

# The Human Cell Atlas

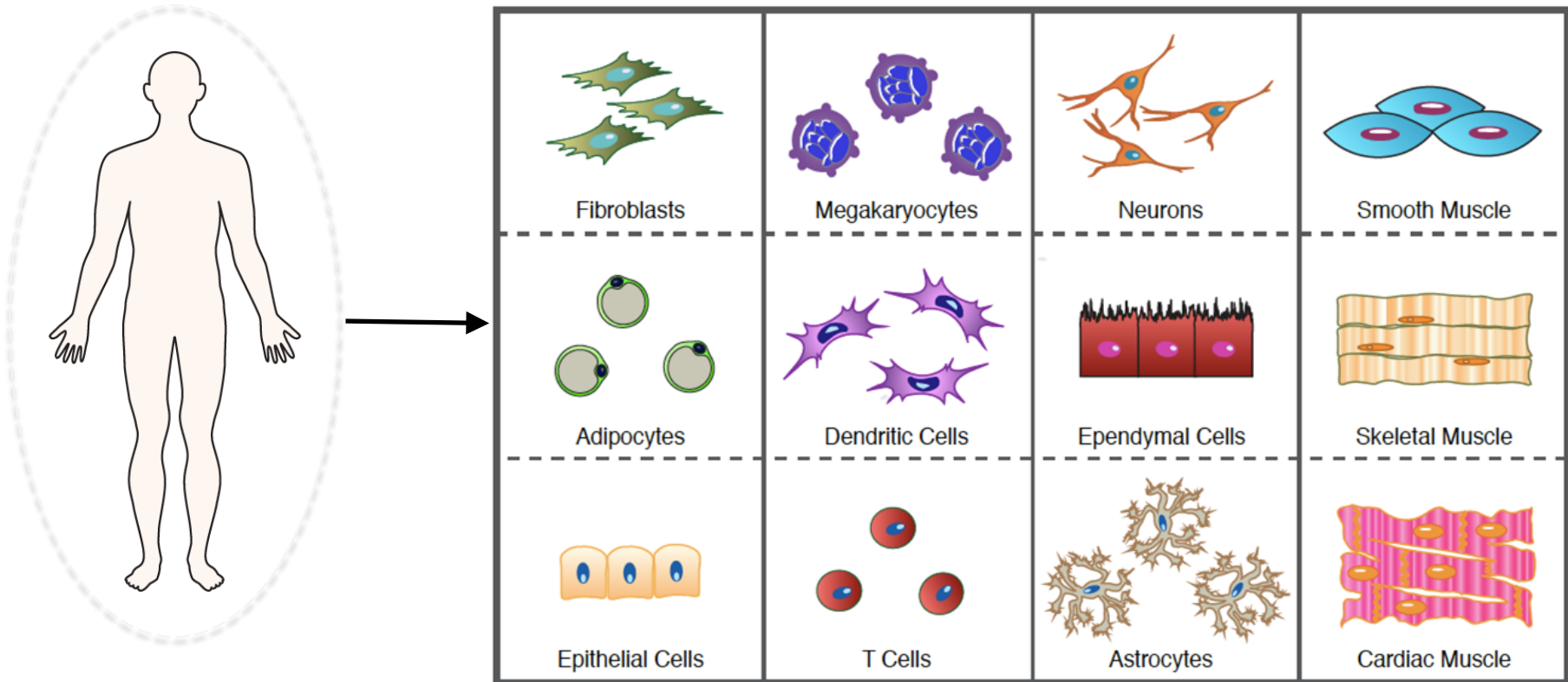
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Department of Biology, MIT

