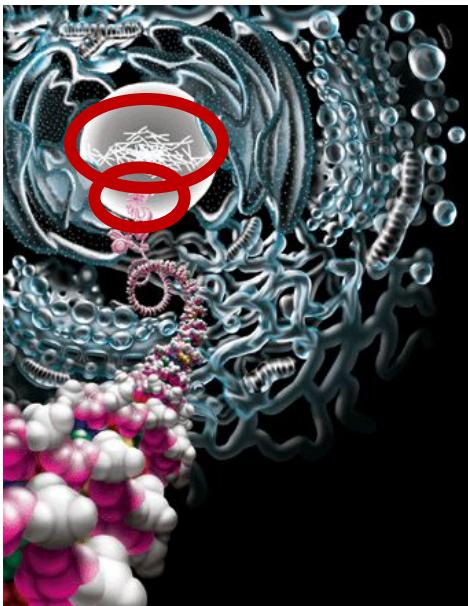
Interactive exploration

ScaleTrotter: Illustrative Visual Travels Across Negative Scales

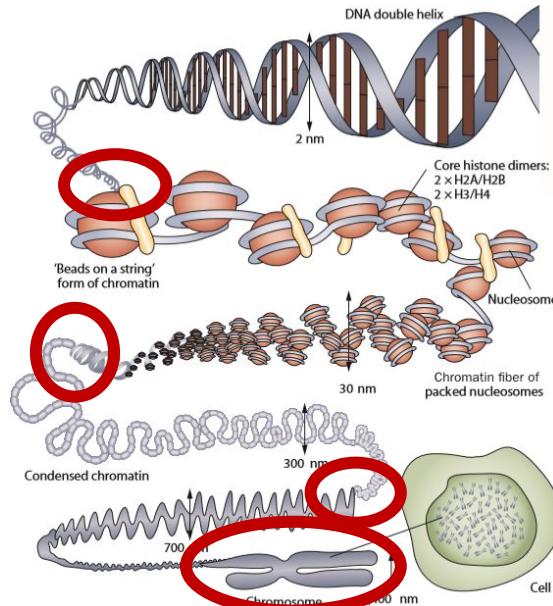
Sarkis Halladjian, Haichao Miao, David Kouřil, M. Eduard Gröller, Ivan Viola, Tobias Isenberg

Inria **université PARIS-SACLAY** **TU WIEN** **VRvis** **Software Cultural Tools Learning platform** **Big Data University of Vienna for Learning**

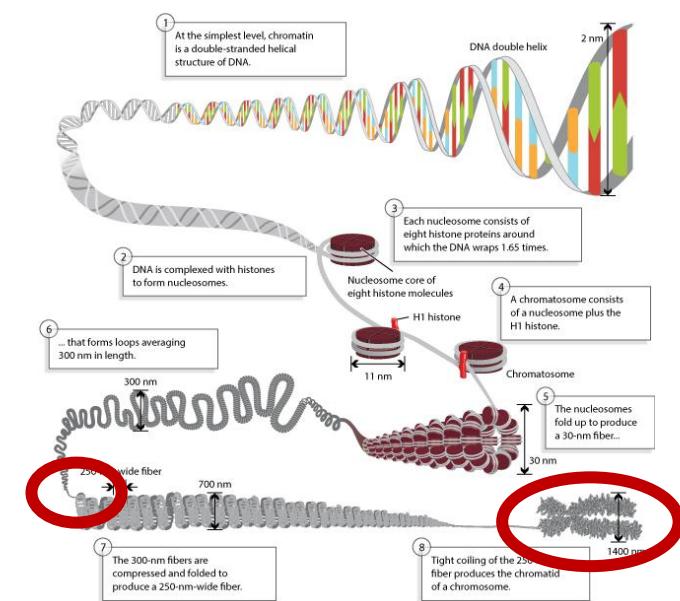
Further Inspiration: Multiscale DNA illustration



[Pennisi, 2001]

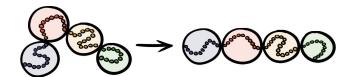


[Tonna et al., 2010]

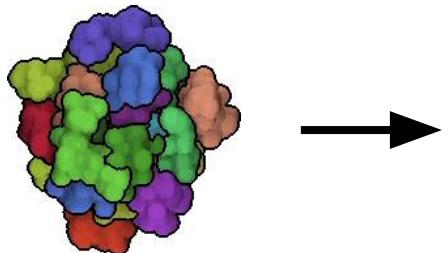


[Nature Education, 2013]

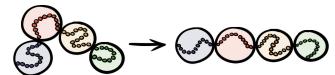
Spatial straightening



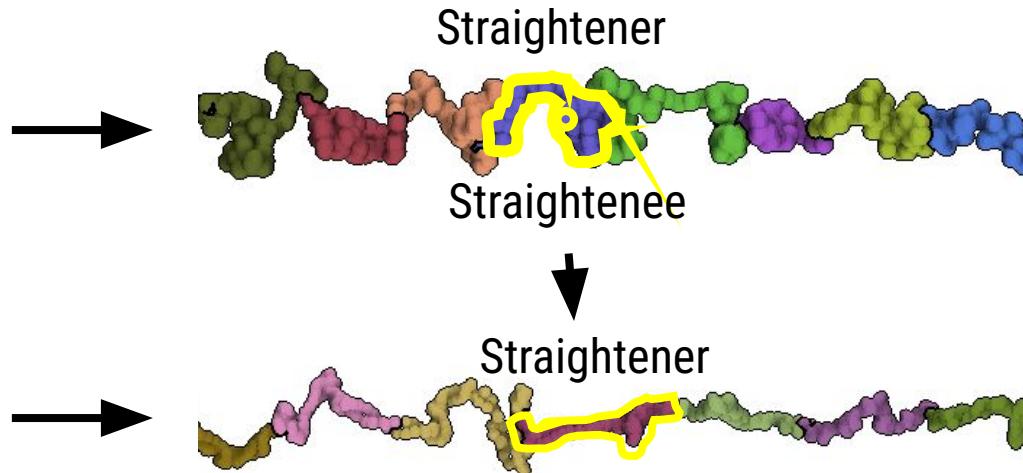
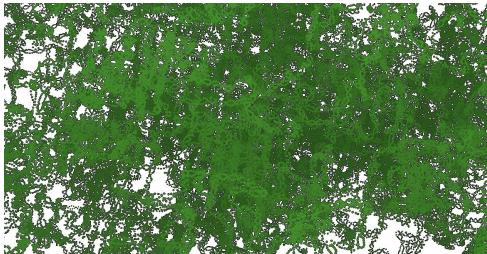
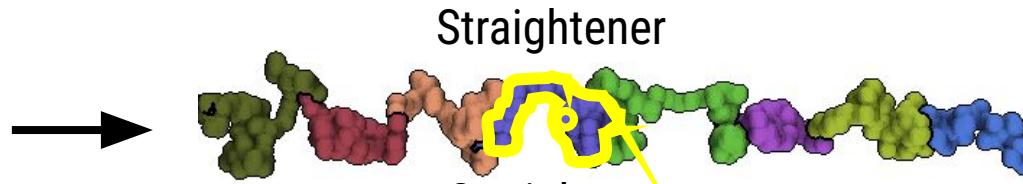
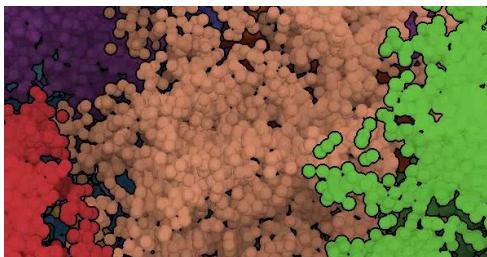
Chromosome scale



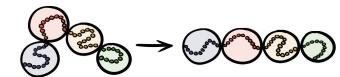
Spatial straightening



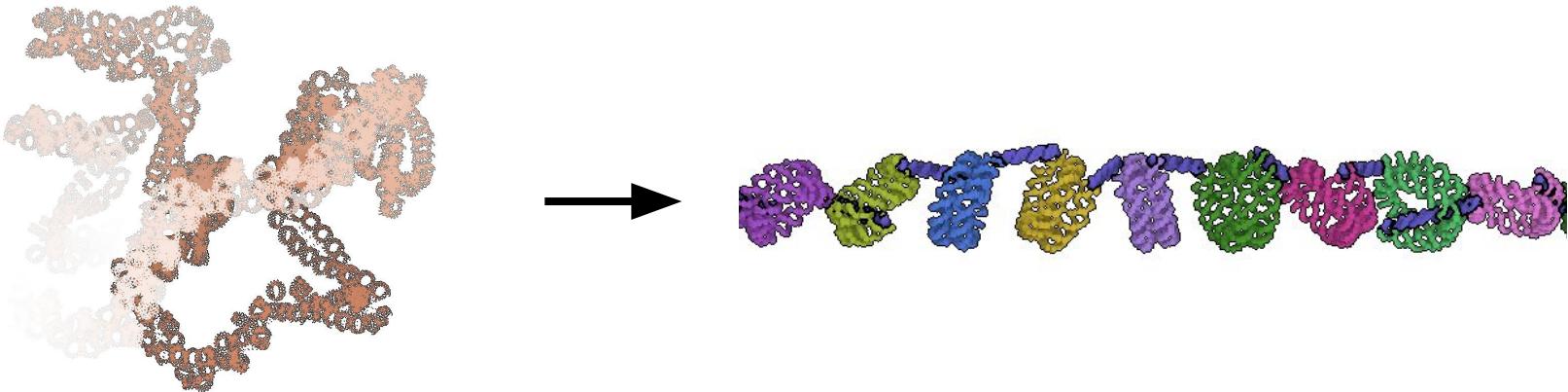
Loci and fiber scales



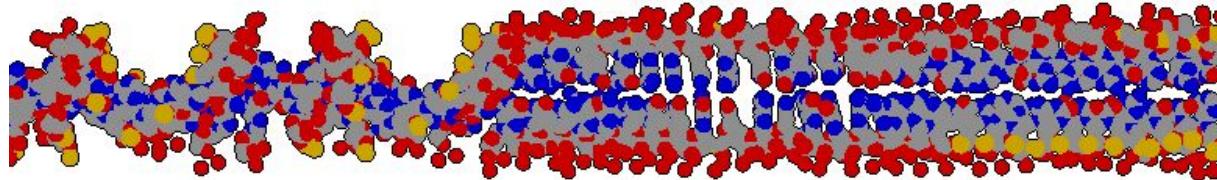
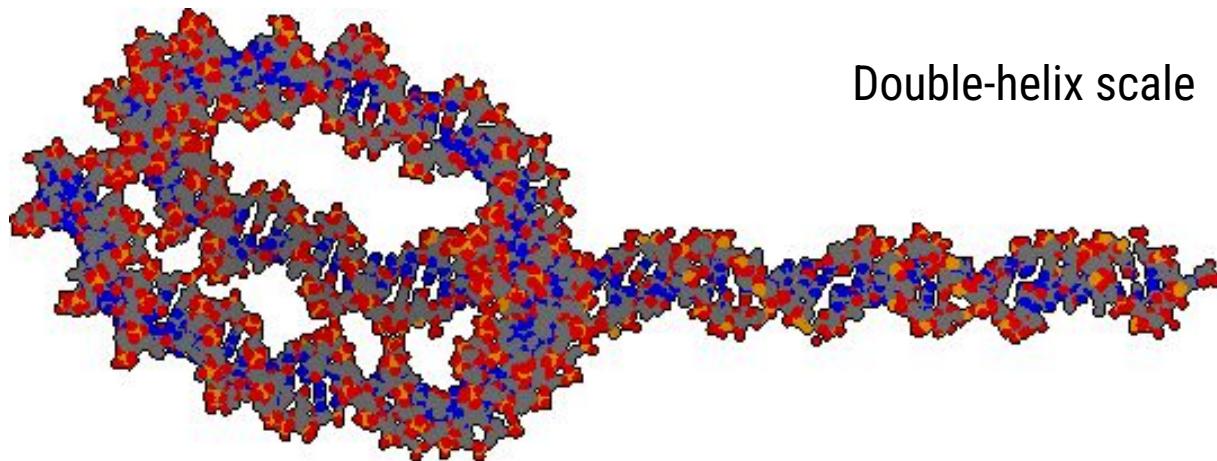
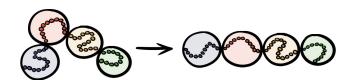
Spatial straightening



