

# Common coordinate systems for the human body

NYGC Mapping

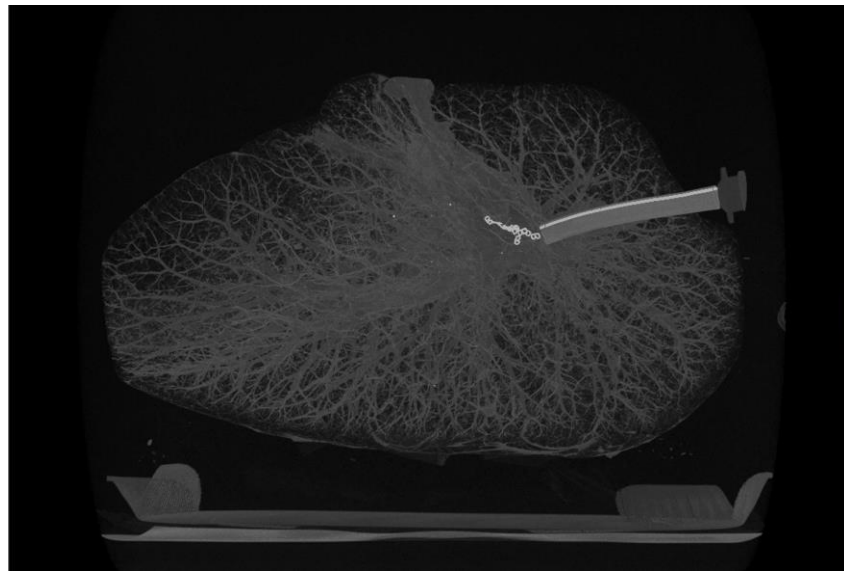
Rahul Satija, John Marioni, Aviv Regev

# Building a reference framework for HuBMAP

*Question: How can we map across individuals to minimize inter-individual variation due to spatial differences?*

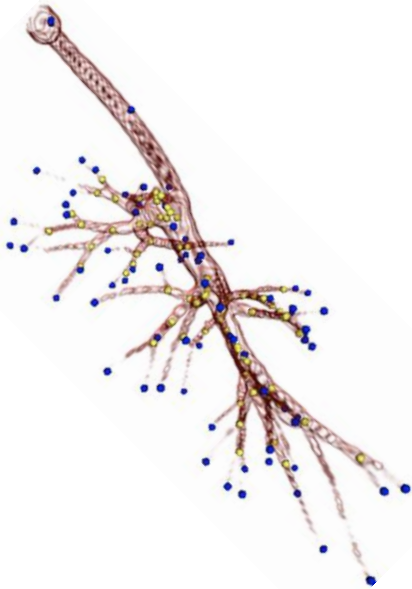
Chose the lung as a pilot to construct a Common Coordinate Framework.