Broad Institute

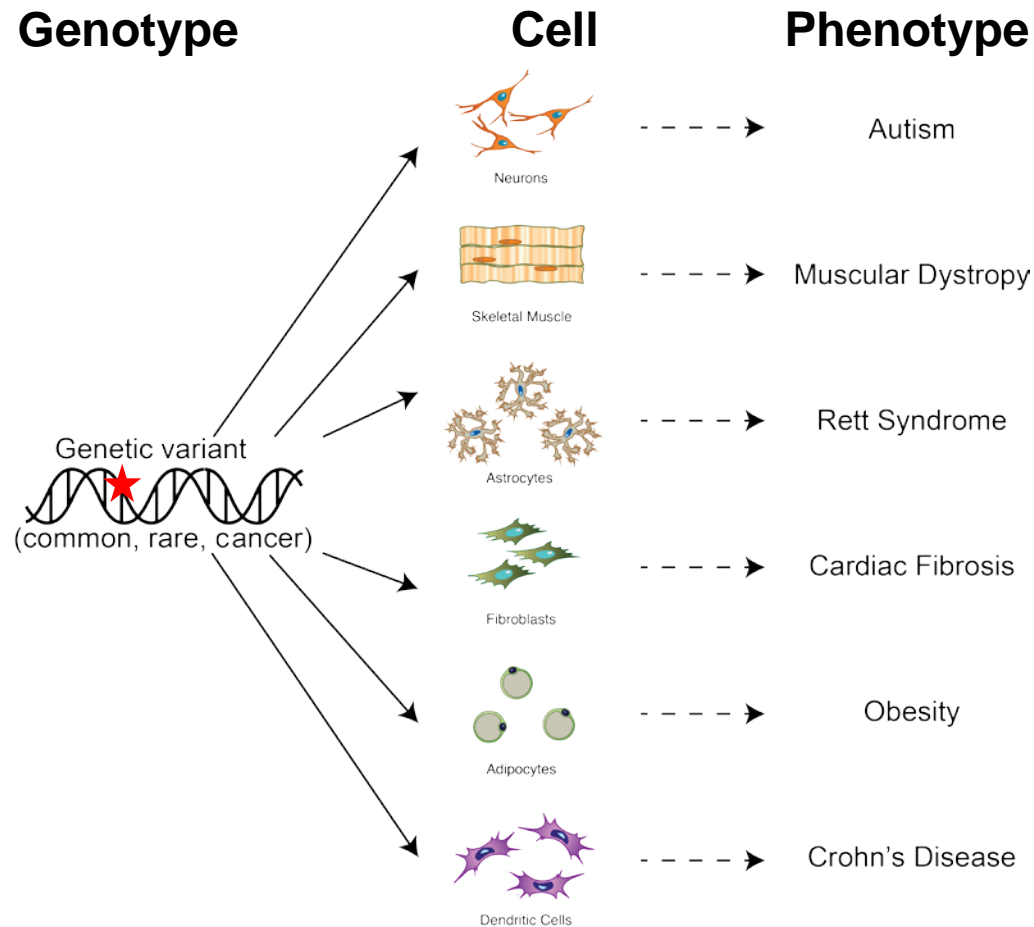
Howard Hughes Medical Institute

# Cells are our core constituents



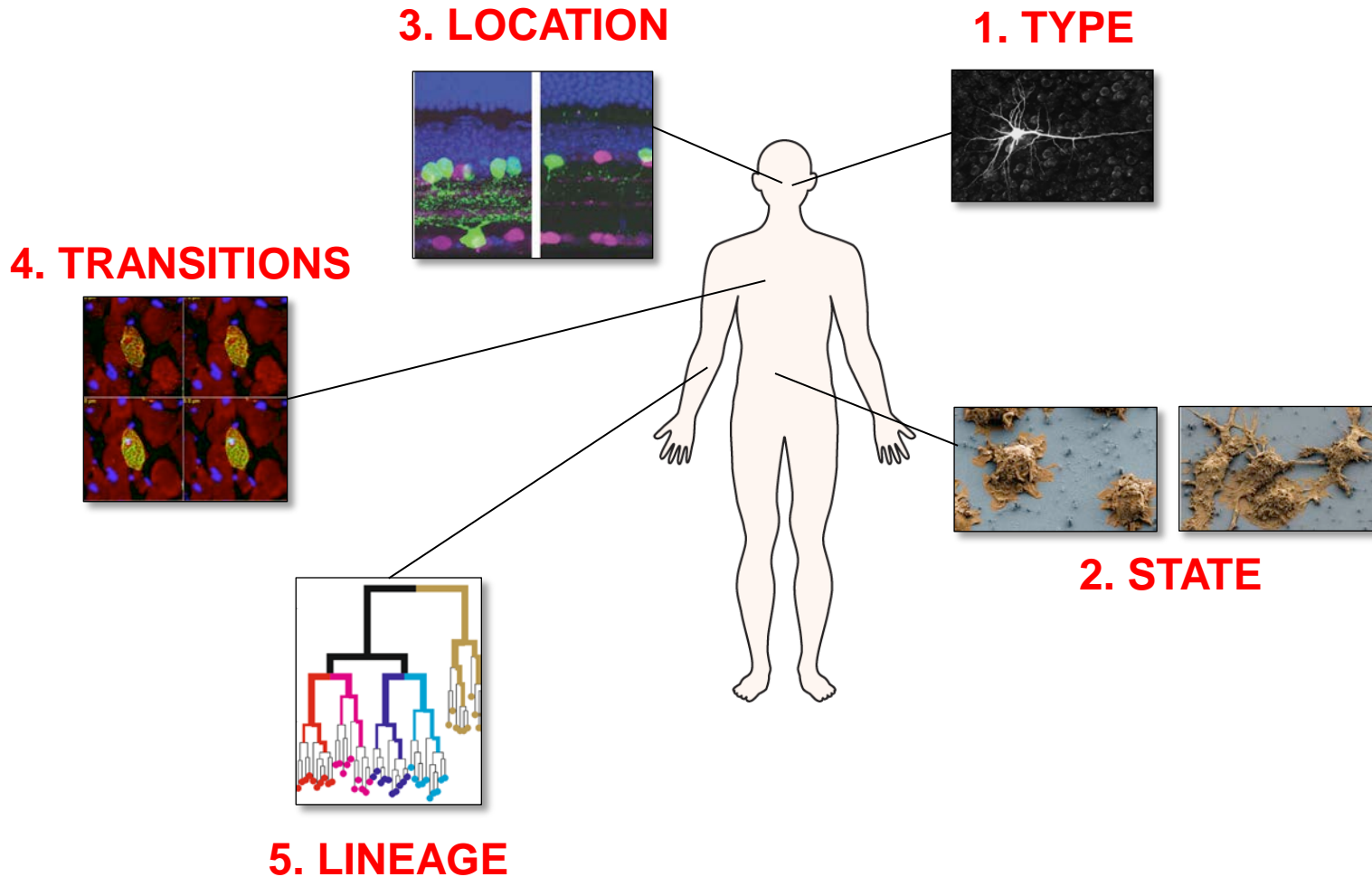
**Cells are classified by characteristic molecules, structures, and functions**

# Cells: a key intermediate from genotype to phenotype



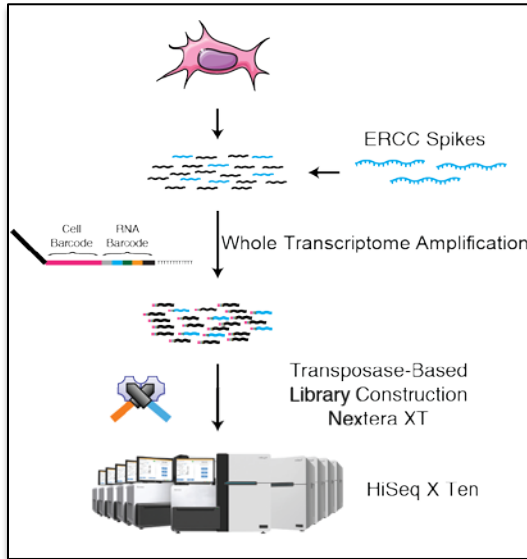
**Knowing our cells is essential for functional dissection of genetic variants**

