

Architecture as defined by
organization of niches and
INTER-CELLULAR relationships

PATHOLOGY

From the Atomic Scale

‘ON-UP’

Garry P. Nolan, Ph.D.

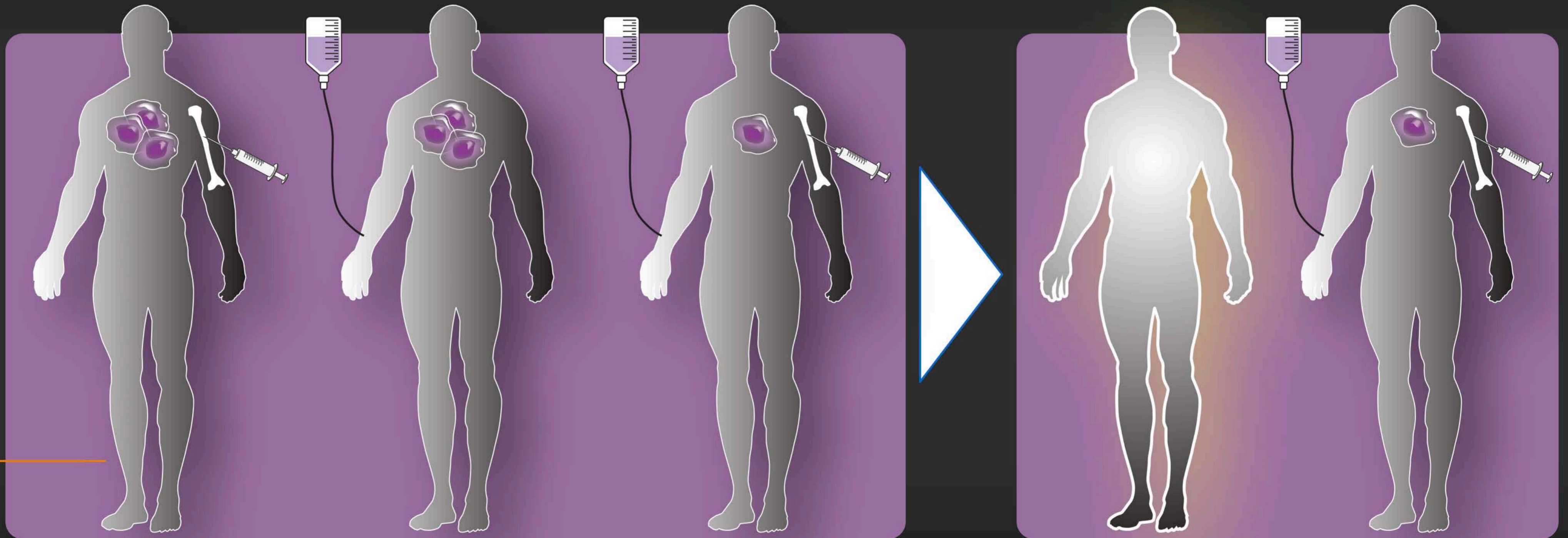
Rachford and Carlota A. Harris Professor
Dept. of Microbiology & Immunology
Stanford University

Parker Institute for Cancer Immunotherapy

ORDERING IN PSEUDOTIME

Can ordering help us with predictors of relapse?

B-cell precursor acute lymphoblastic leukemia (BCP-ALL)