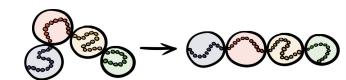
Nucleosome scale



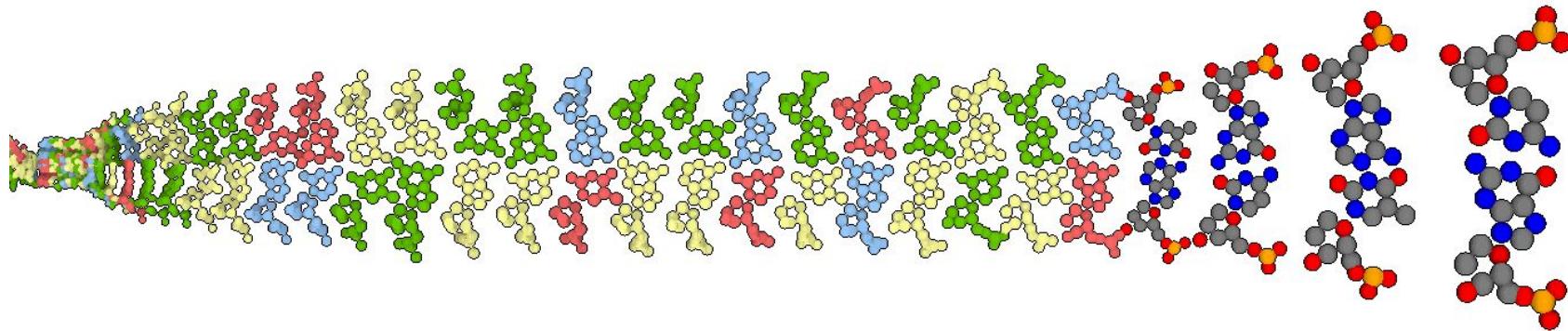
Spatial straightening



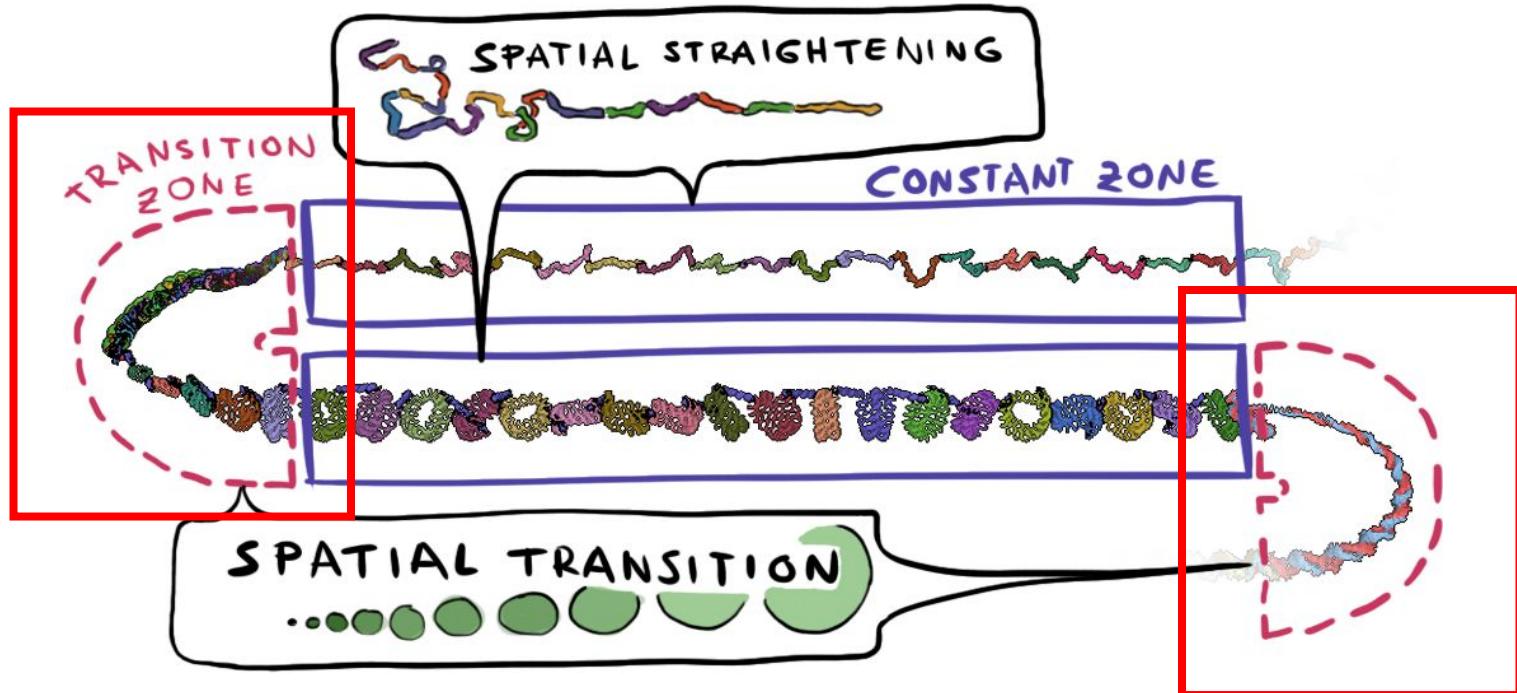
Spatial straightening



Bases scale

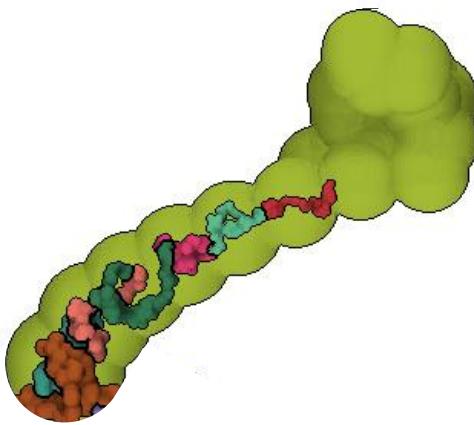
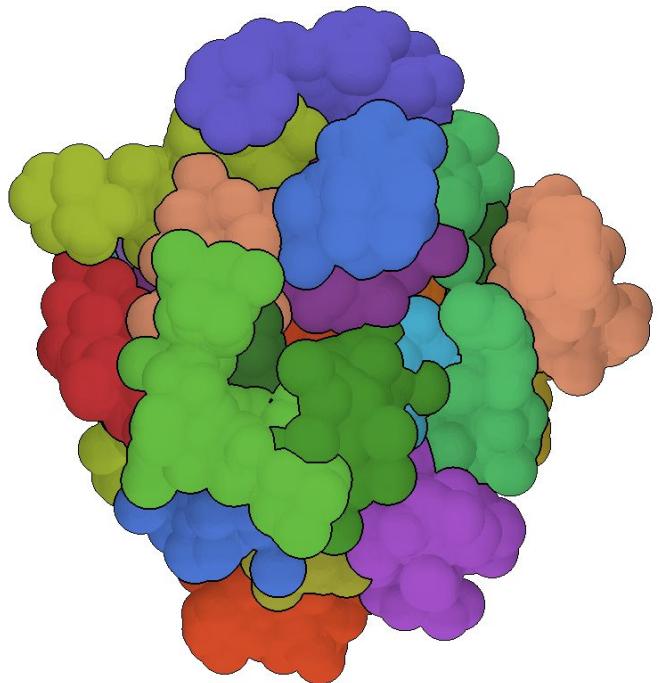