High-resolution CT data available through collaboration with LungMAP

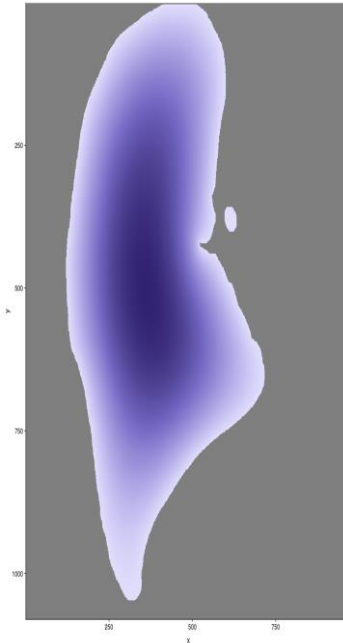


# Progress in Year 1: Methods for CCF construction

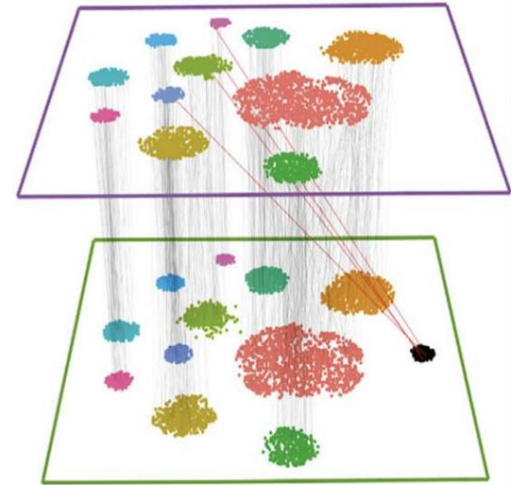
Internal spatial landmarks



Shape-based landmarks



Molecular landmarks



# Next year's deliverable

## 1. Common Coordinate Framework (Lung)

- Given a location in one sample, map to a corresponding position in another sample
- Develop strategies for lung, can be generalized to other tissues
- CCF Workshop (with MC-IU) in 2020

## 1. Molecular integration across individuals and technologies

- Tools for the 'harmonization' of different HuBMAP and community datasets
- Integrate and classify cells across experiments, modalities, and technologies
- Demonstrate in lung and kidney, but methods will broadly apply to many tissues

**Collaboration with LungMAP and KULMAP: Gloria, James Hagood, Xin Sun**

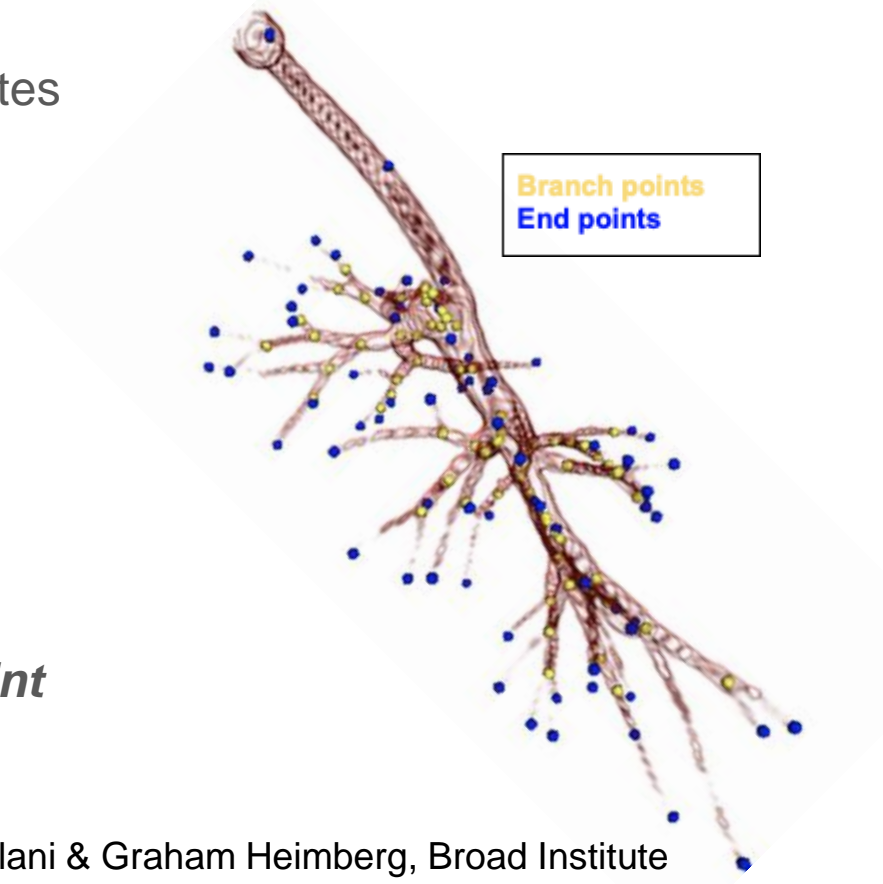
# Internal spatial landmarks - Airway skeletonization

Airway cell studies: Airway-based coordinates are more important.

How can we consistently reference airway locations across individuals?