Diagnosis

Treatment

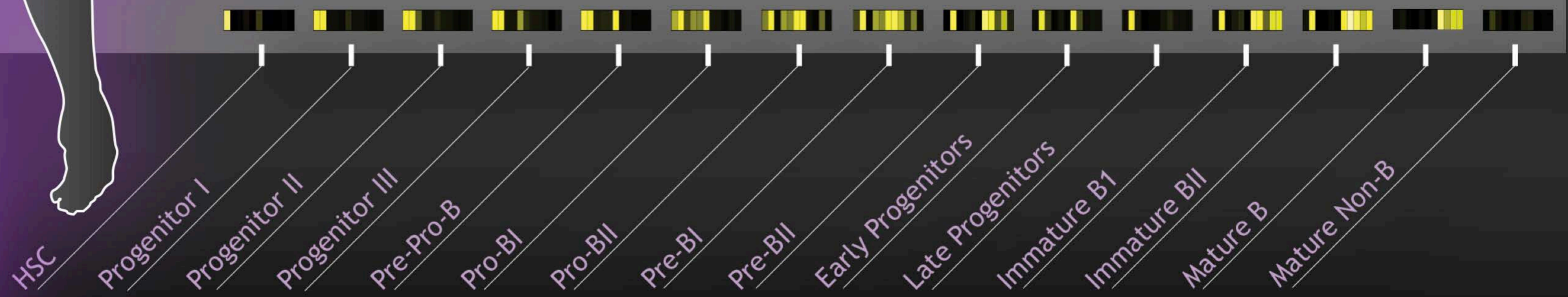
