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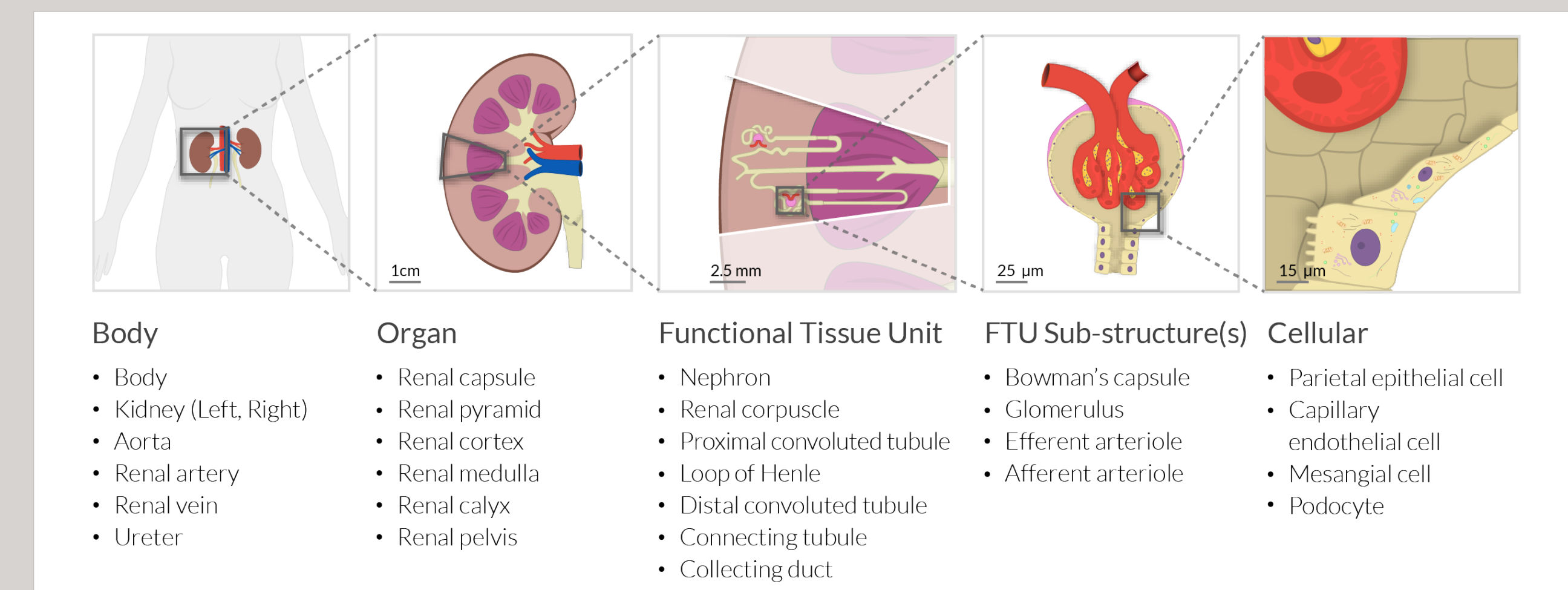
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Abstract