# What is a cell atlas?

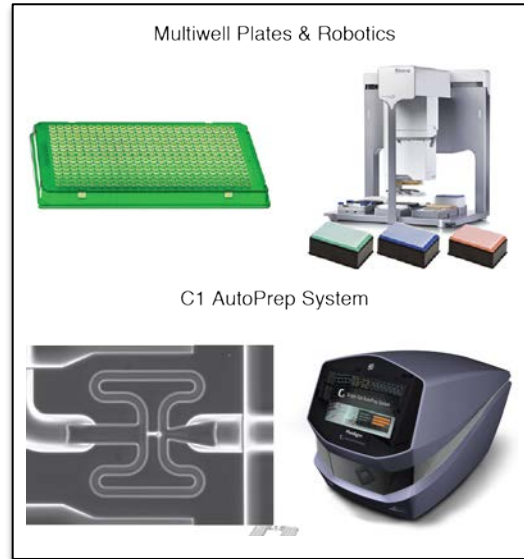


Current knowledge partial, decades and centuries old

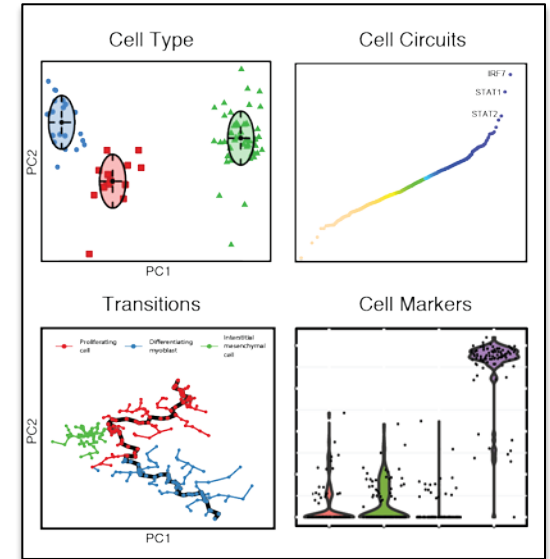
# Single cell genomics makes this possible



✓ Core technology



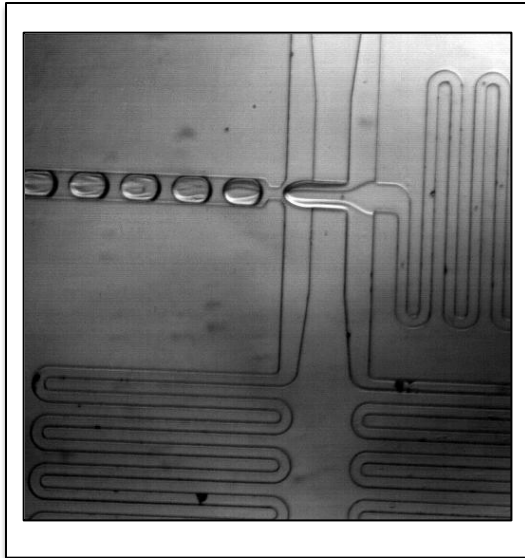
✓ Sample prep



✓ Computation

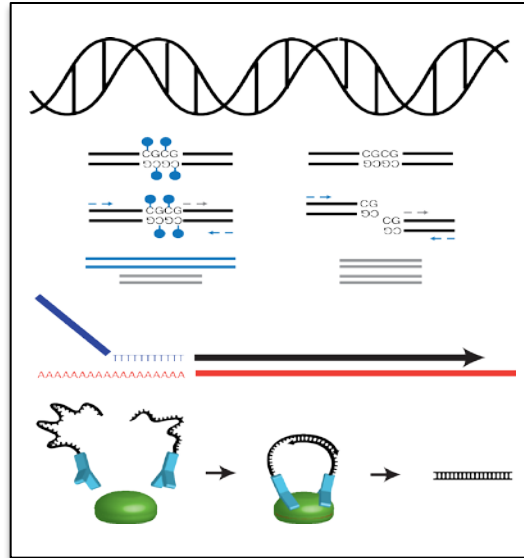
May 2012: 18 cells → July 2014: ~100,000 cells