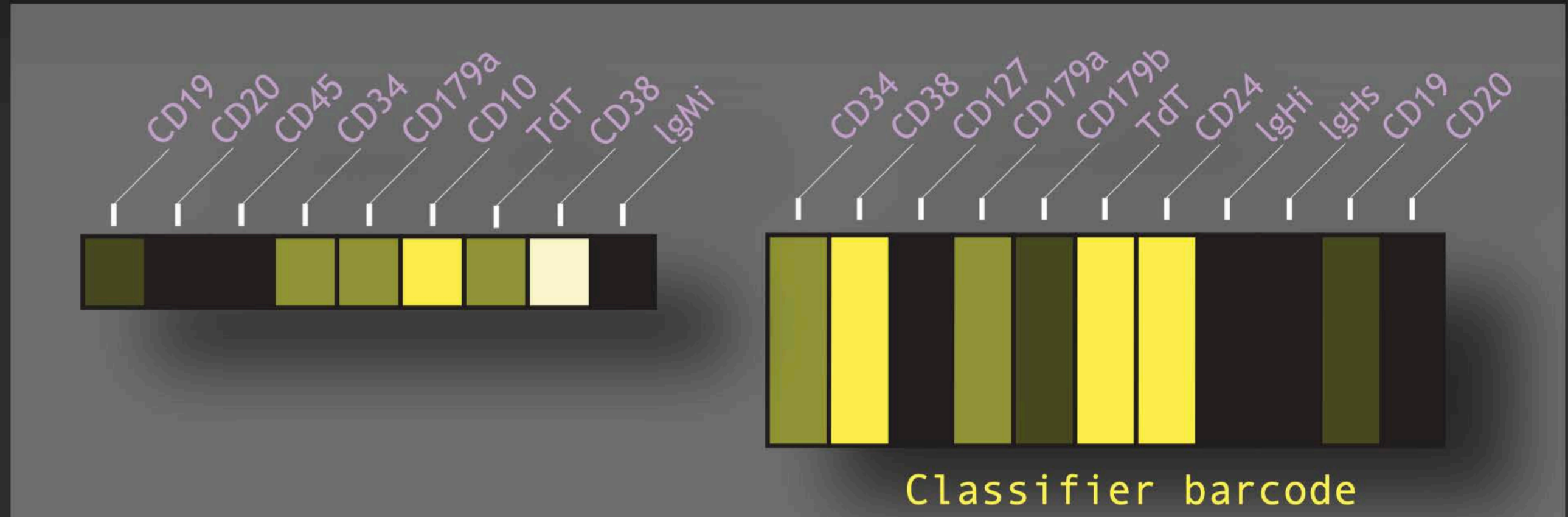
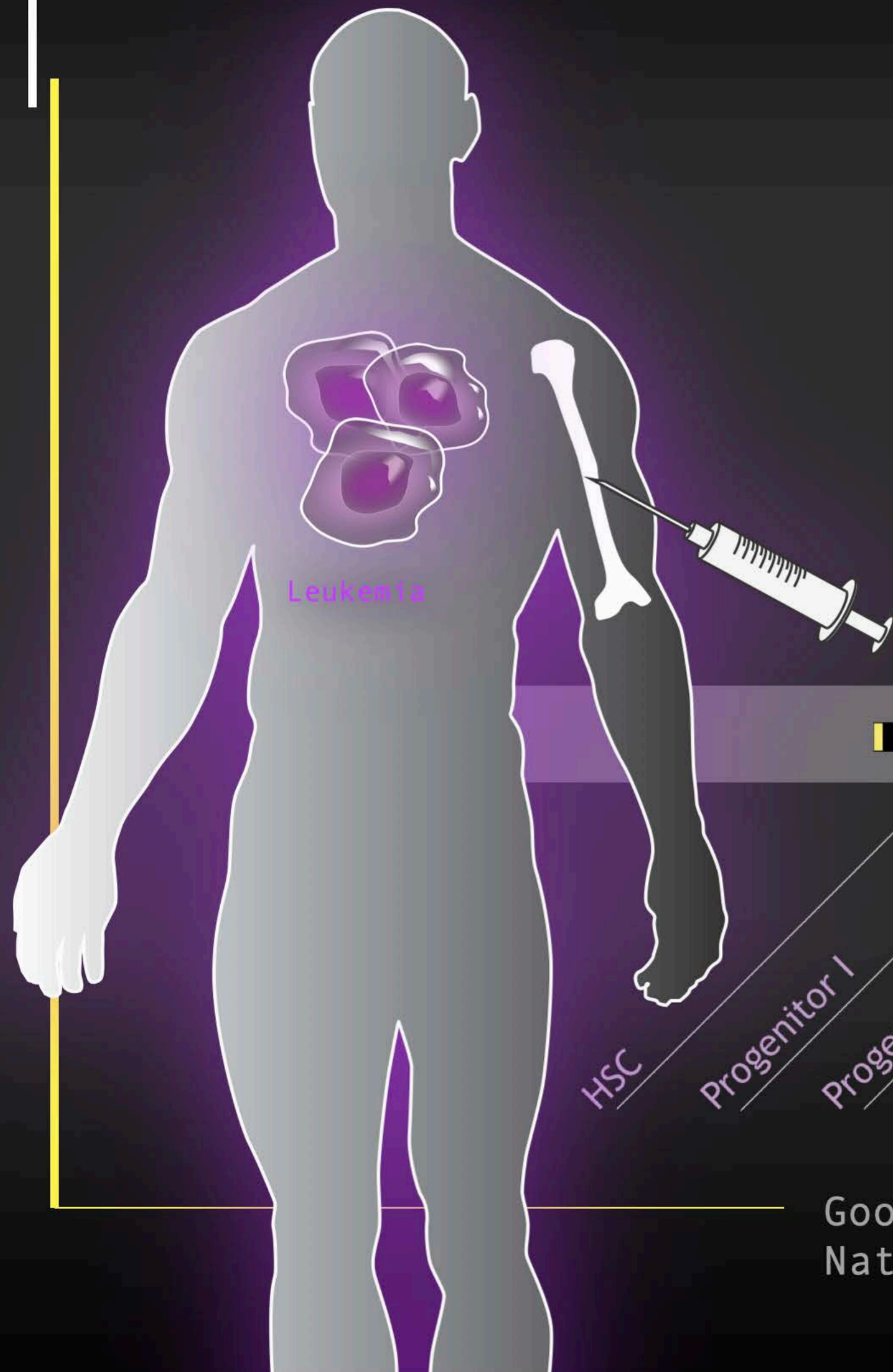
Assess MRD