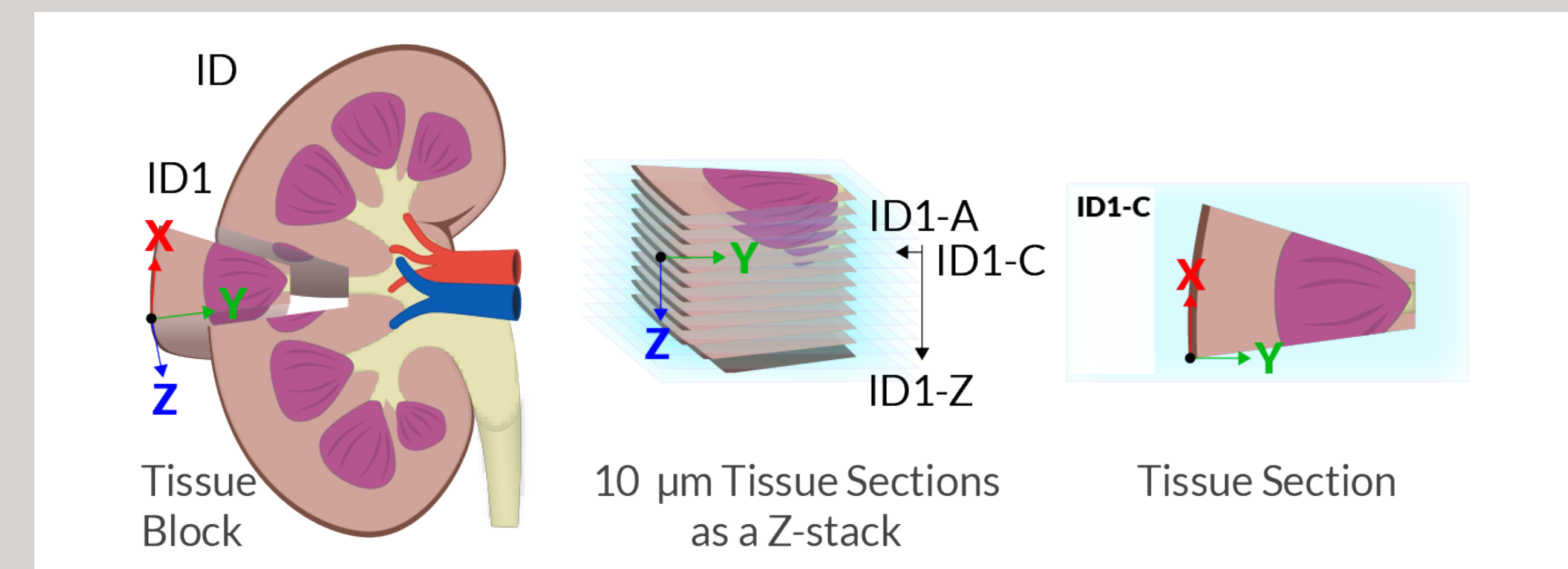
The ultimate goal of the HIVE Mapping effort is to develop a common coordinate framework (CCF) for the healthy human body that supports the cataloguing of different types of individual cells within anatomical structures, understanding the function and relationships between those cell types, and modeling their individual and collective function. In order to exploit human and machine intelligence, different visual interfaces are implemented in support of CCF data generation, exploration, and communication. The CCF and the interactive data visualizations are multi-level and multi-scale. They support the registration and exploration of diverse types of data—from single cell to whole body. In the initial two years, MC-IU ran user needs analyses with stakeholders, compiled an initial CCF ontology and associated 3D object library, developed novel CCF registration and exploration UIs, and explored using the vasculature as a coordinate system to map all cells in the human body.

Common Coordinate Framework

A common coordinate framework (CCF) is a conceptual and computational framework for the storage, analysis, and (visual) exploration of spatially and semantically indexed data—across individuals, technologies, labs.