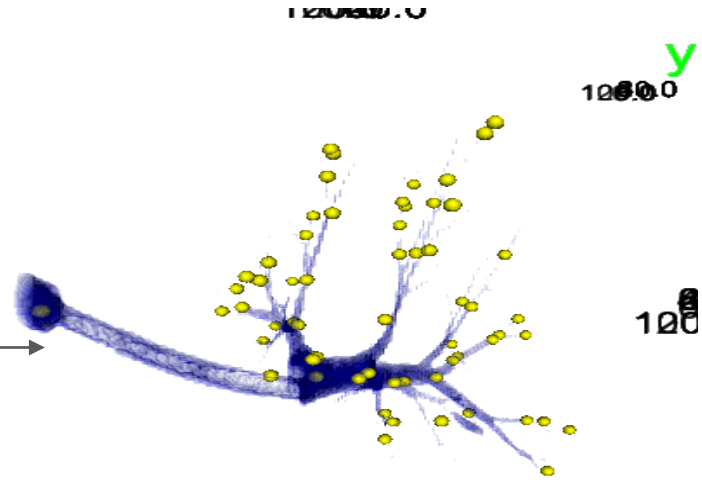
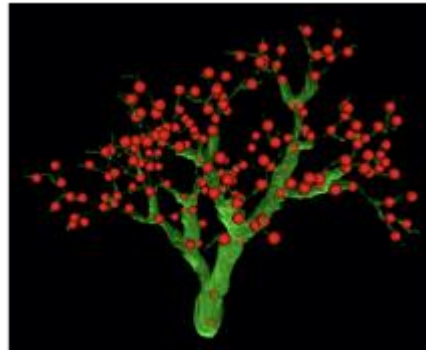
1. Segment airways in CT data
2. Detect branch points and endpoints
3. Align airway trees across individuals

***Problem: simple methods for branchpoint detection fail on lung CT data***



# Internal spatial landmarks - Neural network

- Simulate lung branching
- Train Convolutional Neural Network (3D UNET) to detect branchpoints and endpoints
- Test on simulated data
- Run neural network on real lung data

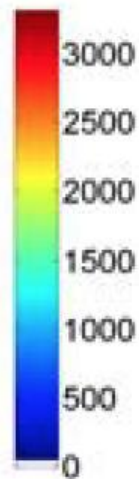


Tommaso Biancalani & Graham Heimberg, Broad Institute

# Poisson distance map

Distance from any given point inside a shape to the edge

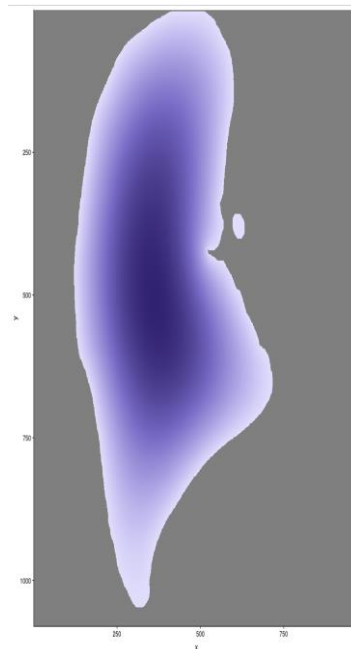
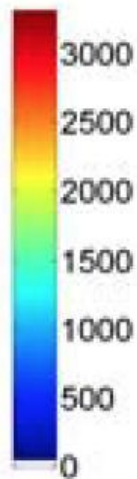
- Random walk to edge of image
- Coordinate position represented by distance to the surface



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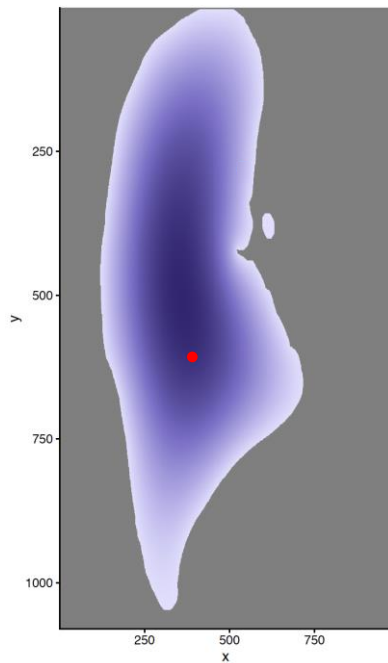
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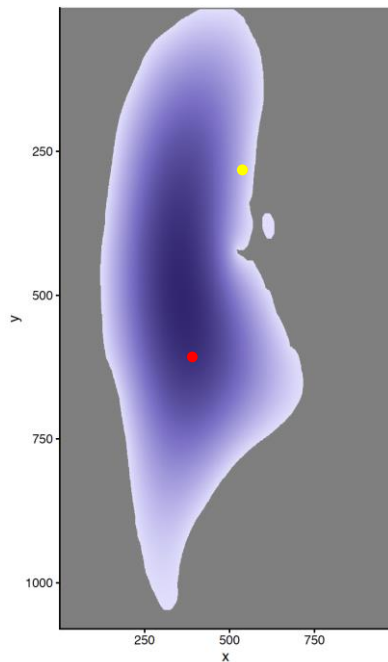
# Poisson distance maps