80% Cure

20% Relapse

MAPPING LEUKEMIA CELLS

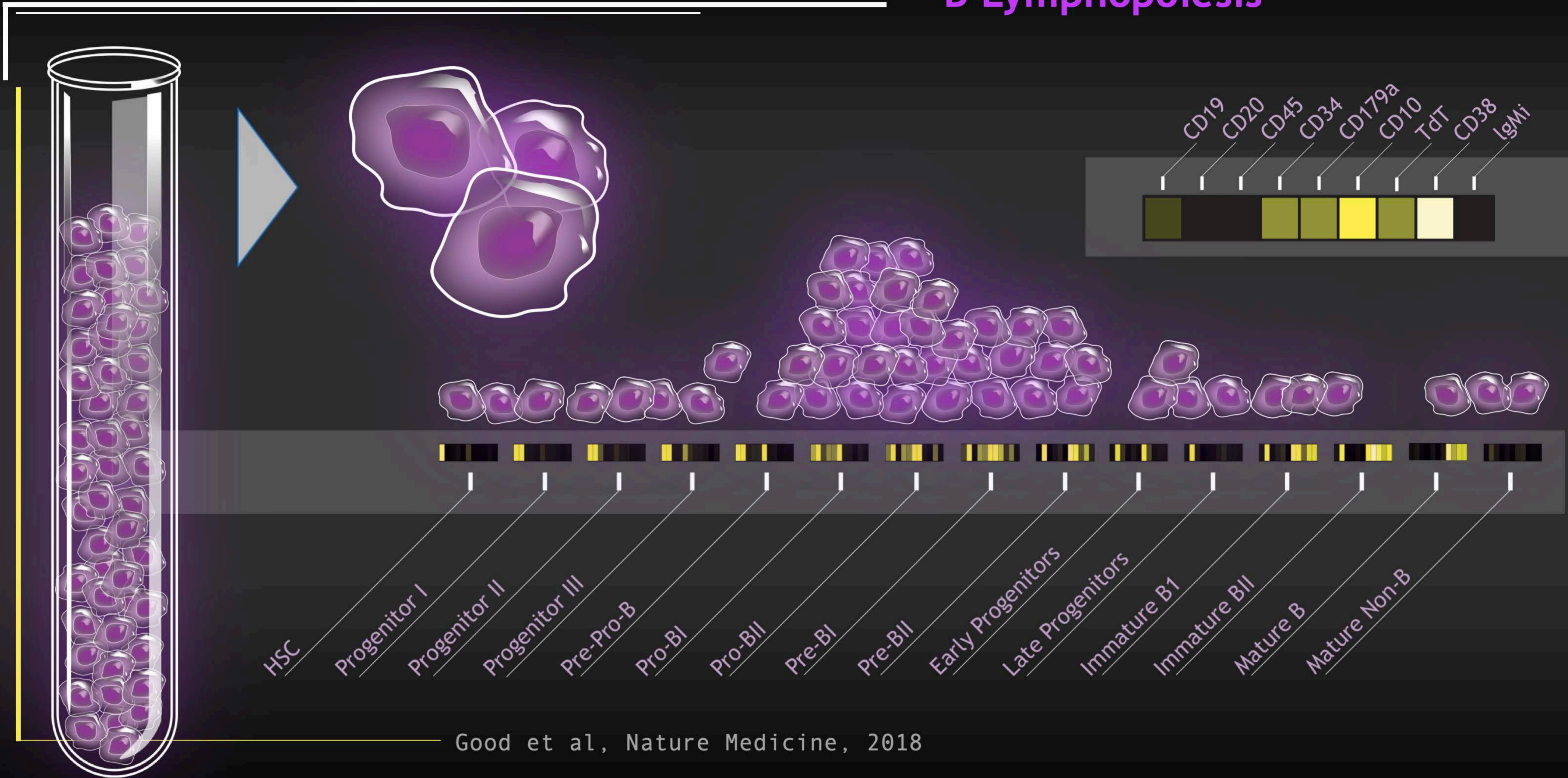
Closest Normal Stage in B Lymphopoiesis