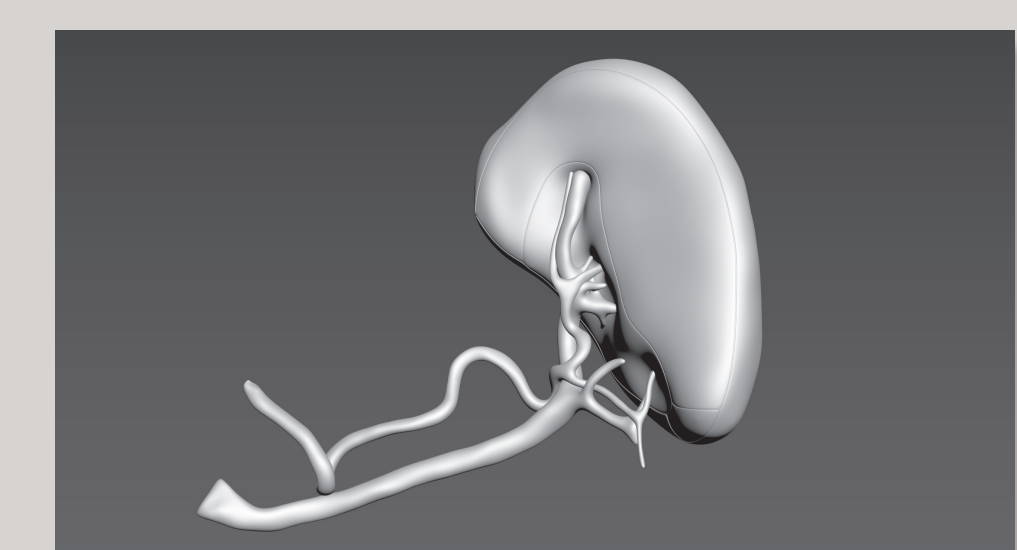
Semantic zoom from whole human body, to organ, to functional tissue units (FTUs), to FTU sub-structures, to single-cell level.



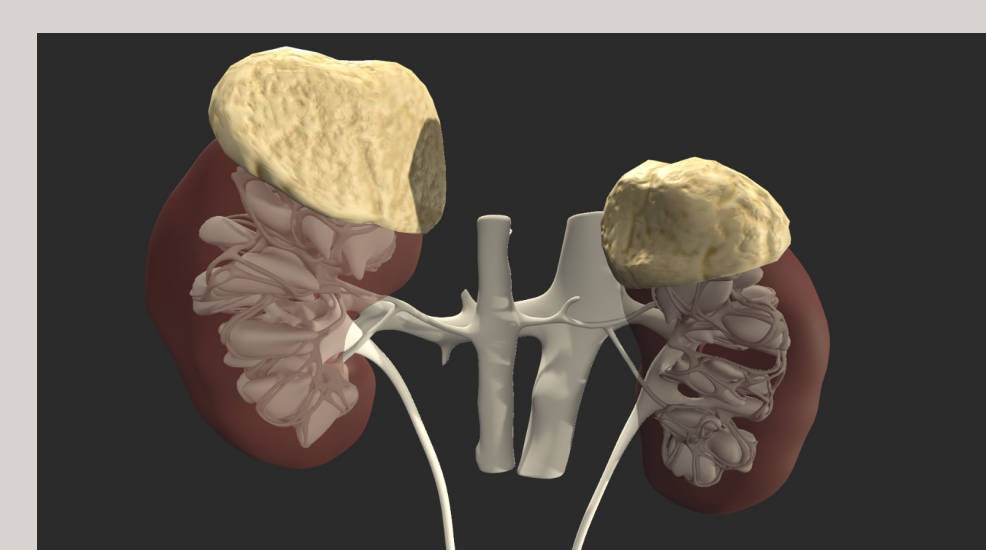
3-step spatial registration of single cells in relation to reference organs.

CCF 3D Object Library

In collaboration with Kristen Browne at National Institute of Allergy and Infectious Diseases (NIAID), NIH we are developing a library of anatomically correct human organ models using data from NLM's Visible Human (VH) dataset.