Poisson Distance  
(Lung 1)



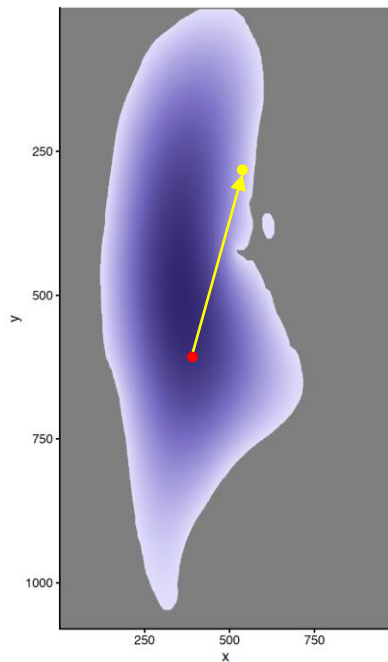
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Poisson Distance  
(Lung 1)



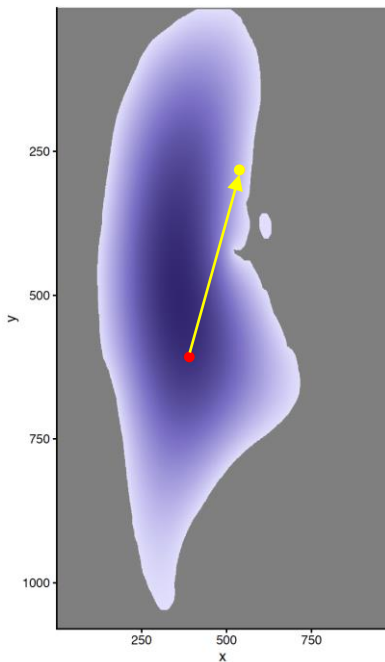
# Poisson distance maps

Poisson Distance  
(Lung 1)

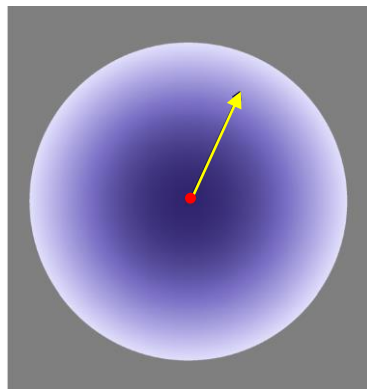


# Poisson distance maps

Poisson Distance  
(Lung 1)