Good et al,
Nature Medicine, 2018

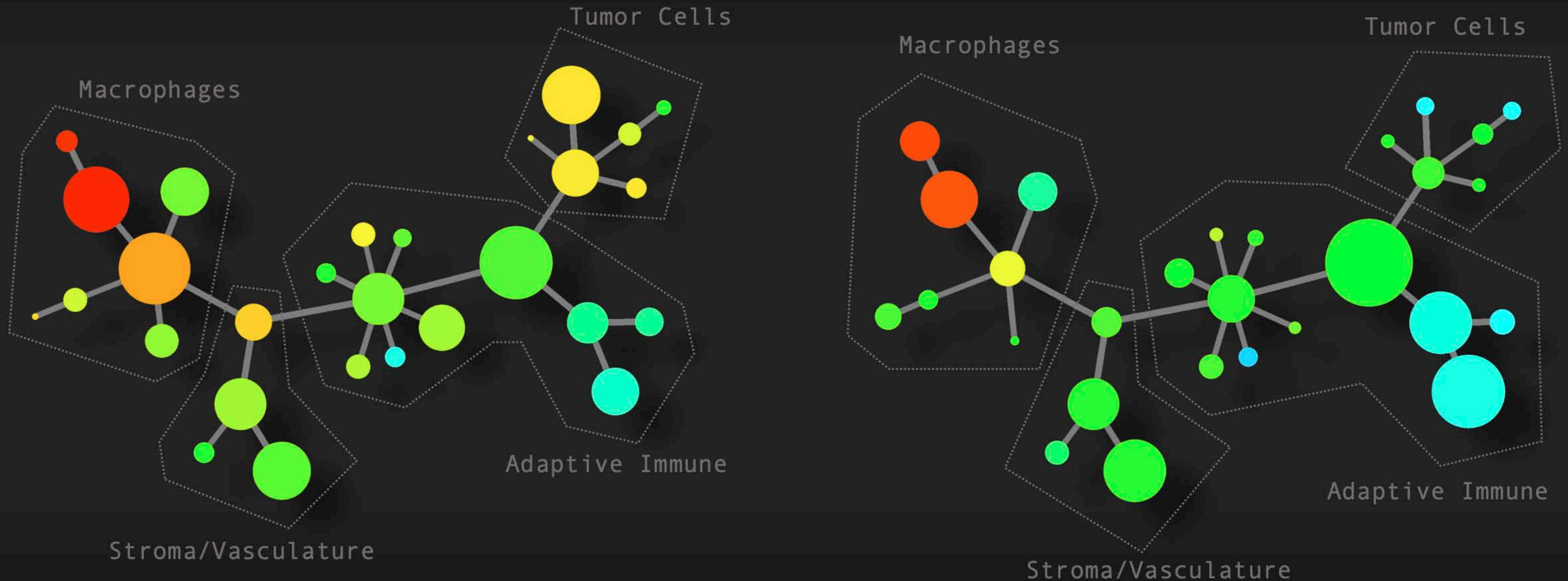
MAPPING LEUKEMIA CELLS

Closest Normal Stage in B Lymphopoiesis