# Emerging capabilities bring scale and resolution



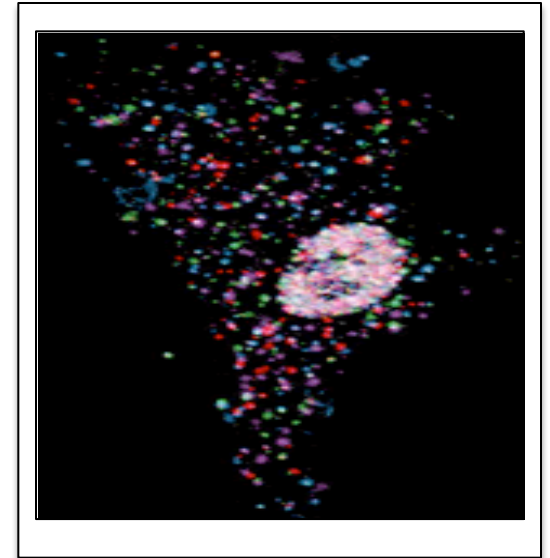
## Scale

5,000 cells/sec;  
ϕ2.8/cell prep



## Measurement

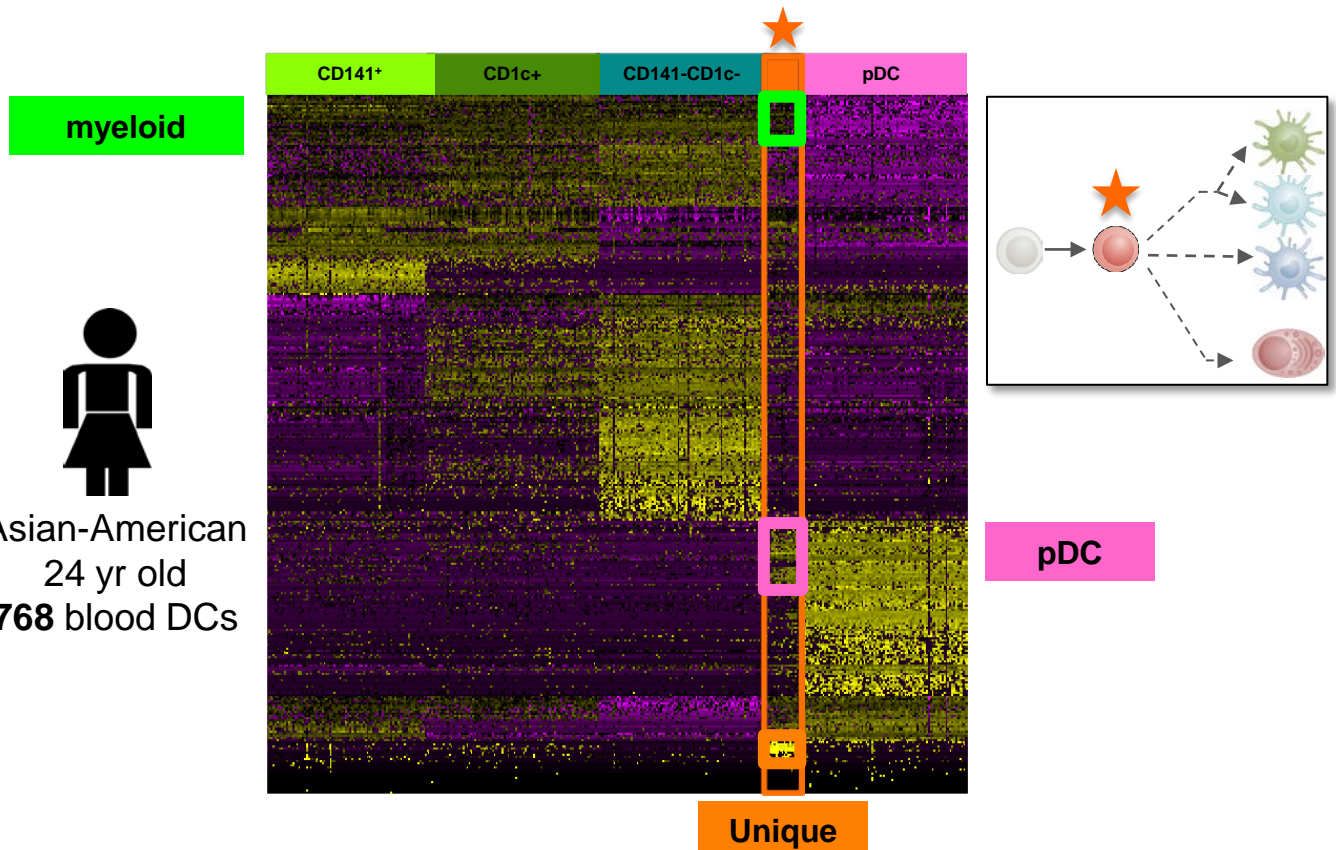
DNA, RNA,  
epigenome, protein



## Location

Registry to 2D, 3D

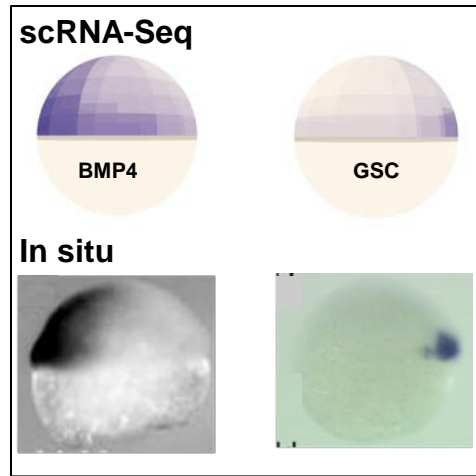
# Already rapidly leading to new insights



- ✓ New population: 1.9-3.2% of DCs; 0.04-0.064% of PBMCs
- ✓ Validated in 10/10 independent individuals
- ✓ Placed in lineage: a blood progenitor?

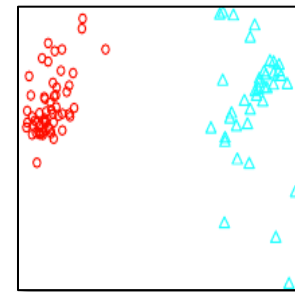