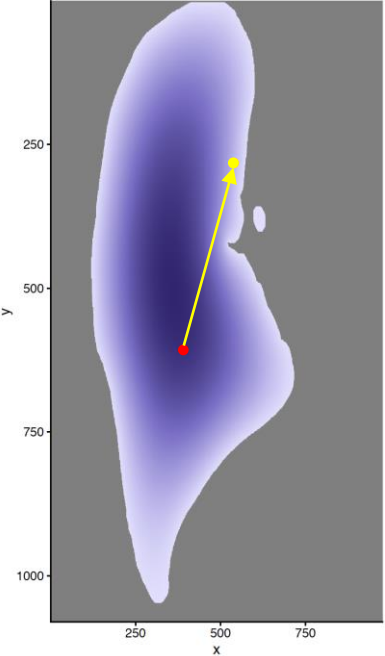


Poisson Distance  
Conformal Mapping

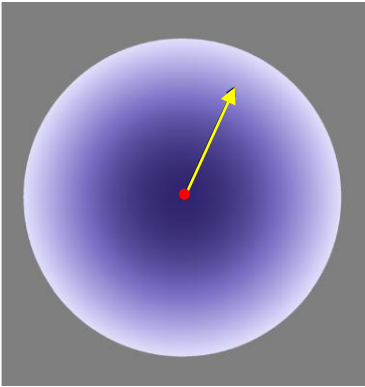


# Poisson distance maps

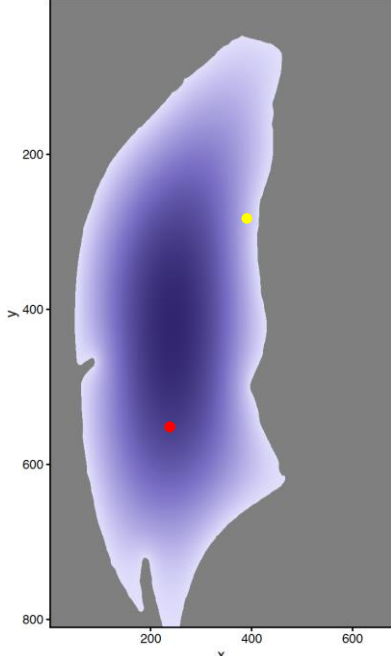
Poisson Distance  
(Lung 1)

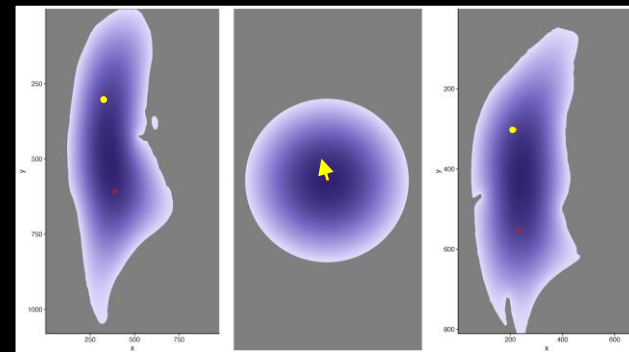
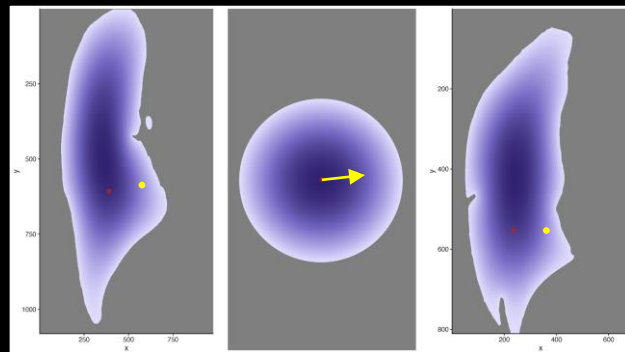
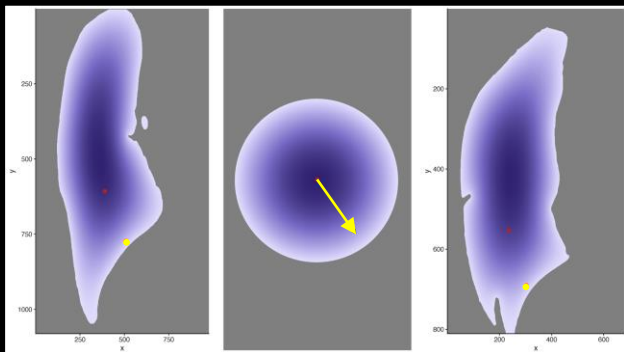
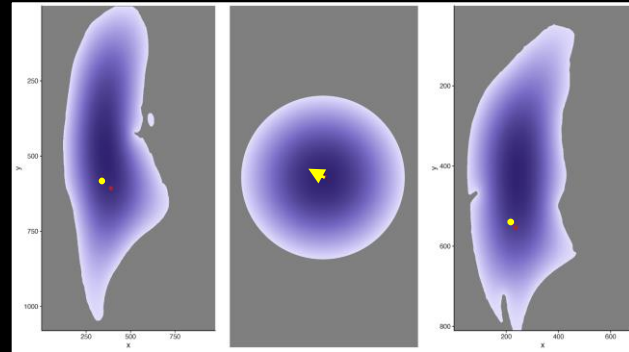
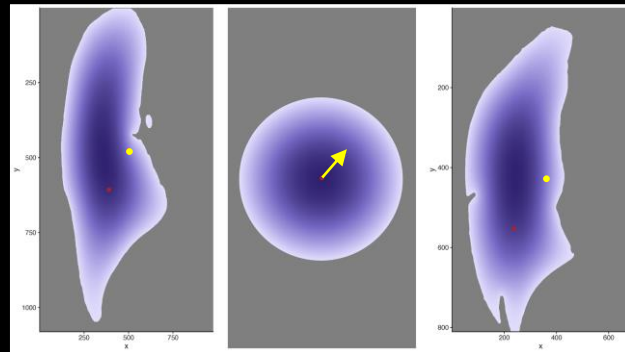
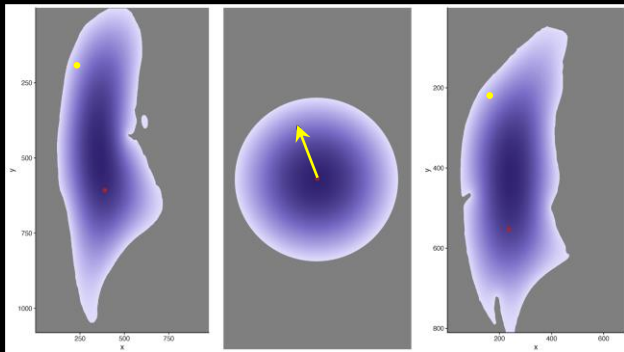