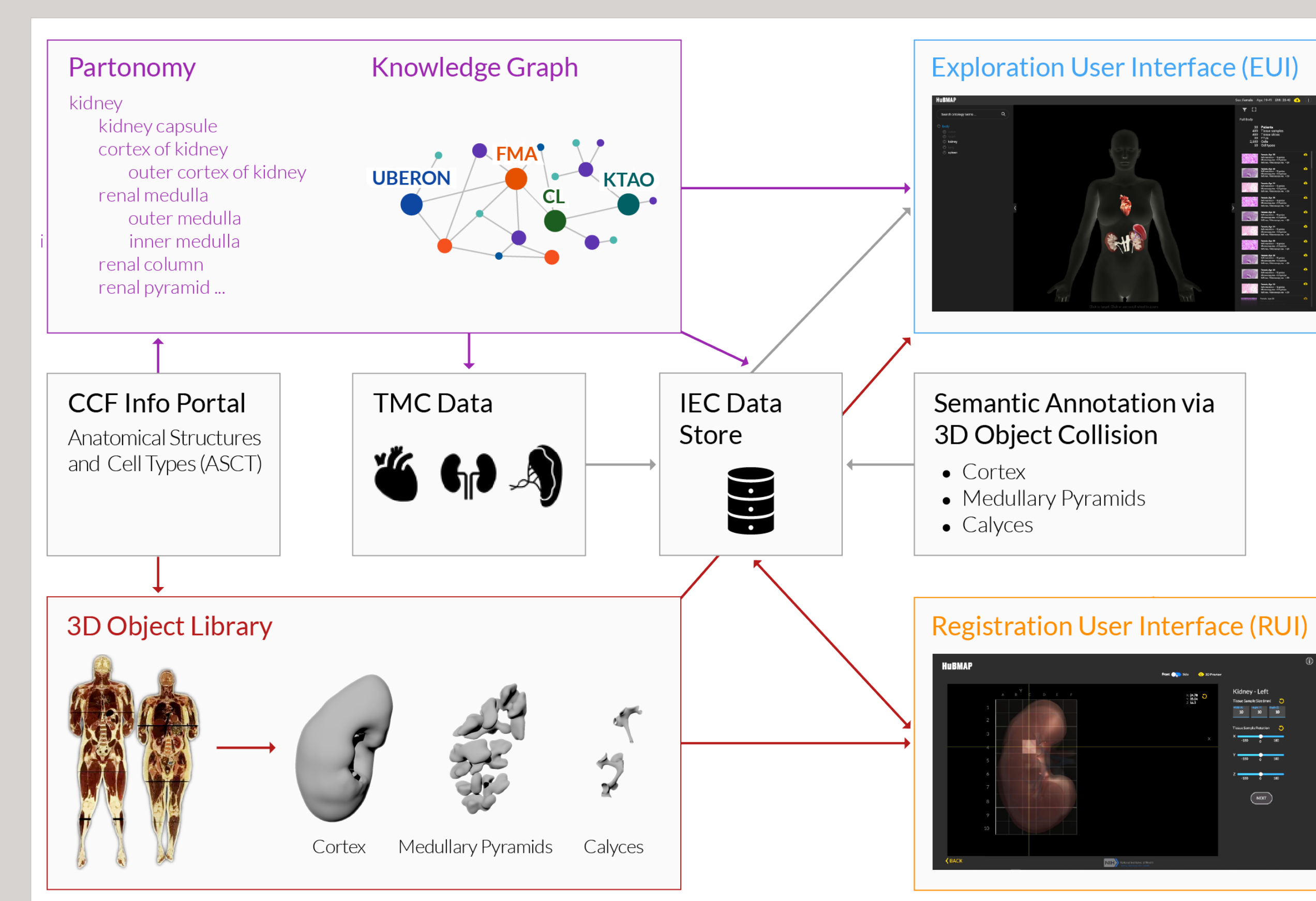


VH Spleen



VH Kidney

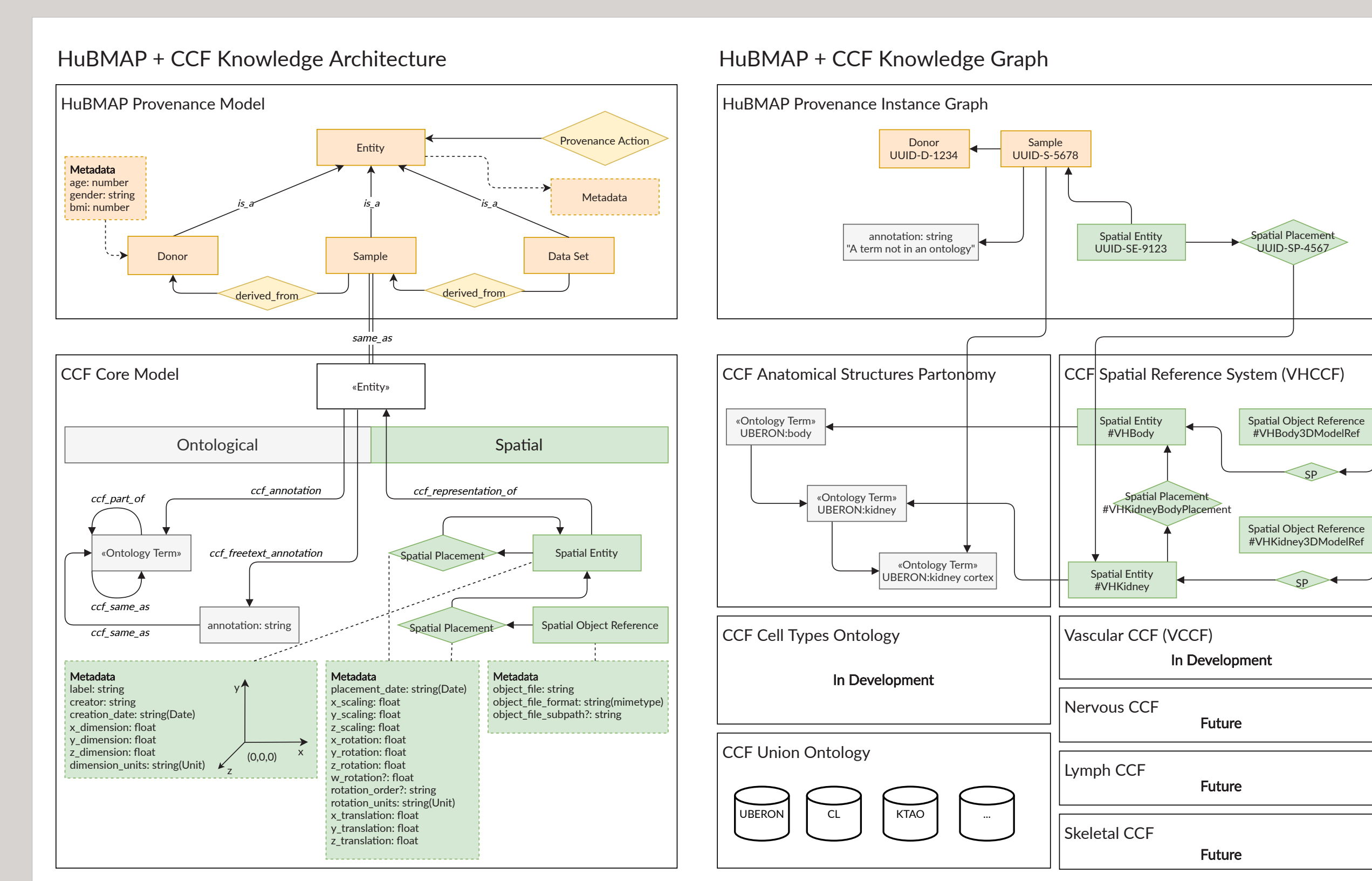
CCF Registration to CCF Exploration Workflow



Overview of CCF Info Portal (left) which systematically captures CCF relevant information, CCF Ontology design (top left) and 3D Object Library construction (lower left), and CCF User Interfaces (right). Arrows indicate data flow.