Conceptual arrangement



Spatial transitions

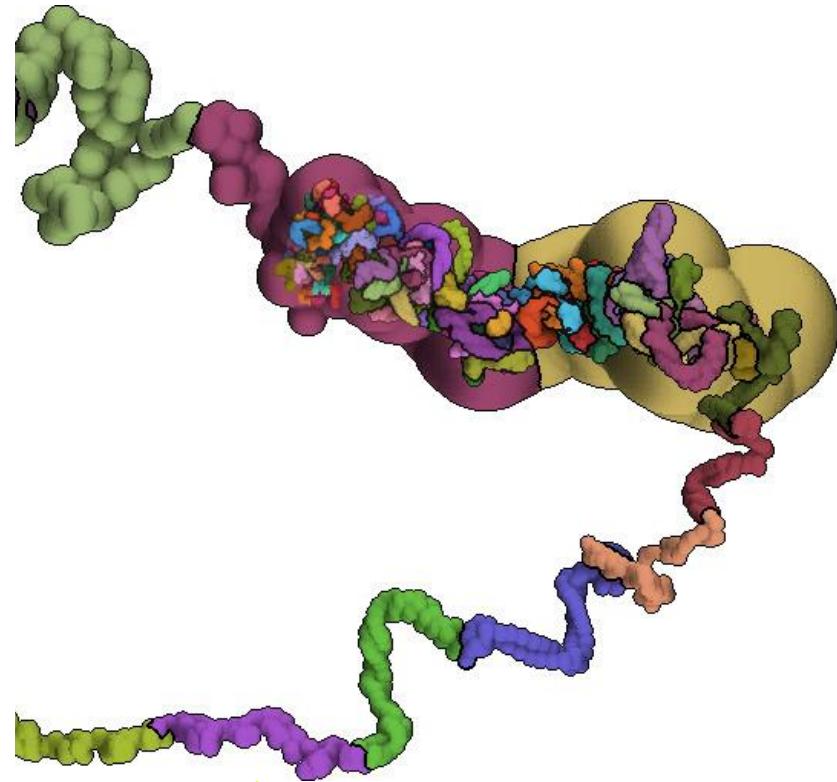
Chromosomes-to-loci



Spatial transitions



Loci-to-fibers



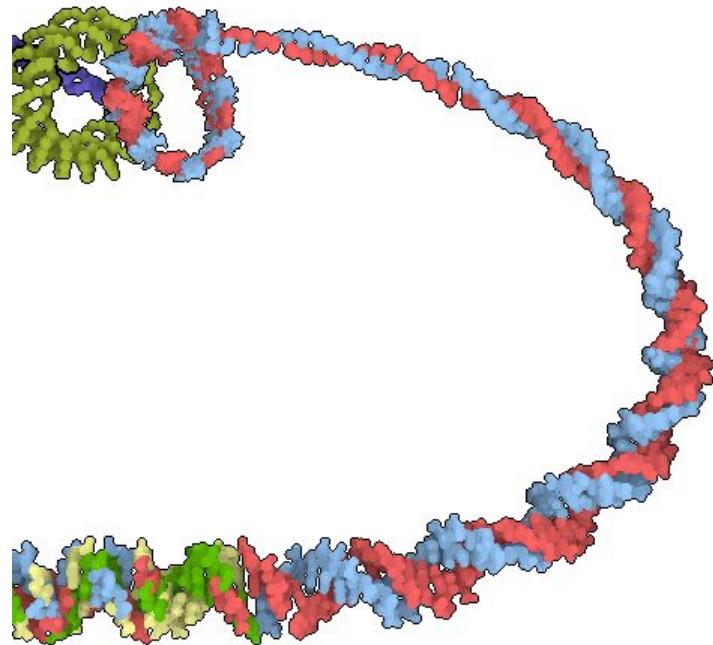
Spatial transitions

Fibers-to-nucleosomes



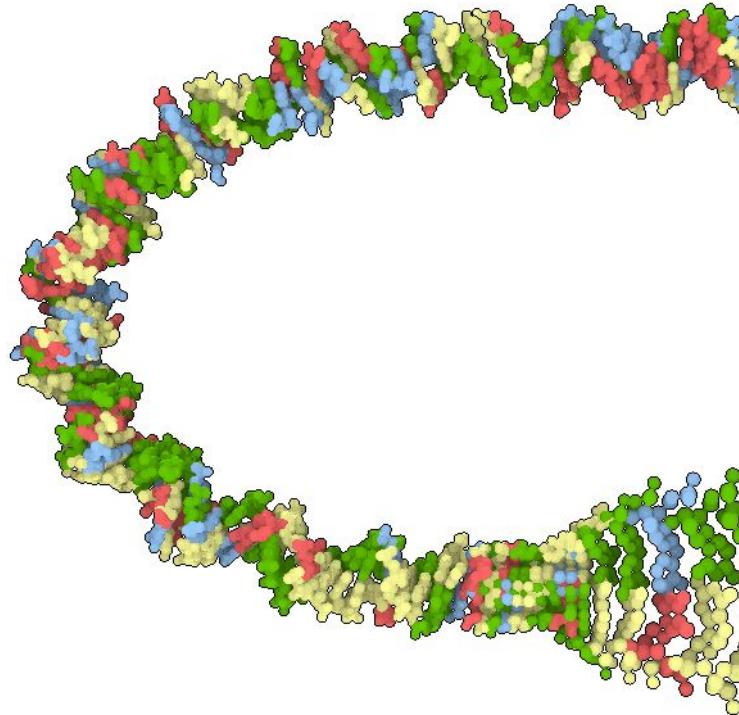
Spatial transitions

Nucleosomes-to-double-helix

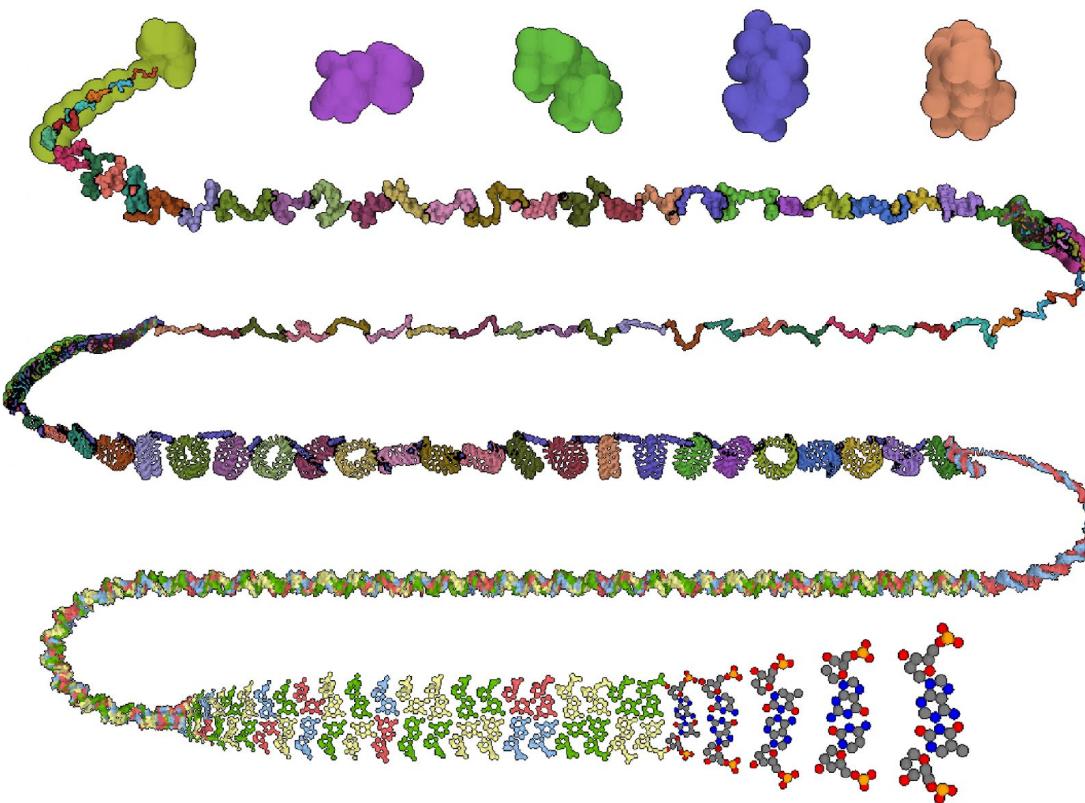


Spatial transitions

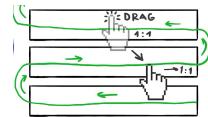
Double-helix-to-bases



Multiscale Unfolding



Panning the scales



Multiscale Unfolding: Illustratively Visualizing the Whole Genome at a Glance

Sarkis Halladjian, David Kouřil, Haichao Miao, M. Eduard Gröller, Ivan Viola, Tobias Isenberg

Inria

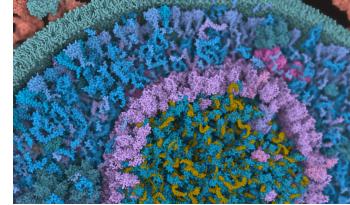
université
PARIS-SACLAY

TU
WIEN

VR VIS



Collaborators



Sarkis
Halladjian



David
Kouřil



Haichao
Miao



Eduard
Gröller



Ivan
Viola

tinyurl.com/scaletrotter

tinyurl.com/multiscale-unfolding

Ludovic Autin



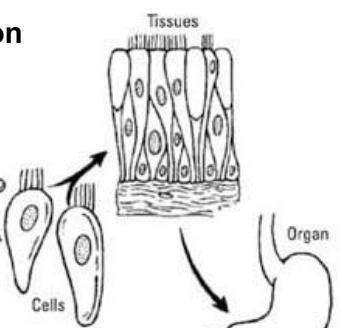
The Mesoscale Challenge



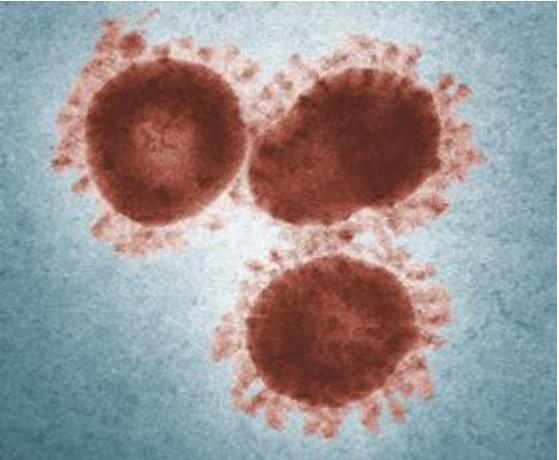
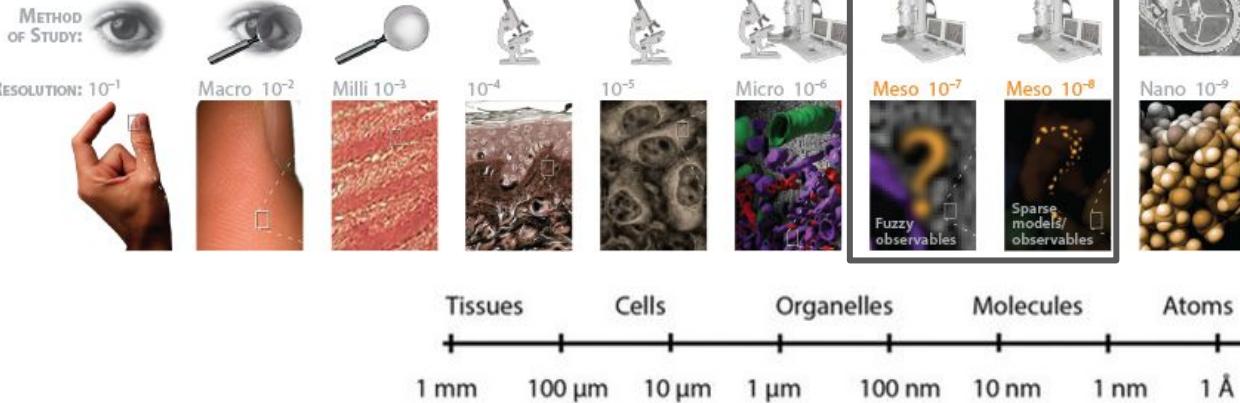
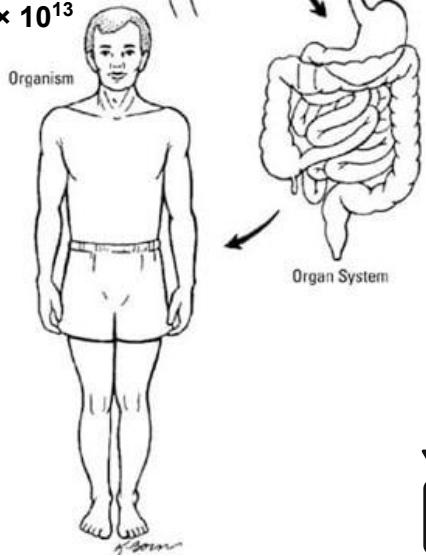
National Institutes of Health
Turning Discovery Into Health

7 octillion
 7×10^{27}

Atoms
Molecules
Cells



30 trillion
 3×10^{13}



Light microscopy

Super-res.

SAXS

Scanning electron microscopy

Electron tomography

Single particle analysis

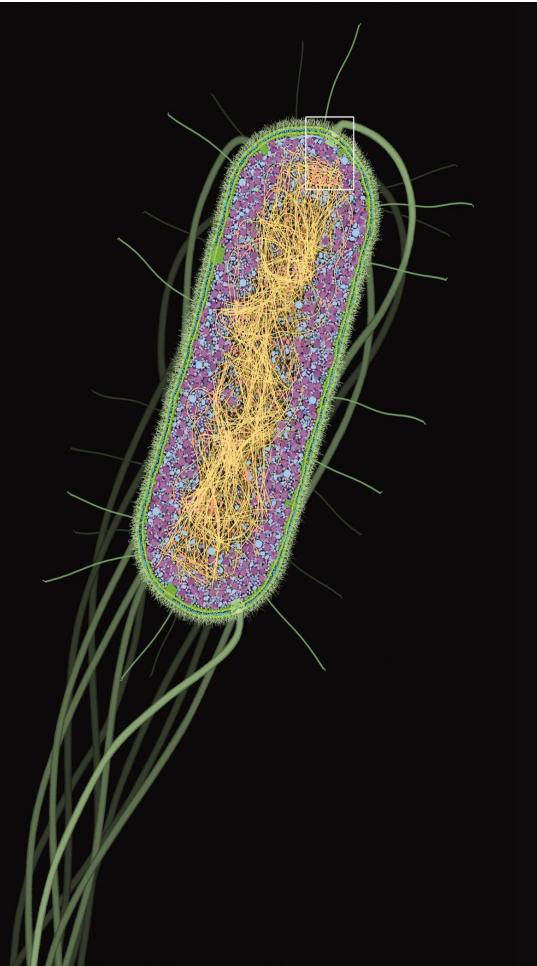
e⁺ cryst./Micro-ED

X-ray crystallography

NMR

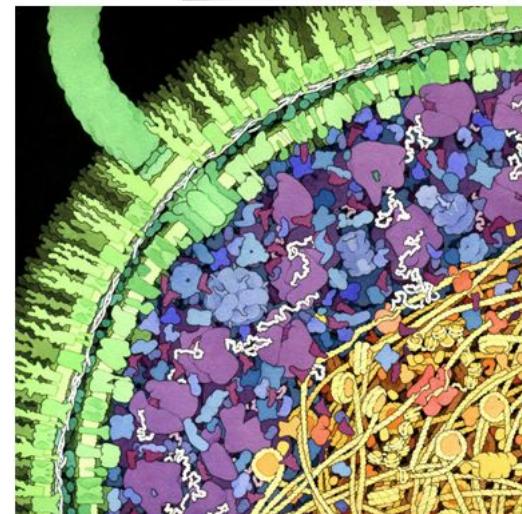
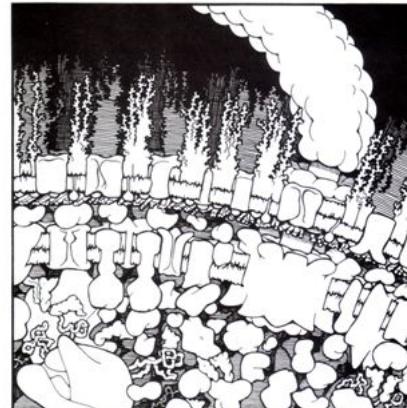
Theodor Escherich 1886