Good et al, Nature Medicine, 2018

CELL CLUSTERING • Brentuximab @CD30, response in CTCL

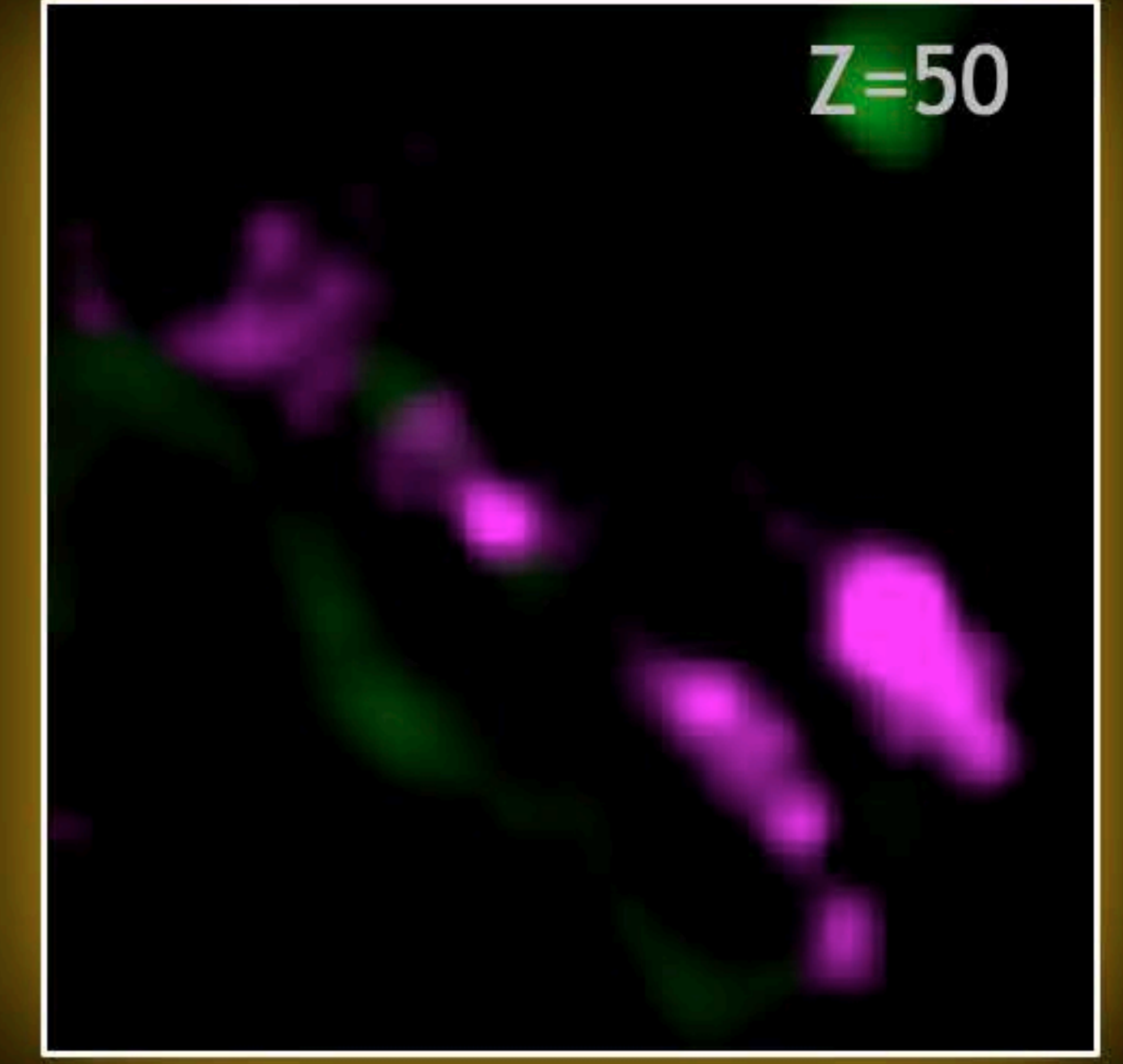