CCF Ontology

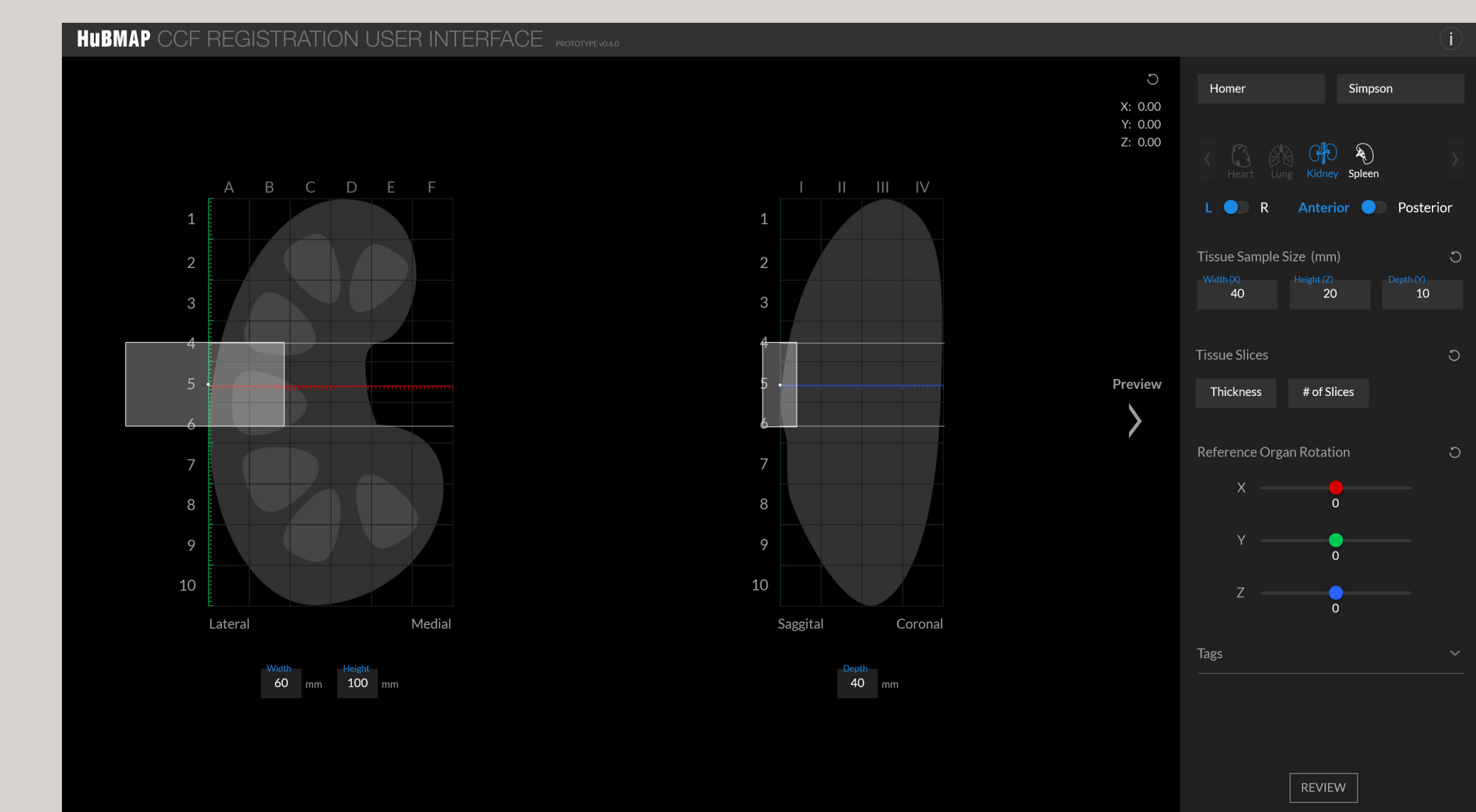
The CCF Core Model has been defined as a formal ontology using Web Ontology Language 2 (OWL) to support compatibility and interlinkage with other ontologies.