Escherichia coli



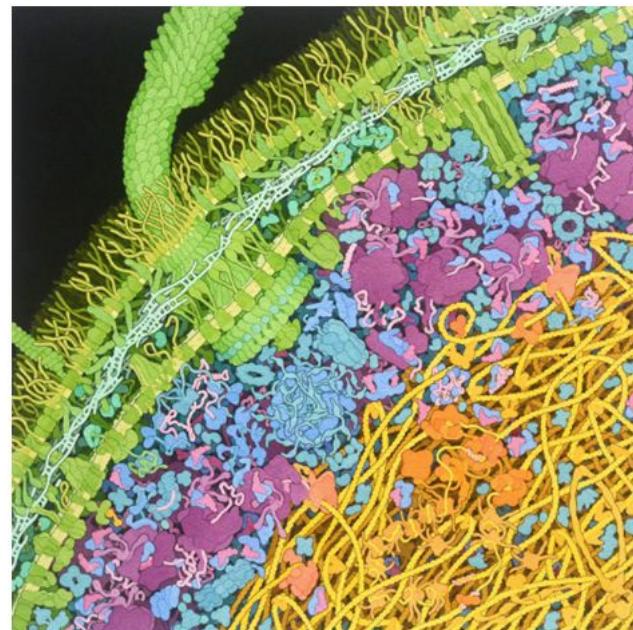
David Goodsell

1991



1999

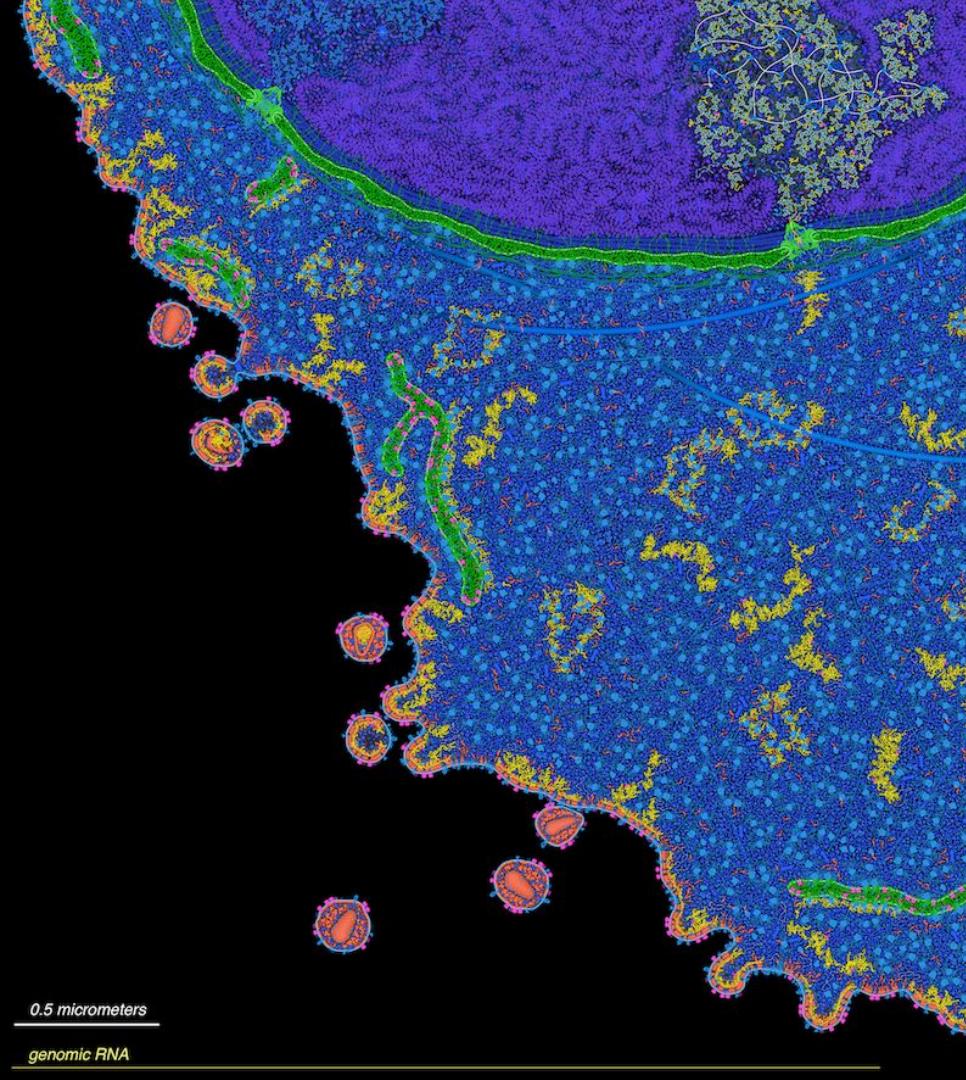
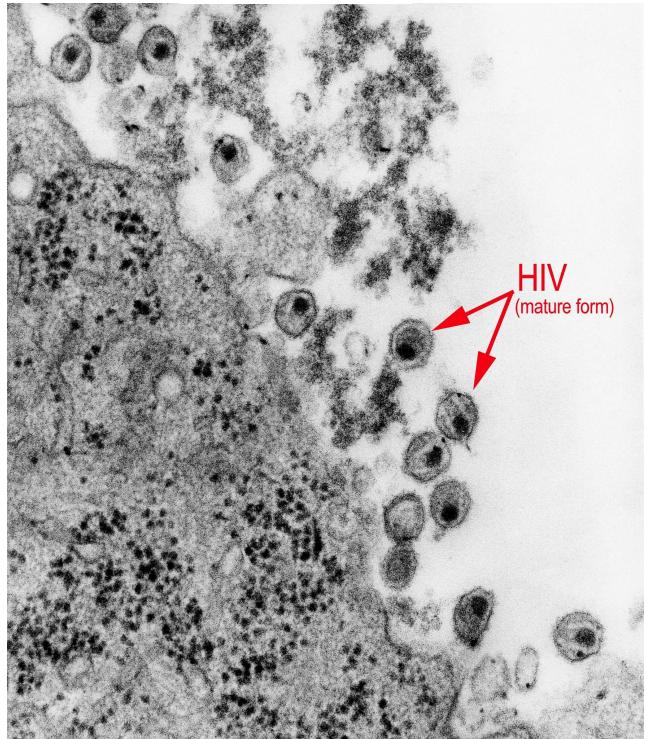
Largely invisible to experiment
Integrates the *current* state of
knowledge



2021



David Goodsell



0.5 micrometers

genomic RNA



David Goodsell

Illustrating Cellular Environments

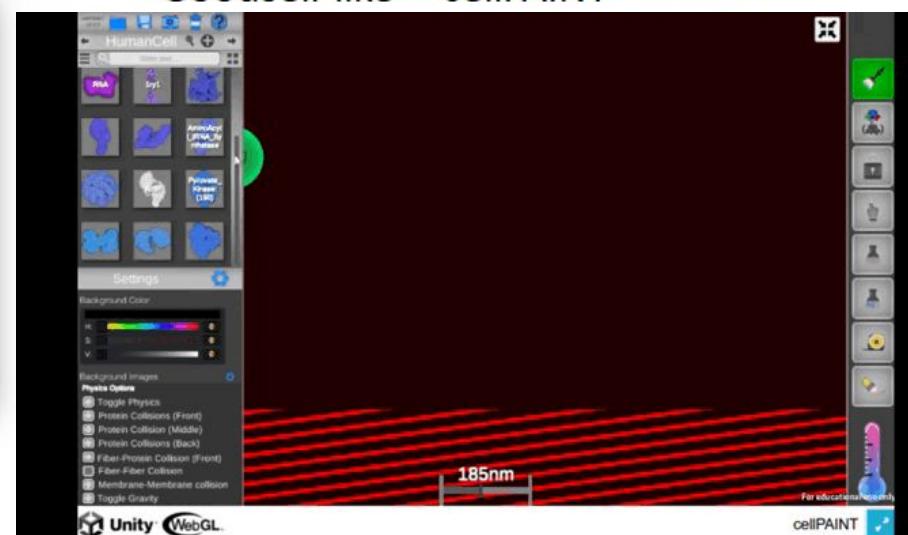


Adam Gardner

User-Friendly Interactive Tools ...CellPAINT Goodsell Watercolor



- Consistent scale
- Simplified iconic shapes
- Cross-sectional view
- Limited depth (3 layers)
- Informative colors



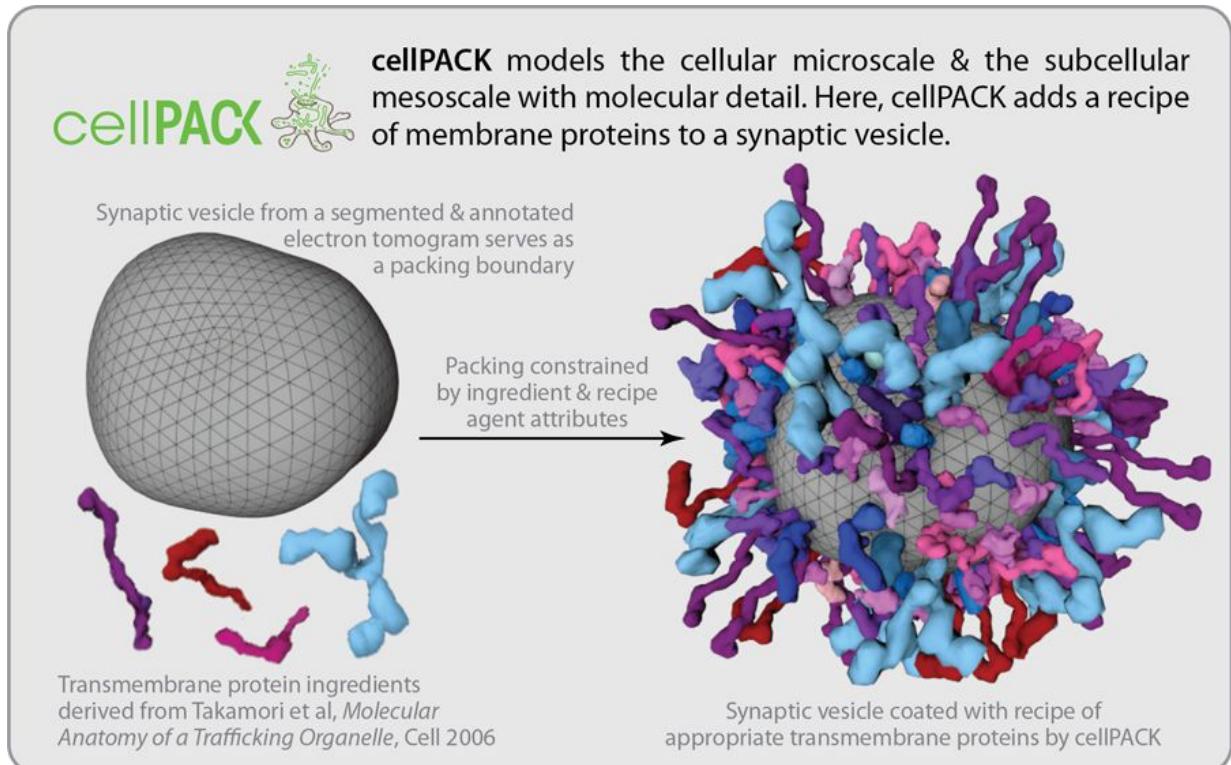
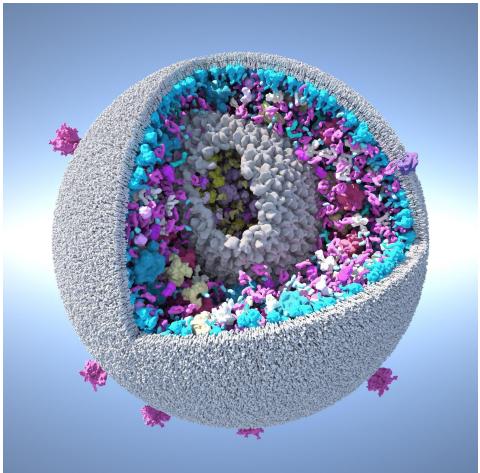
<http://cellpaint.scripps.edu>



cellPACK: a virtual mesoscope to model and visualize structural systems biology

Graham Johnson

2014



Instant Construction and Visualization of Crowded Biological Environments - cellPACK on the gpu



Mathieu Le
Muzic



Tobias Klein



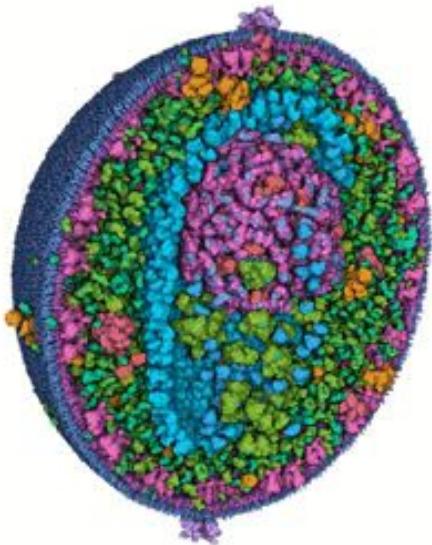
Ivan Viola

2017

The Left Panel interface includes:

- File menu: Loading, D:/Data/HIV/mature_serialized(STAT1).pmln, Load Recipe.
- Buttons: Load Positions, Load Series, Reset.
- Options section:
 - Force align principal axis
 - Show Metadatas
 - Show Bounding Box
 - Show Voids
 - Show Voids Occupied
 - Show Procedural Ingredients Mesh
 - Show Compartments Mesh
 - Populate Exterior Ingredients
 - Populate Surface Ingredients
 - Populate Lipids Surface
 - Populate Interior Ingredients
 - GroB Fiber
 - Populate Procedural Ingredients
 - GroB Fiber real-time
 - Populate Spawning Ingredients
 - Populate Spawning Fiber
 - Visual Jitter
 - Show all Ingredients on a 2D grid
- Buttons: Close File, Update Grid Occupancy.
- Seed and Numbers section:
 - Get Random Seed: A slider from 0 to 1000.
 - Buttons: reset to zero, reset to current volume, Load.
- Protein Number: 156111
- Time: 0.05
- Ingredients List
- Filter Ingredients List
- Overlap Metadatas
- Microscopy

Left Panel



Search for ingredient (select and validate by 'Enter')
Enter HIV

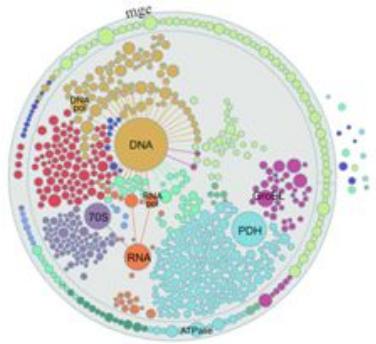
The Right Panel interface includes:

- Search bar: Search for ingredient (select and validate by 'Enter').
- Checkboxes:
 - Enable Metadatas
 - Show metadatas
 - Show resources
- Resolution slider.
- Add one metadatas button.
- Update proteins distribution button.
- CutObject dropdown:
 - Extrude
 - Hide
 - Insert
 - Hard cut
- Buttons: Add, Plane, Remove.
- Reset Cutaway Objects, Reset Camera, Spin Camera, Rock Camera.
- Save Asym:

 - Load Scene State (camera + cutaway)
 - Save Scene State (camera + cutaway)
 - Enable Light
 - Light Follow Camera
 - use HQ coloring
 - Enable Freq / Depth Culling
 - Hide Light+Ingredients overlap
 - Enable Shaders