Poisson Distance  
Conformal Mapping



Poisson Distance  
(Lung 2)

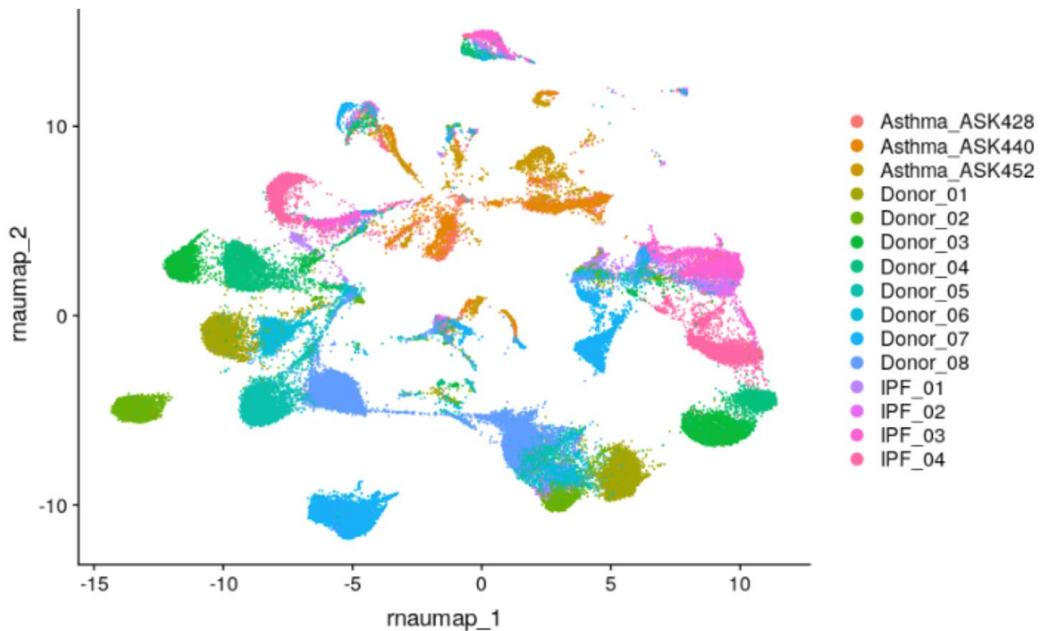




# Single-cell molecular data for the human lung

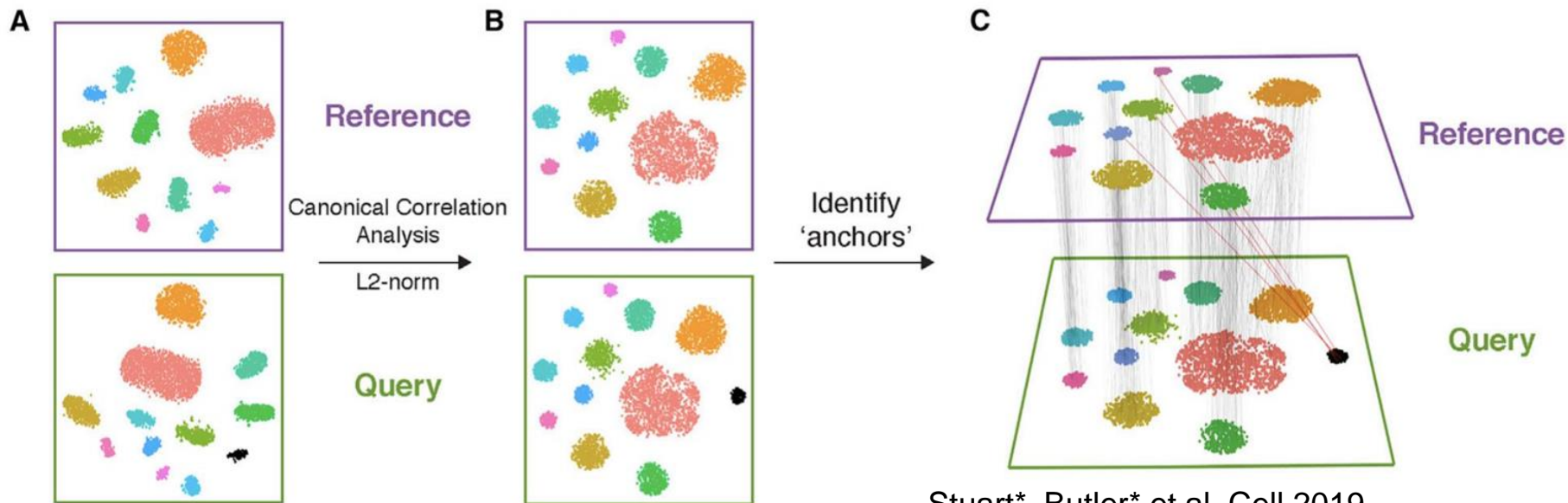