# Already rapidly leading to new insights

## LOCATION



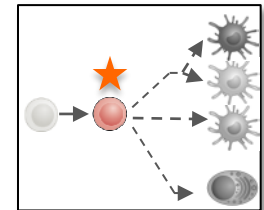
Fish embryo

## TYPE



Retina neurons

## LINEAGE

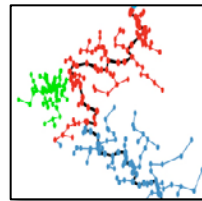


New human immune blood progenitor

## TRANSITIONS

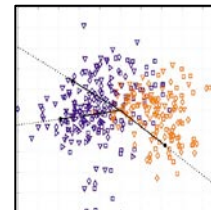


'Early responder' immune cells

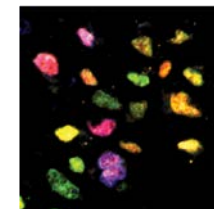


Muscle cell differentiation

## STATE



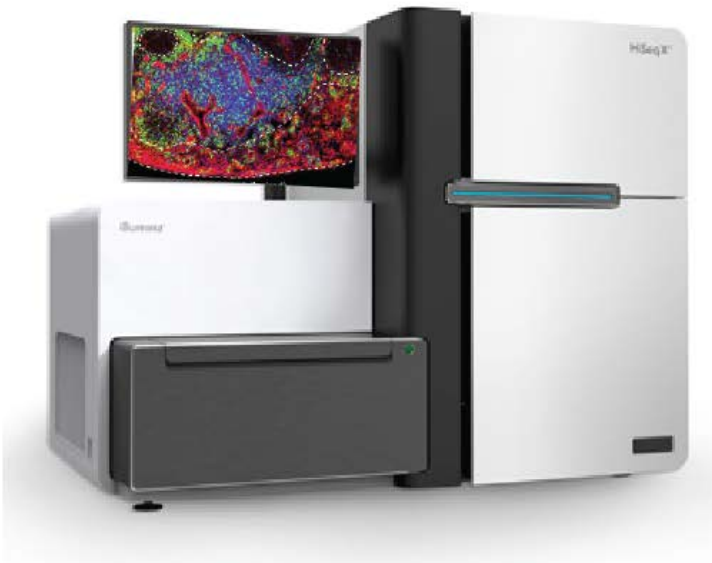
Pathogenic T cells in MS model



CSC spectrum in glioma



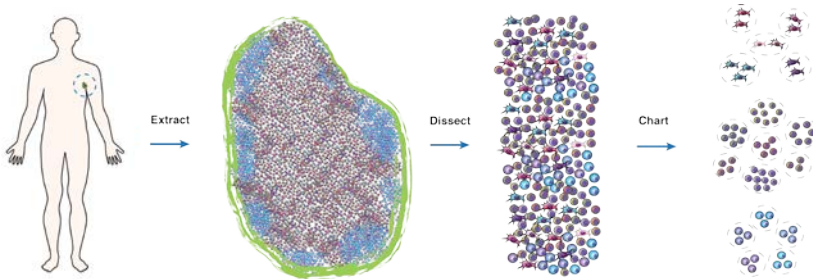
# The sequencer: a microscope for the 21<sup>st</sup> century