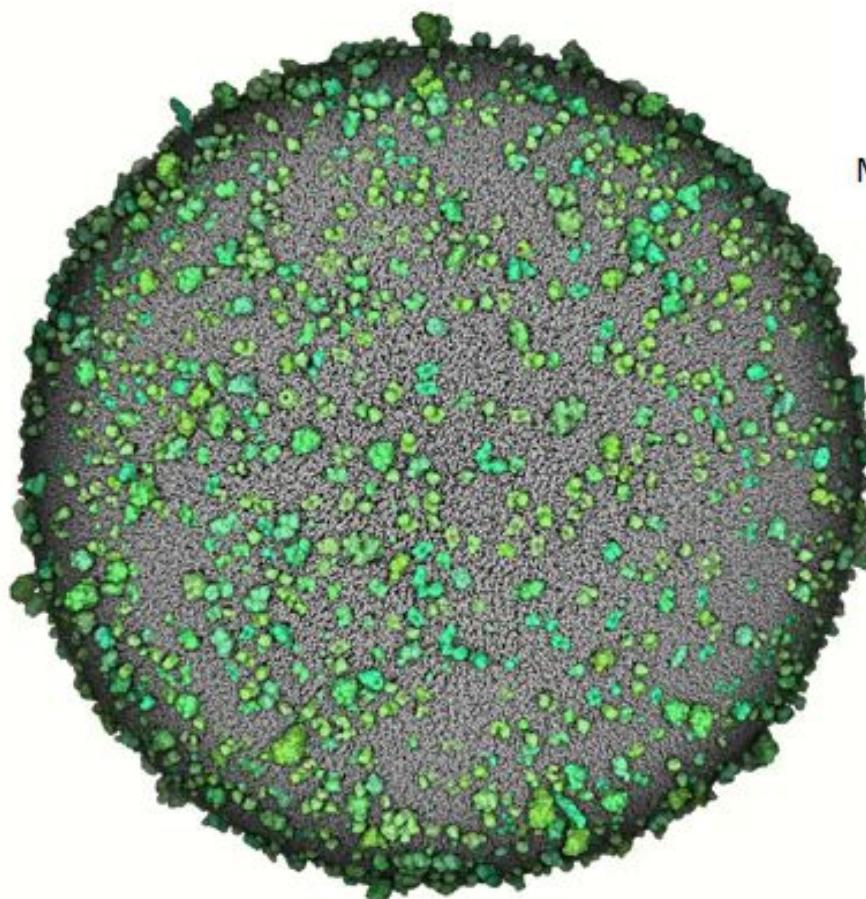
- Scale All Radius slider.
- Color Filter by dropdown.
- Color Protein by dropdown.
- Show LOD info.
- Load Colors Palette (.json), Save Colors Palette (.json), Save Lipids As, Save Model As.
- cellPACK legacy dropdown.
- Object list:
 - root
 - protein
 - HIV_envelope_mature
 - surface
 - interior
 - HIV_capsid
 - surface

Right Panel

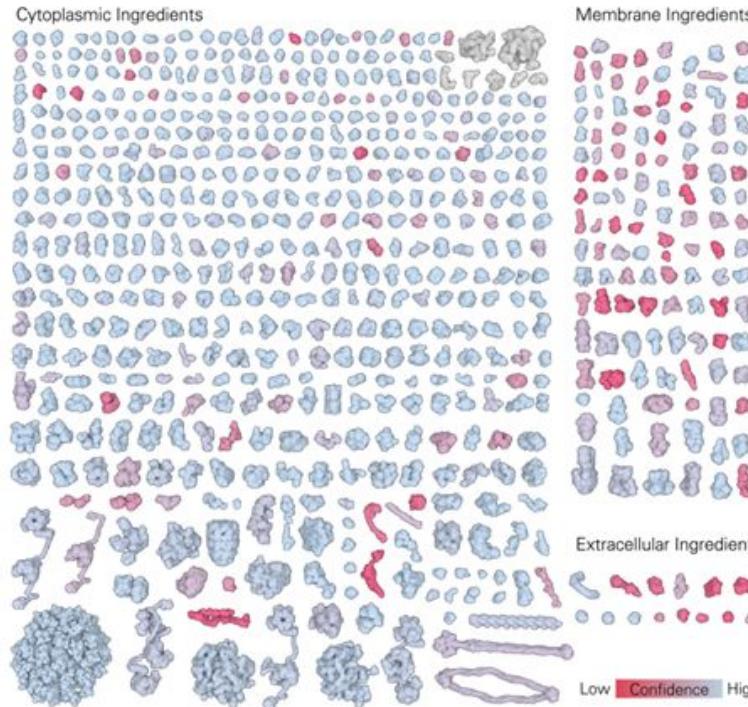


- DNA replication / DNA maintenance
- Transcription
- RNA synthesis / RNA maturation
- Translation
- Protein folding / Maturation
- Metabolism
- Protein transport / Signaling
- Host cell interaction
- Cytokinesis / Motility
- Lipoprotein
- MG-specific
- Uncharacterized

Mycoplasma genitalium, 2022



Martina Maritan





David
Sehnal



Alexander
Rose



Mesoscale Explorer^{BETA}

<https://molstar.org/me/>

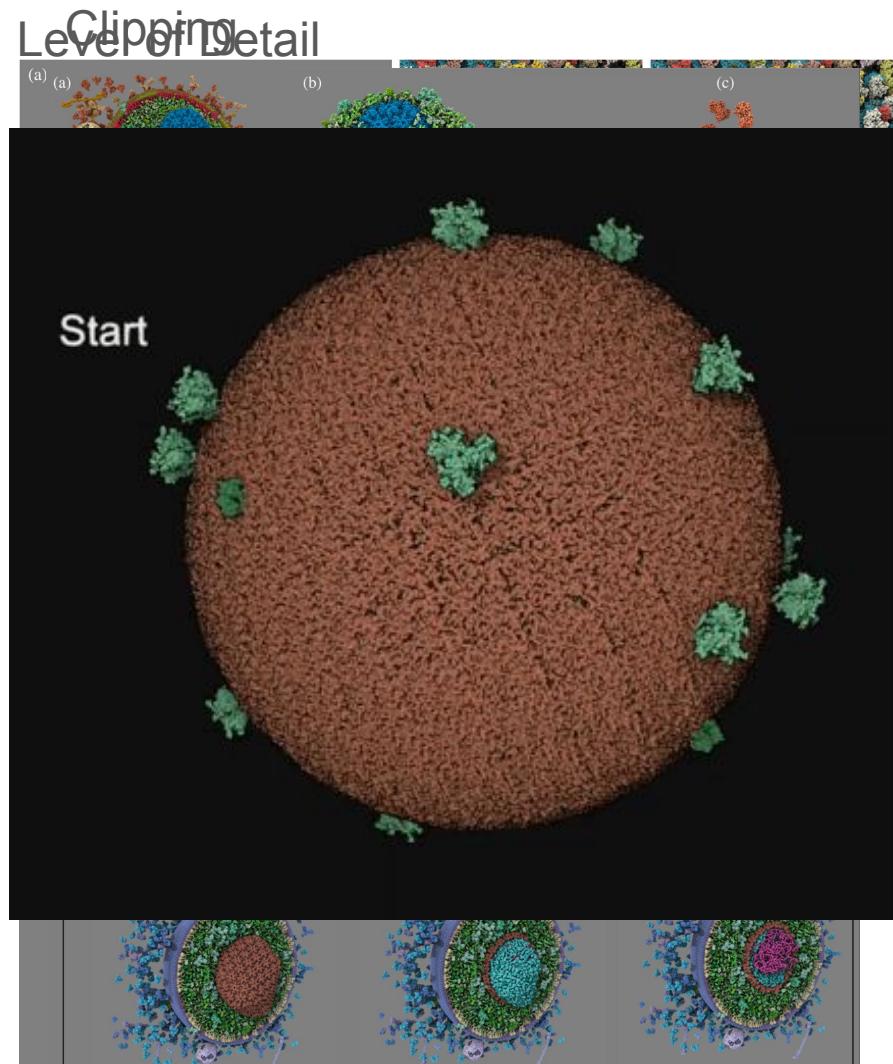
A Mol* app for exploring mesoscale models

The advent of cryo-EM and cryo-ET, coupled with computational modeling, has enabled the creation of integrative 3D models of viruses, bacteria, and cellular organelles. Based on these models, the **Mesoscale Explorer** provides unprecedented access and insight into the molecular fabric of life, enhancing perception, streamlining exploration, and simplifying visualization of diverse data types, showcasing the intricate details with unparalleled clarity.

[Read about Mesoscale Explorer in Protein Science](#)

When using Mesoscale Explorer, please cite:

Alexander Rose, David Sehnal, David S. Goodsell, Ludovic Autin: [Mesoscale explorer: Visual exploration of large-scale molecular models](#), *Protein Science*, 2024; [10.1002/pro.5177](https://doi.org/10.1002/pro.5177).



Collaboration is Key

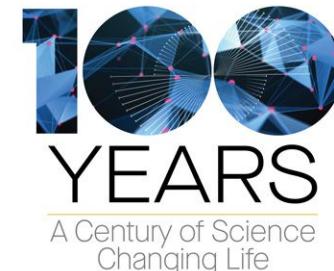
How far are we to visualize a full human cell with molecular details in real-time ?

Data gathering is a bottleneck

Level Of Detail is the common approach for multiscale visualization



Pr. Arthur Olson

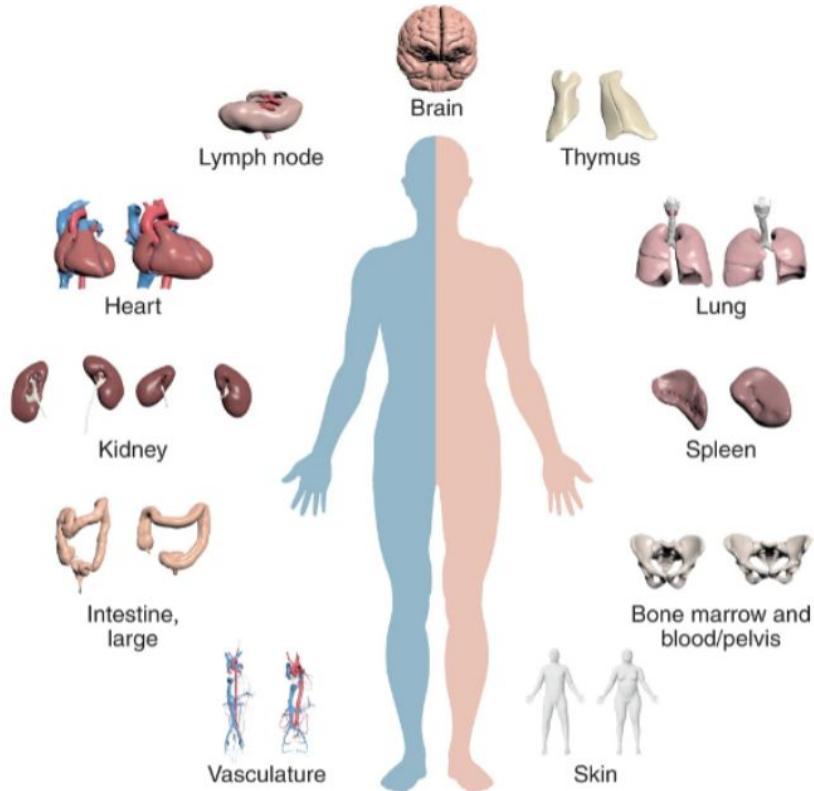


SCRIPPS RESEARCH

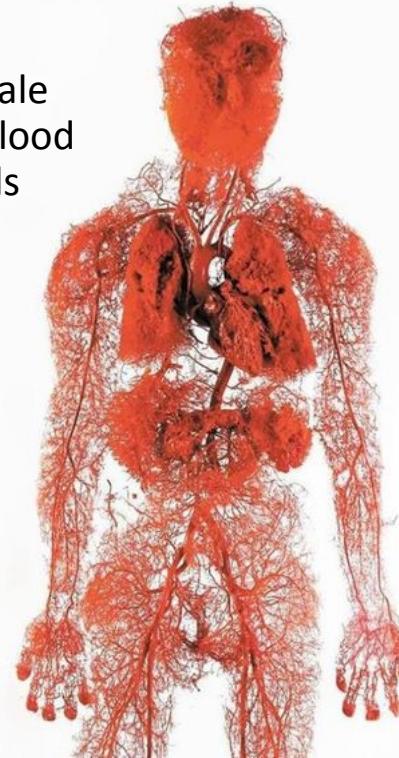
Griffin Weber

Human Reference Atlas (HRA)

<https://humanatlas.io/>



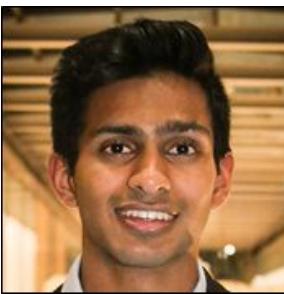
Multiscale
map of blood
vessels



<https://bodyworlds.com/>



Katherine S Gustilo



Avinash Boppana

Sujin Lee

Rajeev Malhotra

Marc Halushka

Ellen M Quardokus

Bruce W Herr II

Ushma Patel

Zorina Galis

Katy Börner

scientific data

OPEN

DATA DESCRIPTOR

Check for updates

Anatomical structures, cell types, and biomarkers of the healthy human blood vasculature

Avinash Boppana¹, Sujin Lee², Rajeev Malhotra^{3,4}, Marc Halushka^{5,6}, Katherine S Gustilo⁵, Ellen M. Quardokus^{1,7}, Bruce W. Herr^{1,8}, Katy Börner^{9,8} & Griffin M. Weber^{1,7,8,10}

More than 150 scientists from 17 consortia are collaborating on an international project to build a Human Reference Atlas, which maps all 37 million cells in the healthy adult human body. The initial release of this atlas provided hierarchical lists of the anatomical structures, cell types, and biomarkers in 11 organs. Here, we describe the methods we used as part of this initiative to build the first open, computer-readable, and comprehensive database of the adult human blood vasculature, called the Human Reference Atlas-Vasculature Common Coordinate Framework (HRA-VCCF). It includes 993 anatomical structures, 150 cell types, and 150 biomarkers, with associated coordinates and links to releasing additional details on vessel type and subtype, branching sequence, anastomoses, portal systems, microvasculature, functional tissue units, mappings to regions vessels supply or drain, geometric properties of vessels, and links to 3D reference objects. Future versions will add variants and connections to the lymph vasculature; and, it will iteratively expand and improve the database as additional experimental data become available through the participating consortia.