Reyfman et al. 2019; Braga et al. 2019

## Naive



# Integration of single-cell molecular data

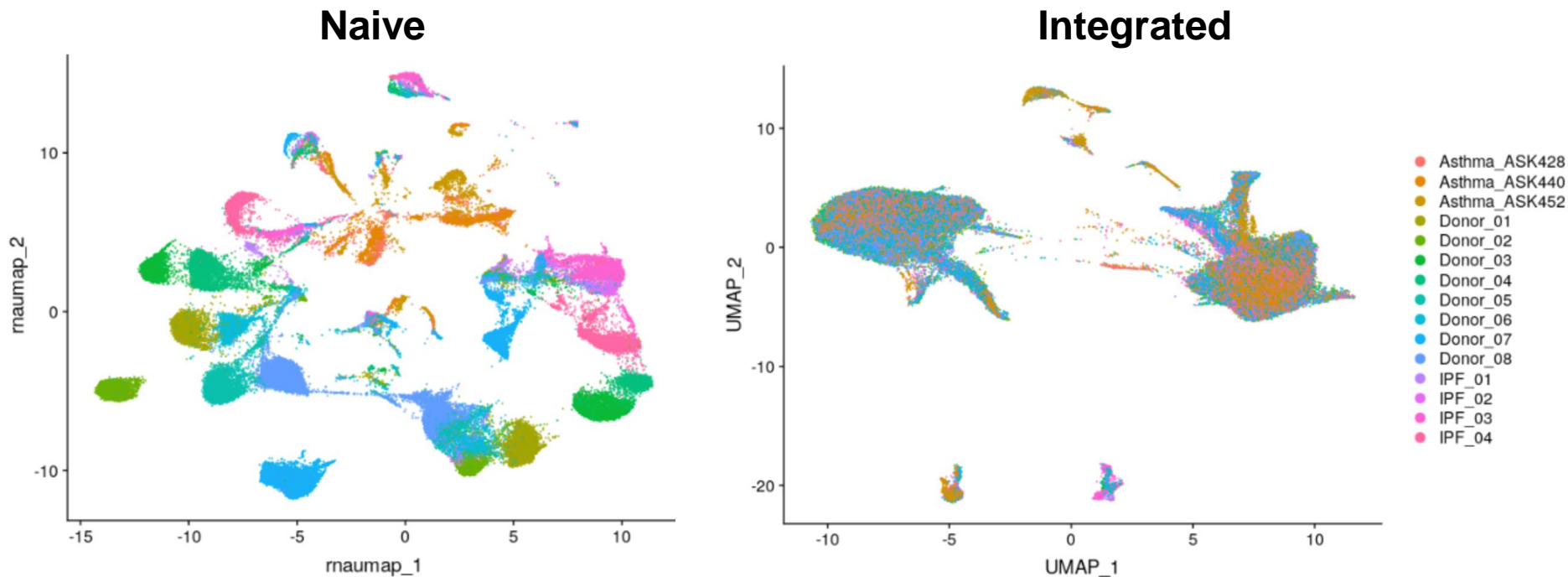
- Shared cell types create anchors
- Multiple datasets harmonized together
- Completely unsupervised