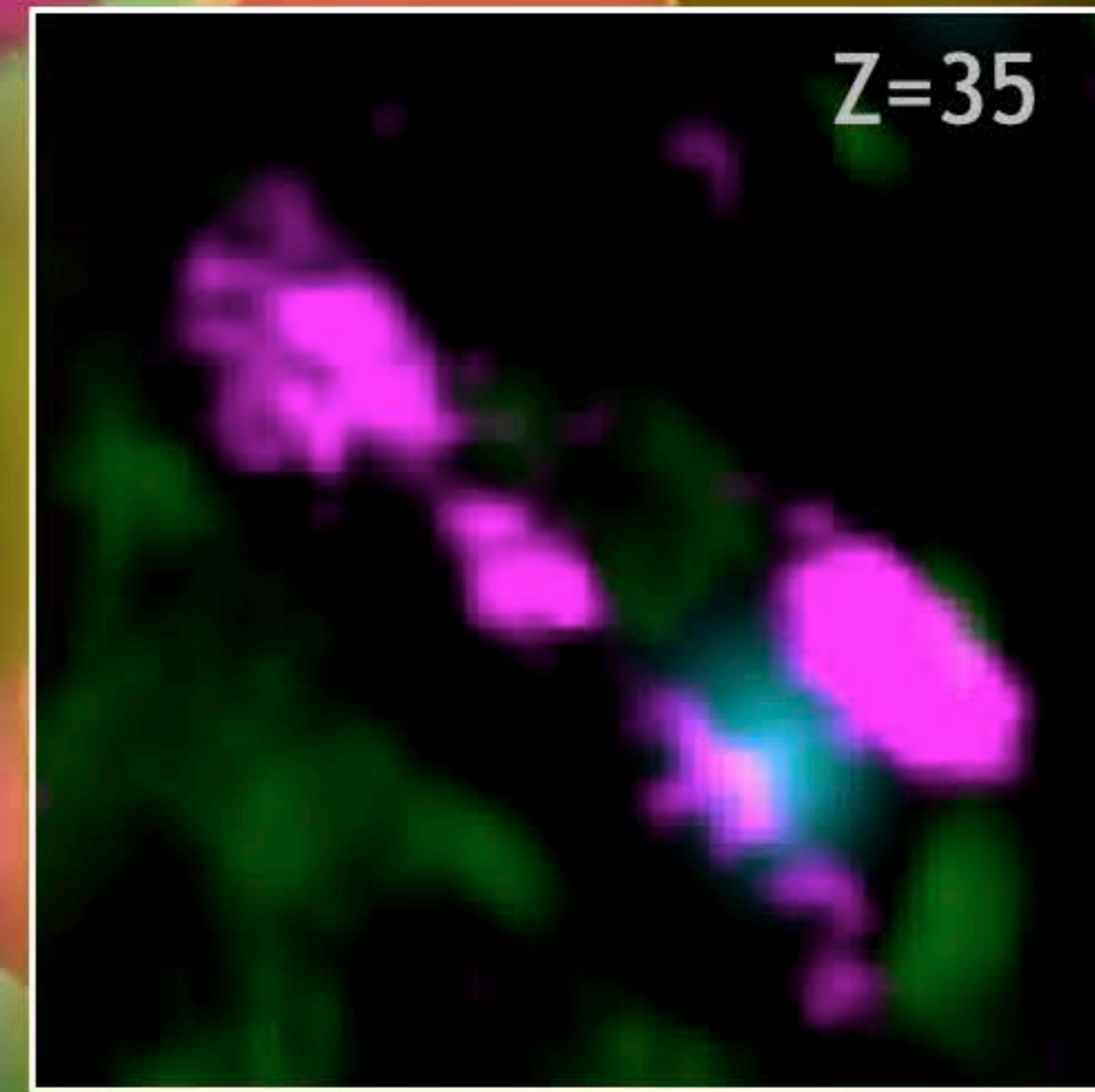
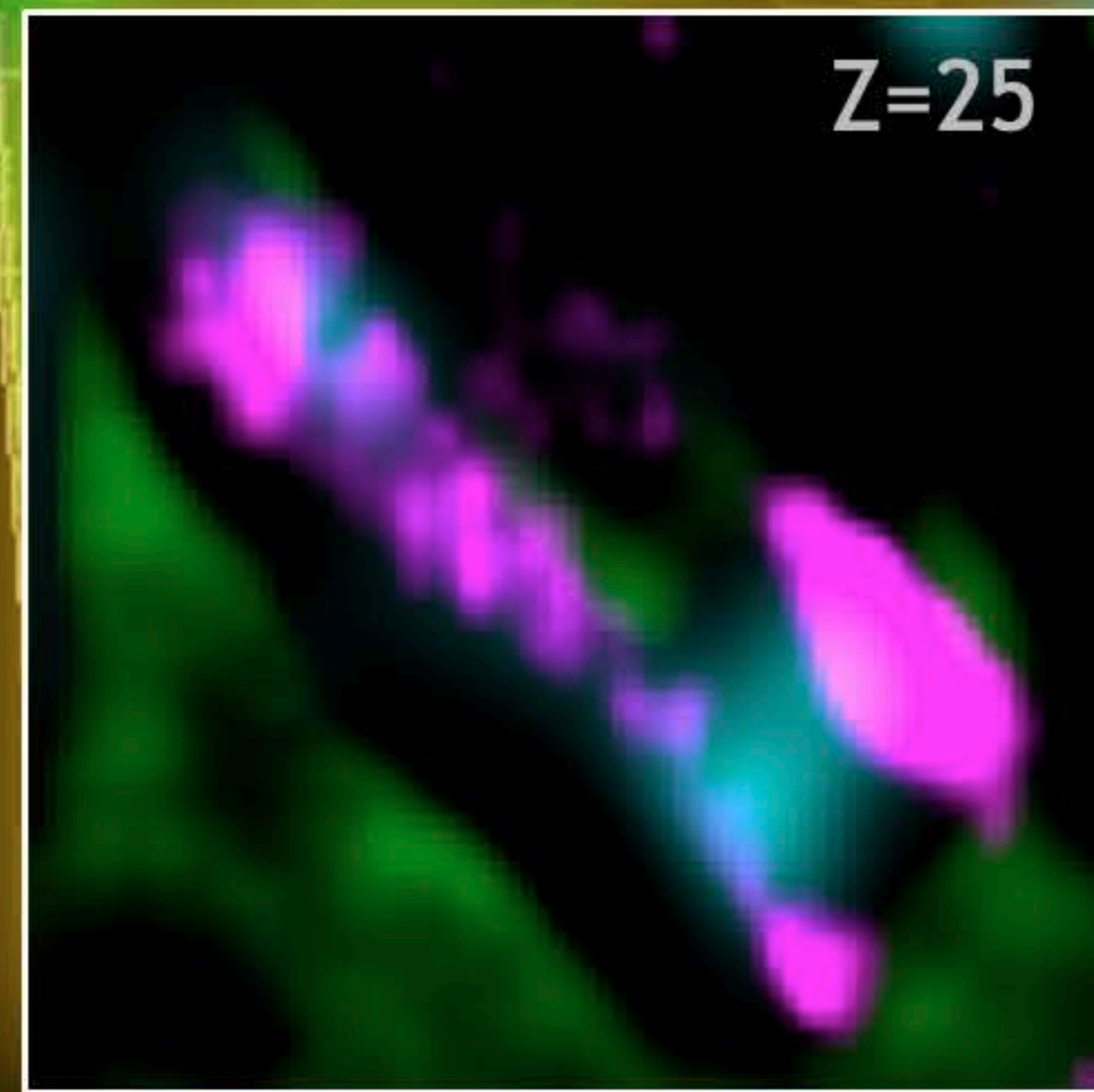
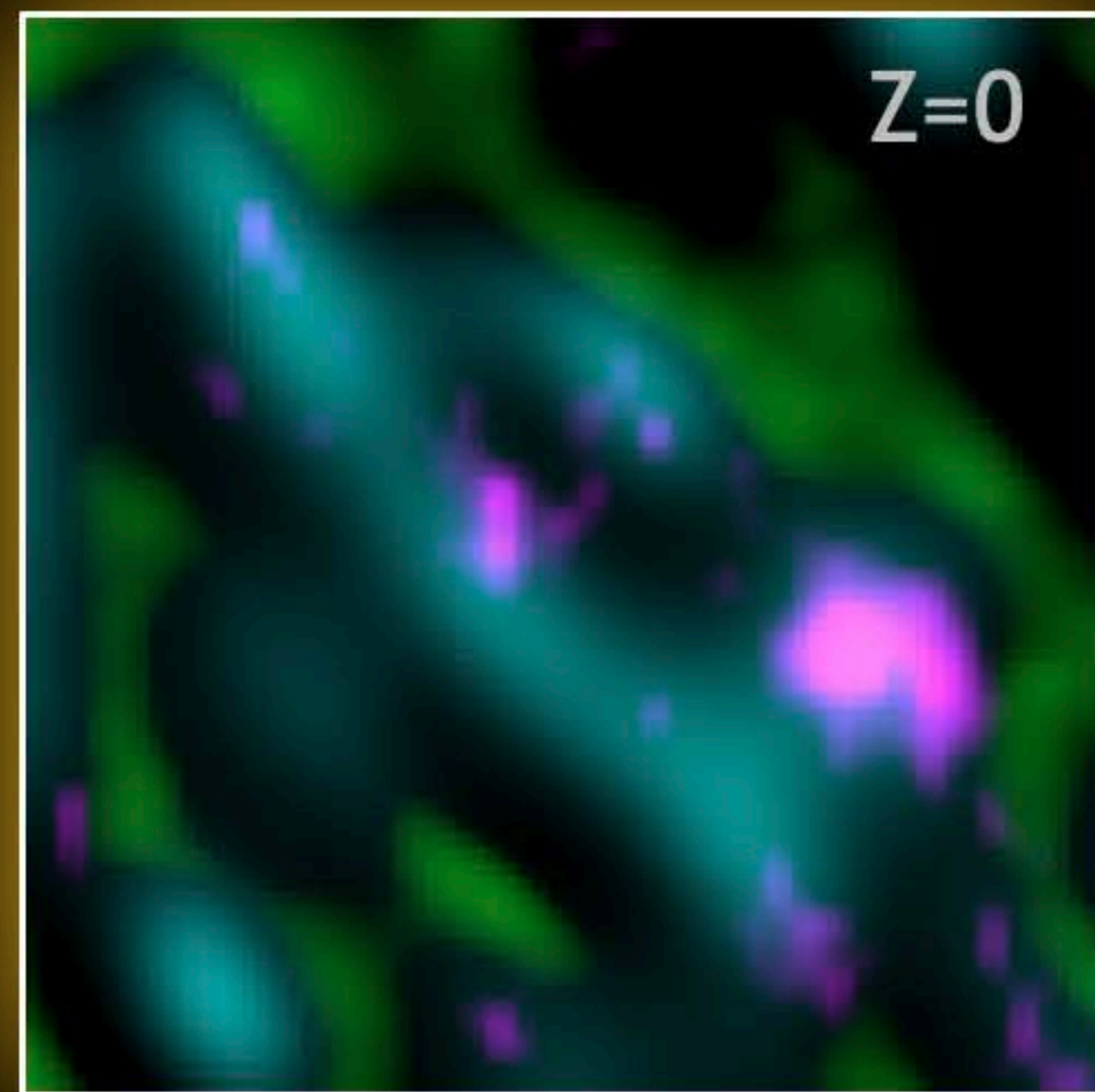