Background & Summary
We recently described an ongoing international effort from 17 consortia to construct a Human Reference Atlas (HRA) that maps the entire healthy adult human body down to the single-cell level. It is a monumental task, considering the body has approximately 37 trillion cells. Currently, more than 150 experts worldwide are collaborating on this effort. At its core, the HRA is a set of ASCT+ B tables, which include a hierarchical list of **Anatomical Structures**, the **Cell Types** they contain, and associated **Biomarkers**. Many of the anatomical structures are linked to 3D reference objects. There are dozens of ASCT+ B tables, each representing an organ or organ system. The completed HRA will provide a picture of tissues in the healthy body and their variability across individuals, as well as the precision, progression, and severity of disease, and it will track the changes that occur during disease. The tables are created by an interdisciplinary team of domain experts who have, to date, based the tables on existing knowledge, literature review, or experimental datasets. Over time, multimodal imaging and tissue assays applied to specimens being collected by the consortia will generate new knowledge that can be used to refine the tables.

One of the ASCT+ B tables represents the blood vasculature. Blood vessels are both the source of life for people, bringing oxygen and nutrients to almost all living cells, as well as pathways that lead to disease, including coagulopathies in COVID-19, vascular complications in diabetes, and the spread of metastatic cancers. For 2018, the ASCT+ B table for the blood vasculature was divided into four parts: the main table, which contained the connected dots to create an open, computer-readable, comprehensive database of all the vessels throughout the healthy adult human body. This paper describes the methods we used to do this. The process involved creating a more extensive vasculature database, called the Human Reference Atlas-Vasculature Common Coordinate Framework (HRA-VCCF), with additional details beyond what is currently included in the ASCT+ B tables.

¹Revela, Woburn, Massachusetts, USA. ²Department of Surgery, Massachusetts General Hospital, Boston, Massachusetts, USA. ³Department of Medicine, Division of General Internal Medicine, Massachusetts General Hospital, Boston, Massachusetts, USA. ⁴Department of Anesthesia, Critical Care and Pain Medicine, Baltimore, Maryland, USA. ⁵Computer Intelligence and Engineering, Luddy School of Informatics, Computing, and Engineering, Indiana University, Bloomington, Indiana, USA. ⁶Department of Biomedical Informatics, Harvard Medical School, Boston, Massachusetts, USA. ⁷Department of Medicine, Beth Israel Deaconess Medical Center, Boston, Massachusetts, USA. ⁸E-mail: weber@hms.harvard.edu

SCIENTIFIC DATA | (2023) 10:452 | https://doi.org/10.1038/s41597-023-02018-0

Boppana A, et al. Anatomical structures, cell types, and biomarkers of the healthy human blood vasculature. *Sci Data.* 2023 Jul 19;10(1):452. doi: 10.1038/s41597-023-02018-0.

Multiscale Maps of Roads

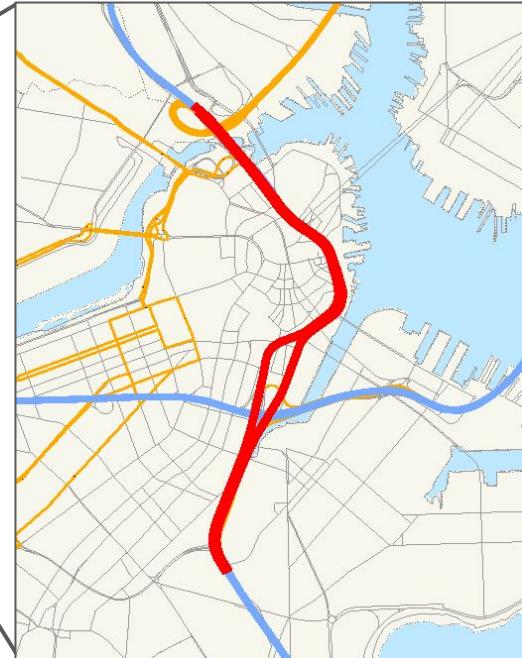
Daily traffic, U.S. National Highway System



Note: Major flows include domestic and international freight moving by truck on highway segments with more than twenty five FAF trucks per day and between places typically more than fifty miles apart.

Source: U.S. Department of Transportation, Federal Highway Administration, Office of Freight Management and Operations, Freight Analysis Framework, version 4.3, 2017.

Boston “Central Artery”



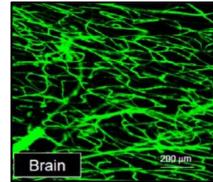
John F. Fitzgerald Expressway, By Sswonk, Public Domain, <https://commons.wikimedia.org/w/index.php?curid=4538754>

Back Bay



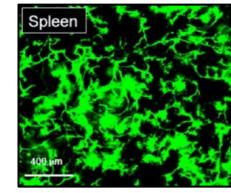
Skeletal muscle

Beacon Hill

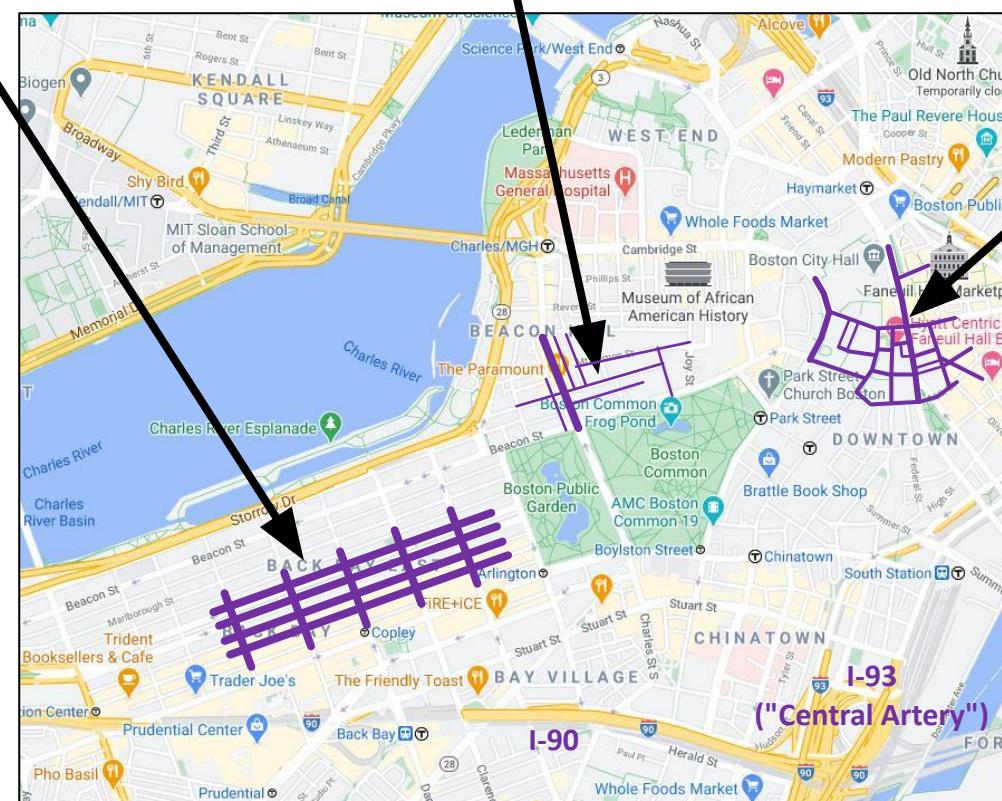


Brain

Downtown



Spleen



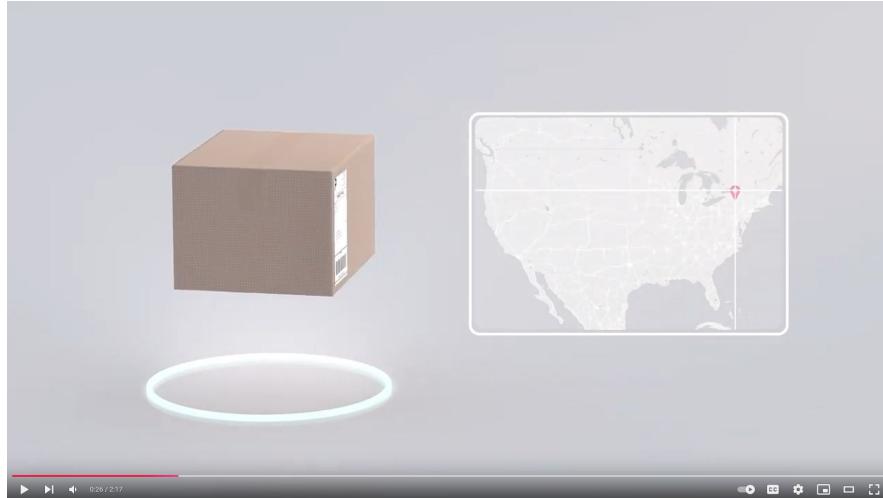
<https://www.zumper.com/blog/best-neighborhoods-in-boston-for-newcomers/>

<https://www.75statestreetgarage.com/nearby-destinations/financial-district/>

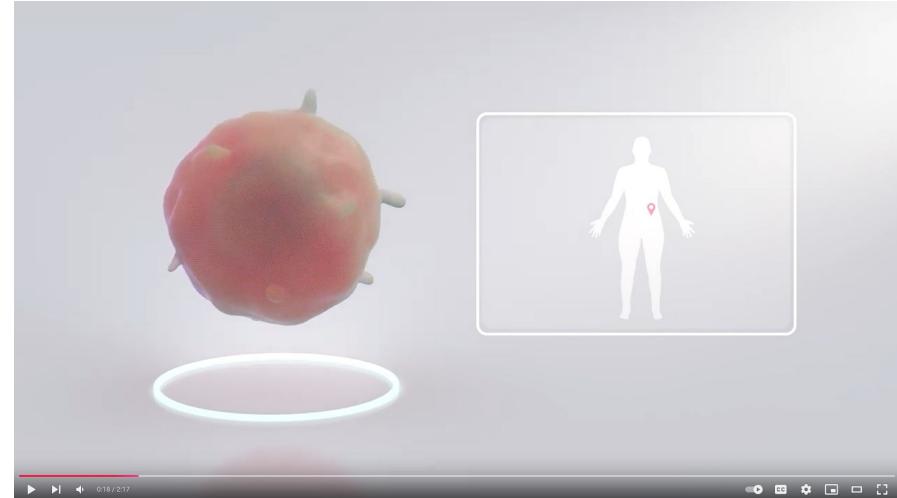
<https://pubmed.ncbi.nlm.nih.gov/27815267/>

Green lines show microvasculature in different tissues

Trucks follow roads to deliver a package to a house



Blood cells follow vessels to deliver oxygen to organs

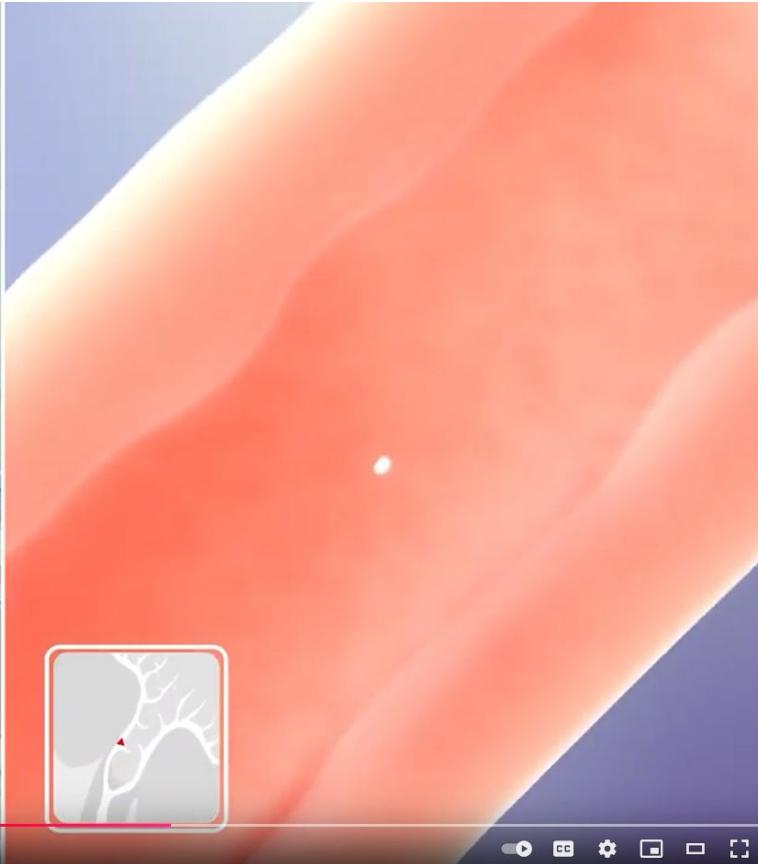


https://www.youtube.com/watch?v=zQeMgxo8n_U

Highway (1000 km)