CCF Core Model, see <https://hubmapconsortium.github.io/hubmap-ontology/ccf.owl>

CCF Registration User Interface (RUI)

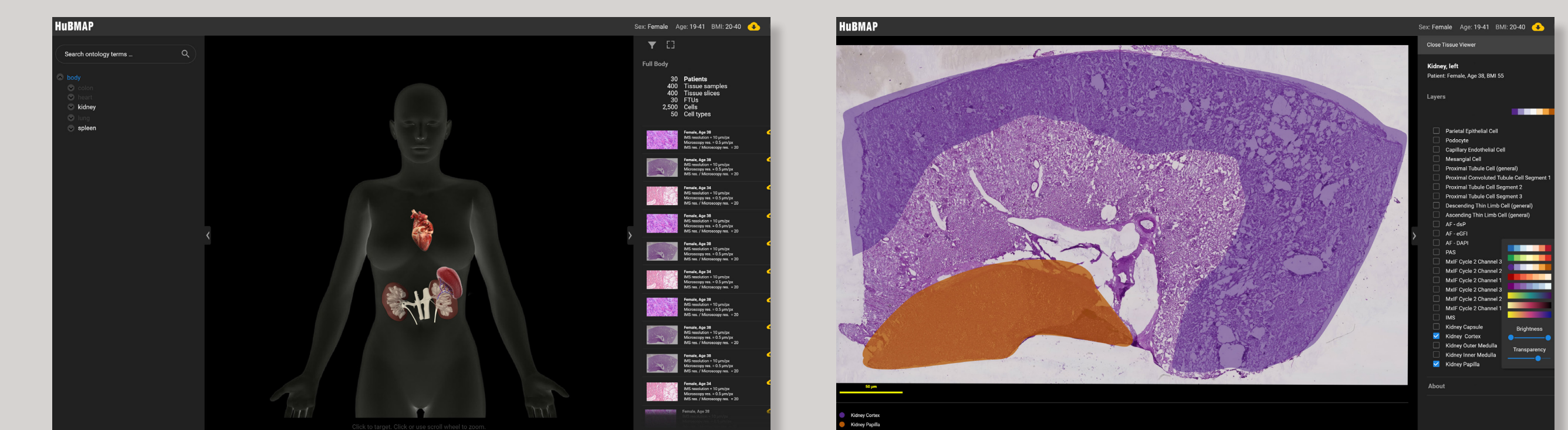
The RUI was designed for usage by experts that collect human tissue and need to document the tissue extraction site. It requires about 5 minutes of training time and 2 minutes for each tissue registration. Currently, the RUI supports gross anatomical tissue registration of tissue blocks. When biomolecular data becomes available, it will be extended to support placement based on biomolecular markers and patterns.



RUI functionality can be examined at <https://hubmapconsortium.github.io/ccf-3d-registration>

CCF Exploration User Interface (EUI)

The EUI makes it possible to explore 2D/3D tissue data semantically and spatially across multiple scales. Spatial data generated by the RUI is used to position tissue blocks. Cell segmentation algorithm results will soon support cell position and cell type exploration. Semantic and spatial search, browsing, filtering, and details on demand are supported.



EUI functionality can be examined at <https://hubmapconsortium.github.io/ccf-ui/>

Publications

- Griffin M. Weber, Yingnan Ju, Katy Börner. Considerations for Using the Vasculature as a Coordinate System to Map All the Cells in the Human Body. *Frontiers in Cardiovascular Medicine*. 7 (29): doi: 10.3389/fcvm.2020.00029
- Michael P. Snyder et al., 2019. The Human Body at Cellular Resolution: The NIH Human Biomolecular Atlas Program. *Nature*. 574: 187-192. doi: 10.1038/s41586-019-1629-x.
- MC-IU, TMCs, NIAID. Conceptualization, Construction, Validation, and Usage of a Human Body Common Coordinate Framework. Draft available for expert comments.

Acknowledgements

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