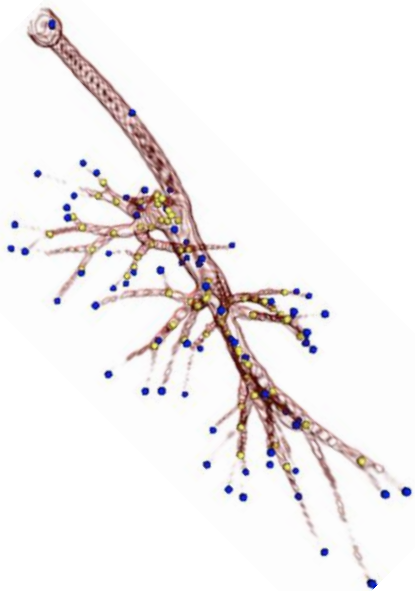
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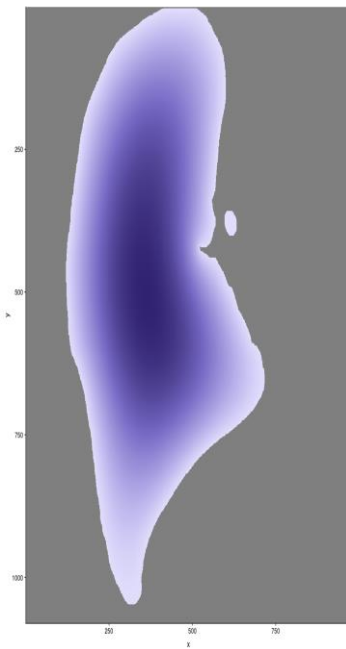


Collaboration with Sanjay Jain, Kun Zhang, Peter Kharchenko

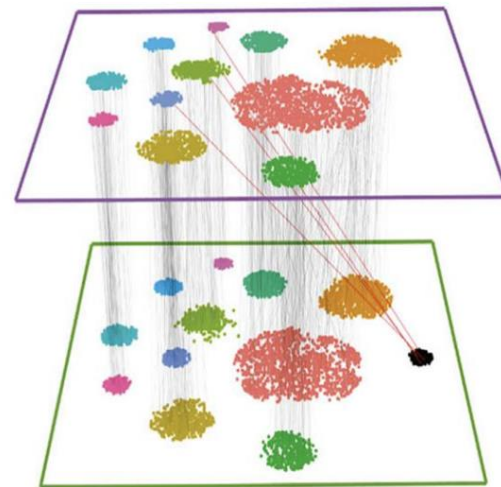
## Internal spatial landmarks



## Shape-based landmarks



## Molecular landmarks



# HuBMAP: HIVE Mapping Components (MC)



Rahul Satija  
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EMBL/EBI



Aviv Regev  
Broad/MIT/HHMI

- **NYGC**

- Bill Mauck
- Andrew Butler
- Tim Stuart

- **EMBL / EBI**

- Eyal Fisher
- Shila Ghazanfar

- **Broad Institute**

- Tommaso Biancalani
- Graham Heimberg