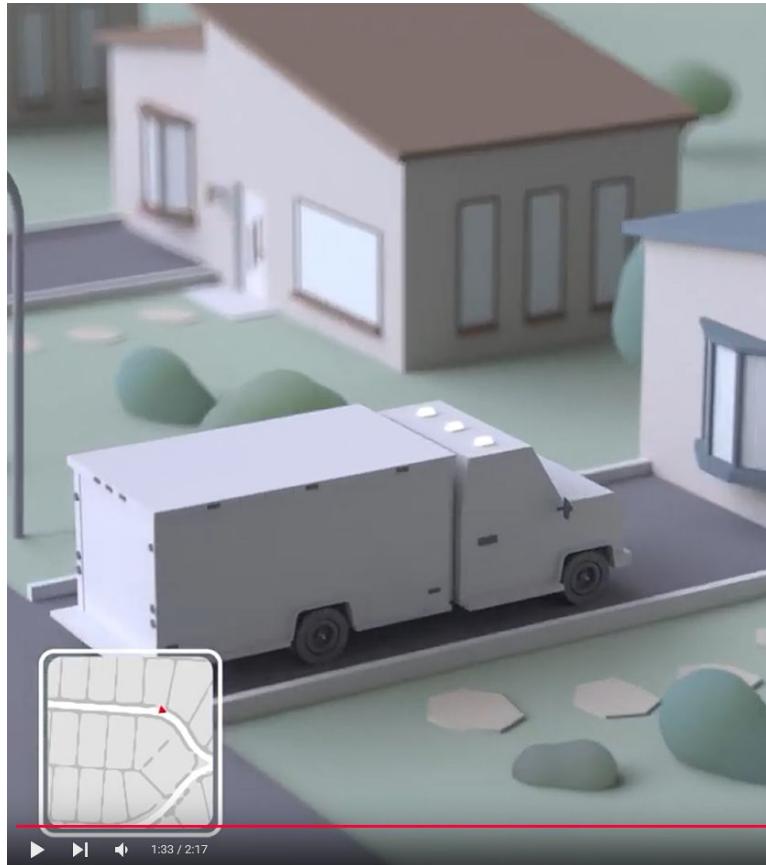


Artery (1 m)

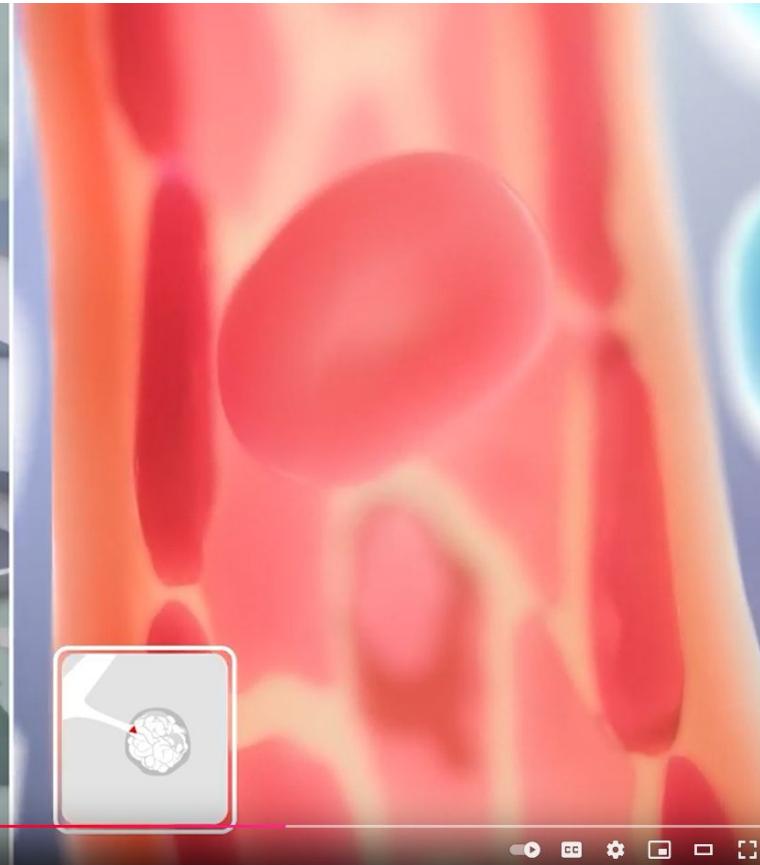


https://www.youtube.com/watch?v=zQeMgxo8n_U

Street (1 km)

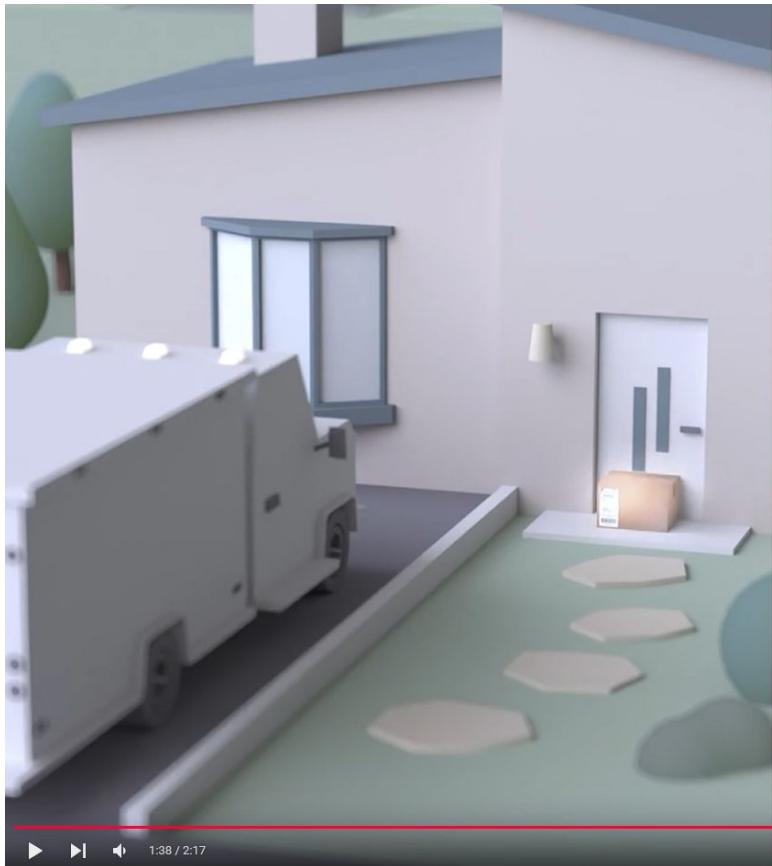


Arteriole (0.5 cm)

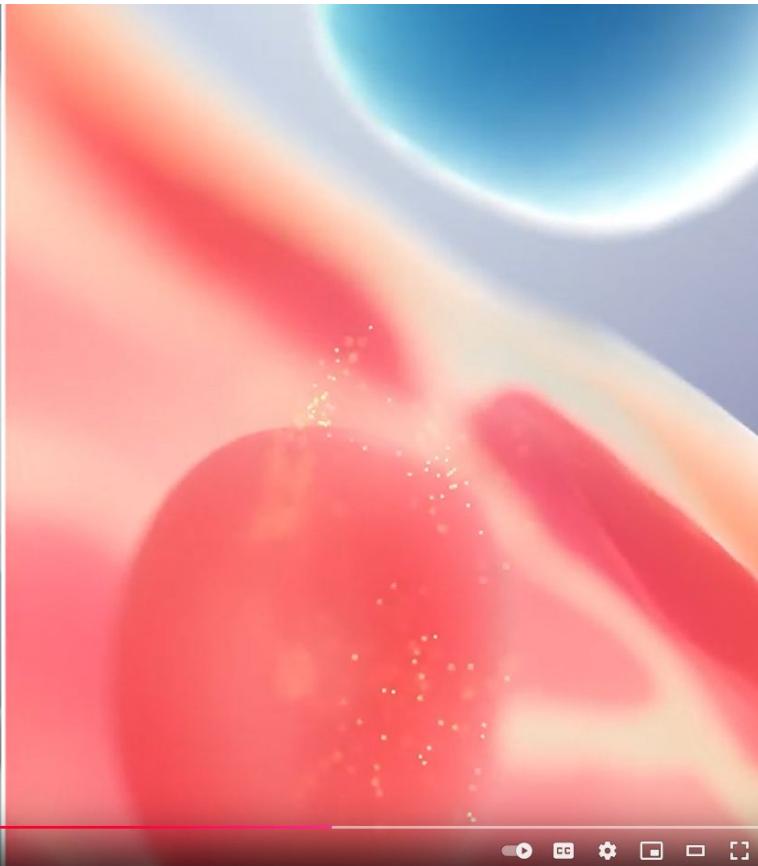


https://www.youtube.com/watch?v=zQeMgxo8n_U

Driveway (10 m)



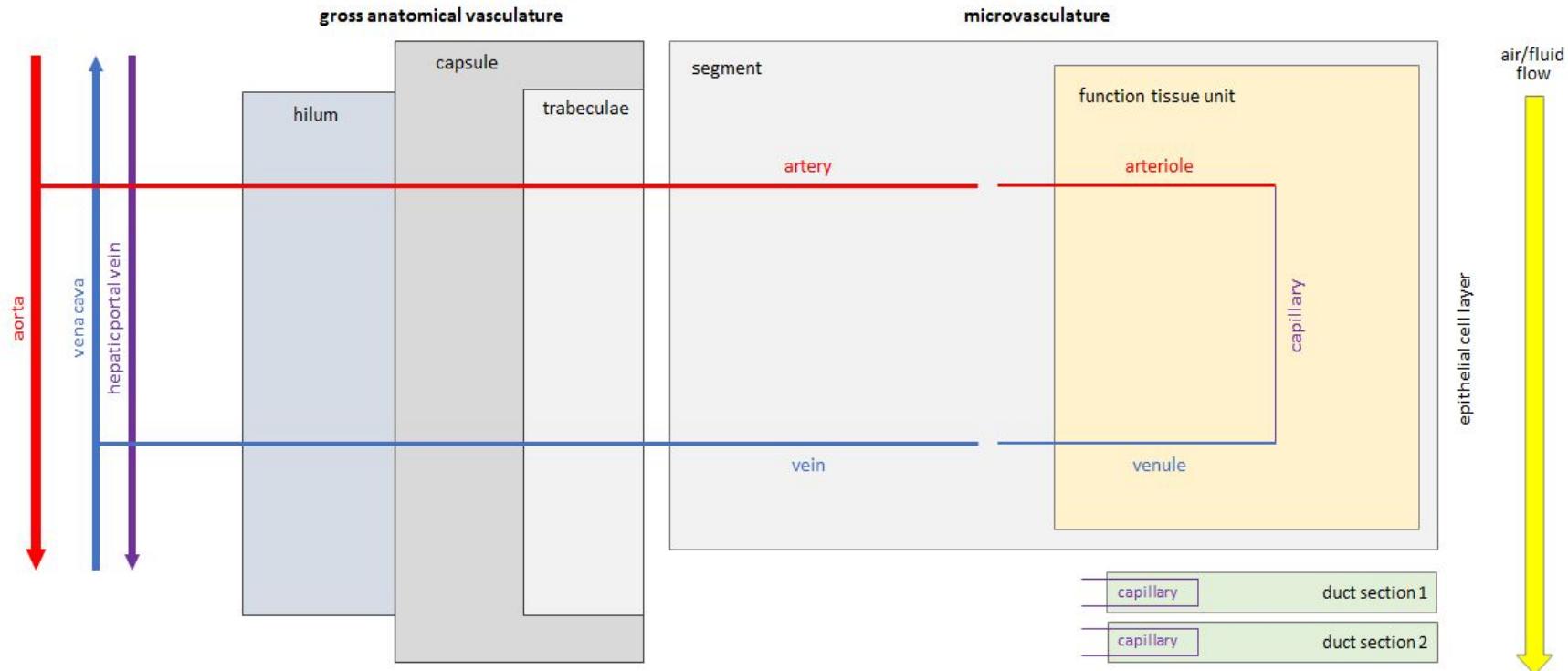
Capillary ($0.5 \text{ mm} \times 0.01 \text{ mm}$)