# The Human Cell Atlas Project

## What? A census of all cells

- **Types**
- **States**
- **Transitions**
- **Location**



## Why? A reference map for function

- **Foundational, fundamental knowledge**
- **Function of genetic variants**
- **Heterogeneity in disease**
- **Eventually characterize in individual patients**

# The Human Cell Atlas Project

## How? A unified project

- **Pilot project in complementary systems (e.g., blood, gut, liver)**
- **Consortium with expert communities**
- **Standard, controlled process**
- **Shared analytical tools**
- **Drive costs to ~\$0.15/cell\***

## Why? Standardized and impactful

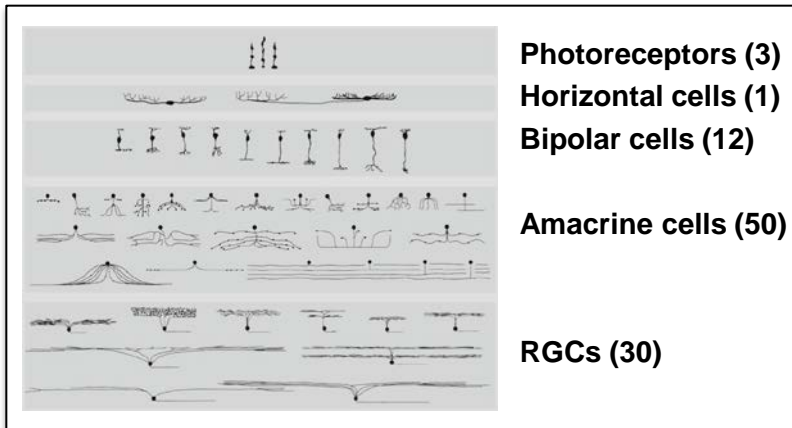
- **Managed: Only standard process ensures we are not deceived by noise**
- **Scale: Drive cost down**
- **Technology advancing: Novel sample prep, cell isolation, analytical tools**
- **Resources for entire community**
- **Commensurate with clinic**

\* with drops,  $10^5$  reads/cell, @ \$0.4-0.6/cell with HiSeq X10 yields

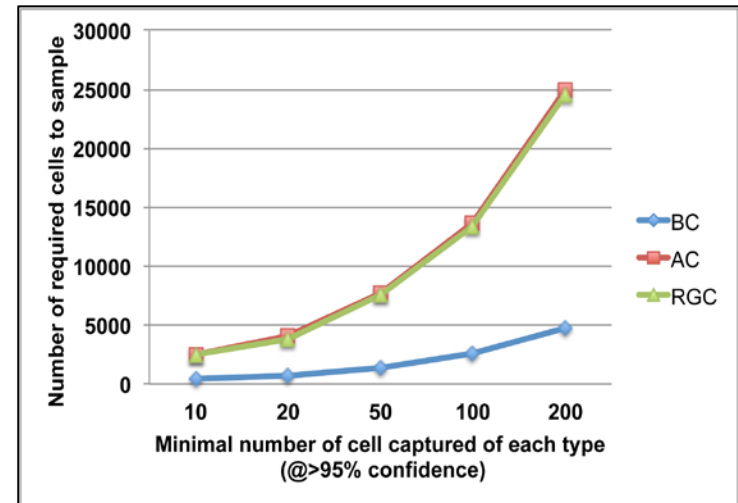
# The Human Cell Atlas Project

## Large

- Adult human:  $\sim 2 \times 10^{13}$  cells (excluding red blood cells)
- 300 'major' cell types
- ....but,  $\sim 100$  sub-sub-types just of retinal neurons



## ... but finite



**150M** neurons in retina,  
**~40K** required for survey

Retinal bipolar cell (BC): 12 sub types, rarest @5%  
Retinal amacrine cell (AC): 50 subtypes, rarest @1%  
Retinal Ganglial cell (RGC): 30 subtypes, rarest @1%