https://www.youtube.com/watch?v=zQeMgxo8n_U

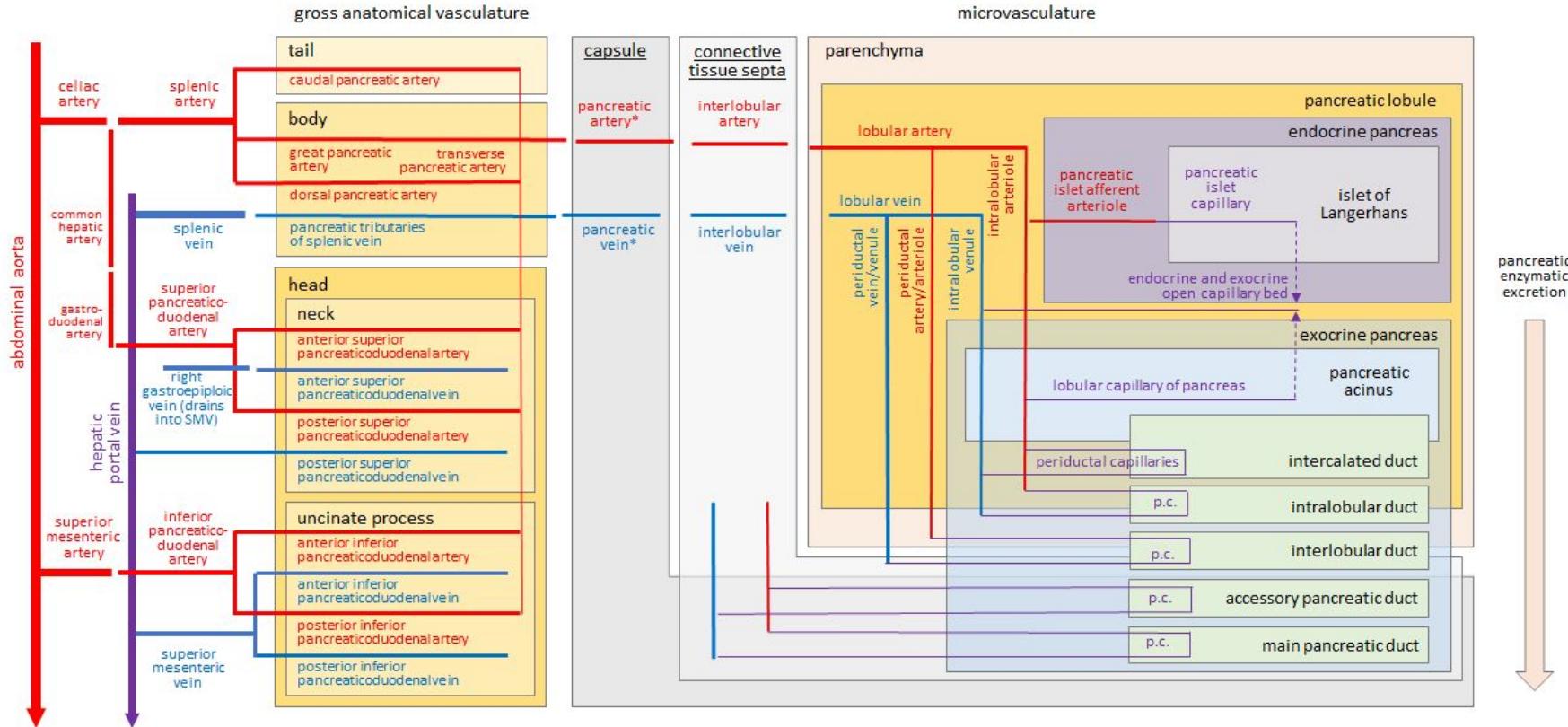
Blood Vasculature to Organ Crosswalk Diagrams

Template



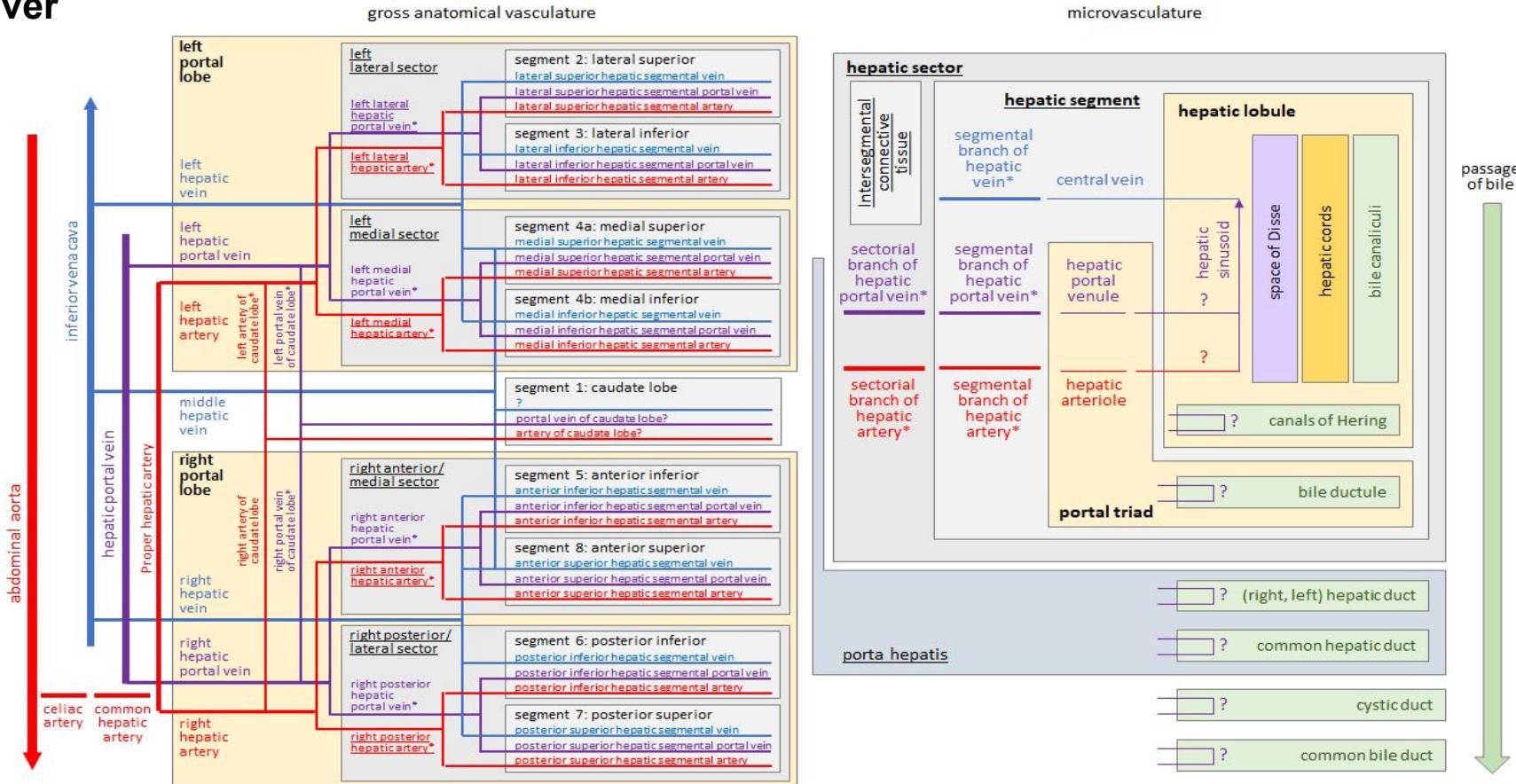
Blood Vasculature to Organ Crosswalk Diagrams

Pancreas



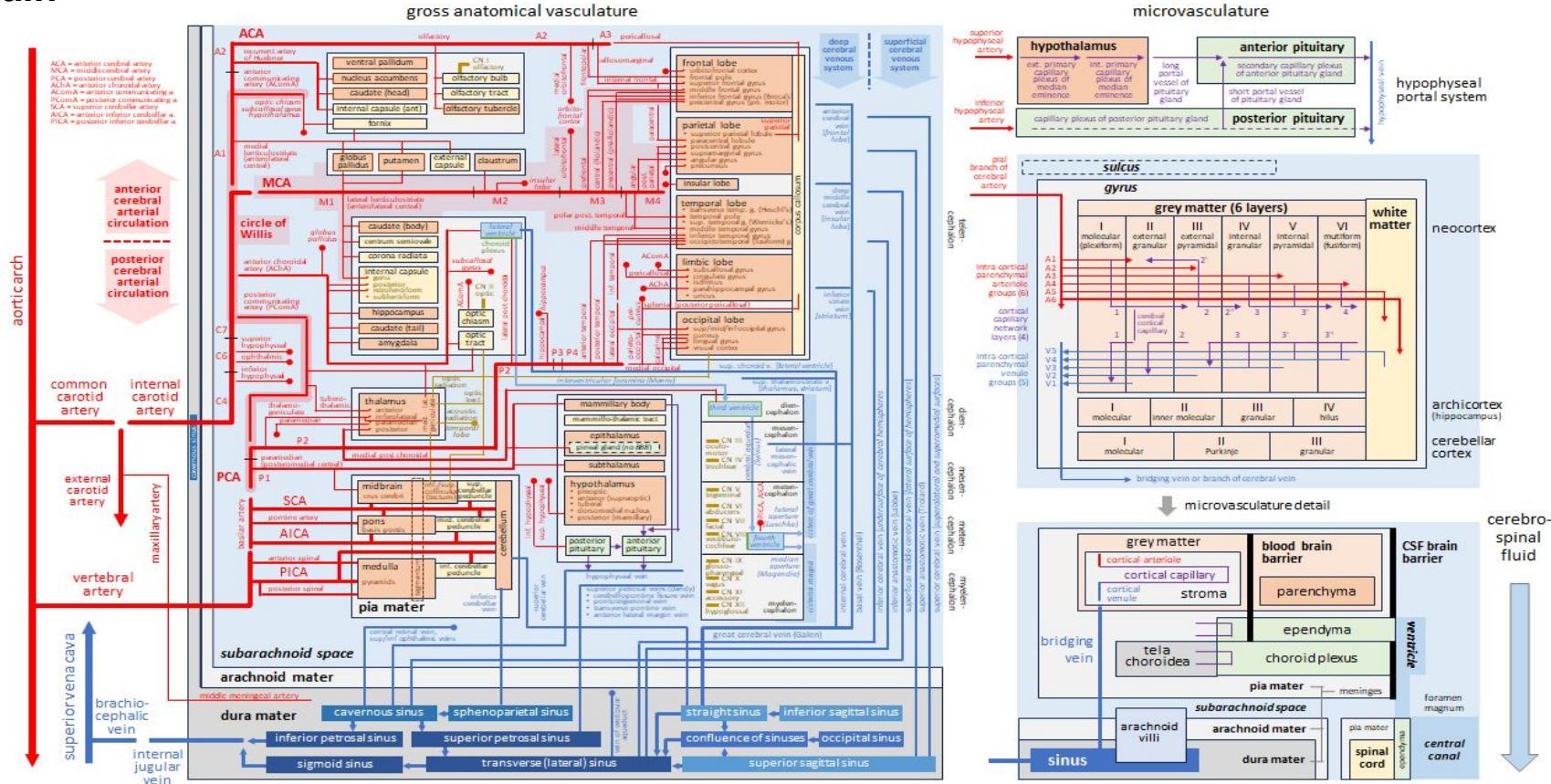
Blood Vasculature to Organ Crosswalk Diagrams

Liver



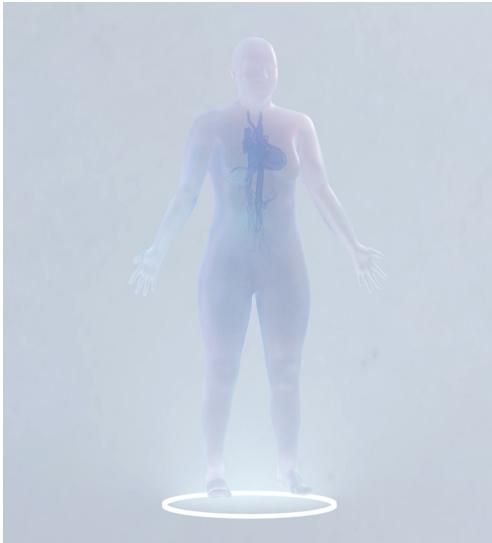
Blood Vasculature to Organ Crosswalk Diagrams

Brain





Q&A



- What is the best definition of “multiscale visualizations”?
- What application domains are best and worst served by multiscale visualizations?
- What kinds of conceptual and cognitive challenges exist for constructing and reading multiscale visualizations?
- What kinds of visual encodings make sense only in multiscale visualizations, and which ones do not make much sense?
- What kinds of interactions are needed to make multiscale visualizations usable?
- **What are the opportunities afforded and challenges posed by using extended reality (XR) technologies such as virtual, augmented, and mixed reality (VR, AR, MR) for multiscale visualizations?**
- **What is the relationship between multiscale visualizations and one-scale visualizations, such as bar graphs, scatter graphs, or line graphs? How can one be served by the other?**
- **When are stepless and stepped zooms better, respectively?**

Q&A

<https://humanatlas.io/events/2024-24h>



Thank you