• Pre-treatment

Post-treatment (responder)

3D NANO-IMAGING • Mitochondria & Protons

DR Congo
1995 (Kikwit)



Single Mitochondria

Nitrogen $^{12}\text{C}^{14}\text{N}$ Tom20 ^{197}Au Proton $^{12}\text{C}^1\text{H}$

CTCL Plaque •

Pan Keratin

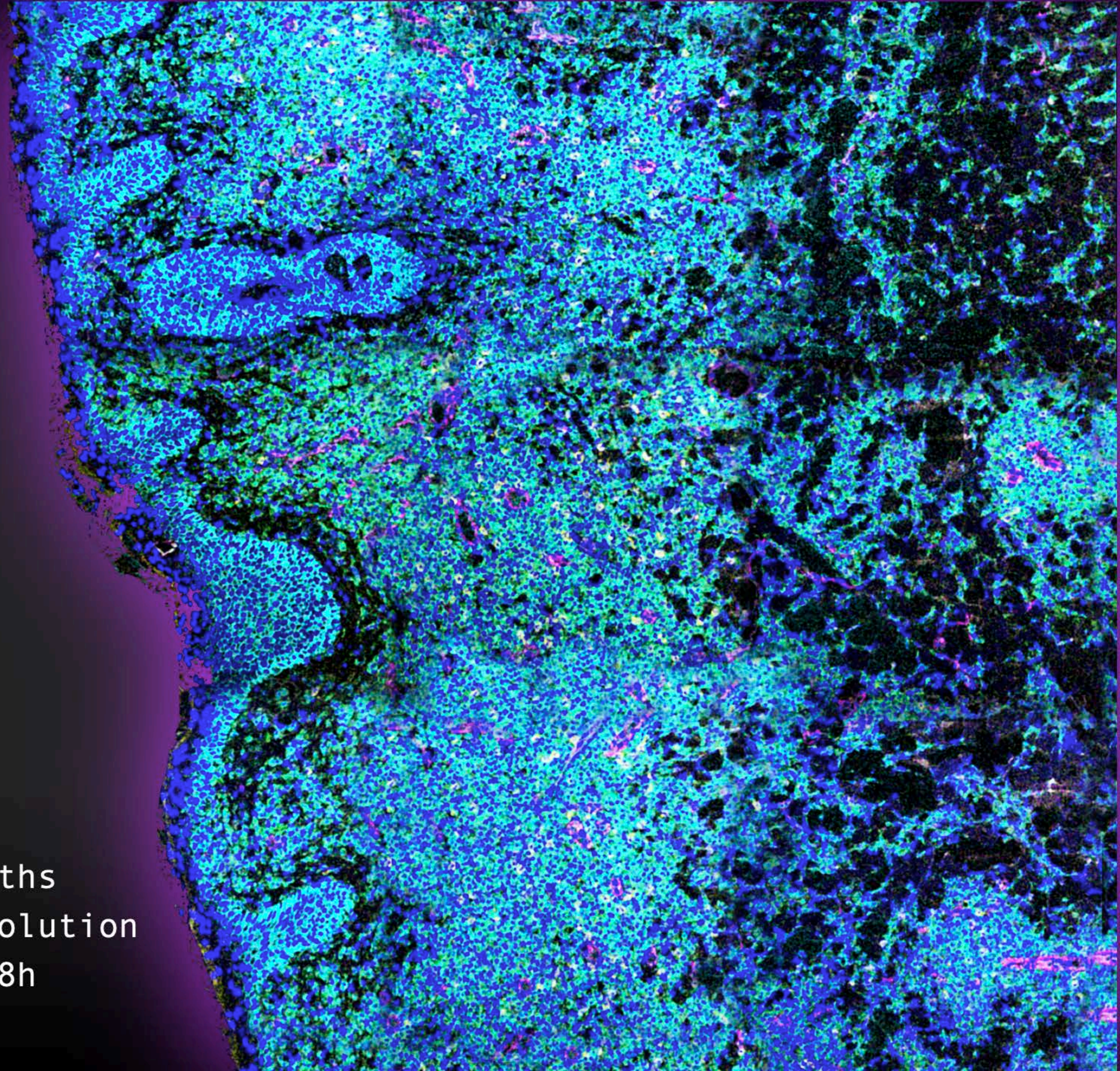
Vimentin

HLA-DR

dsDNA

CD8

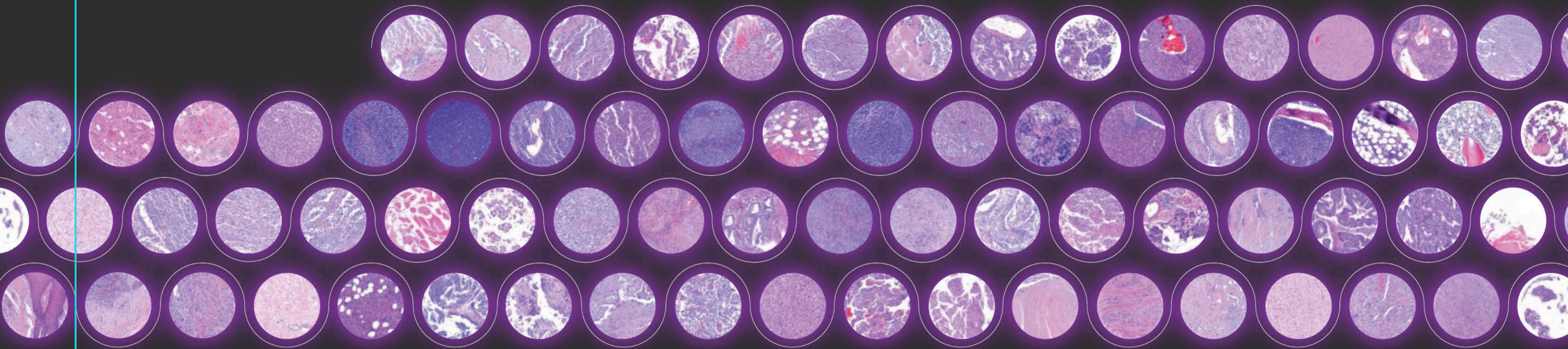
- 1.5mm * 1.5mm
- 42 Channels
- 3 x 3 F0Vs, 3 depths
- 300-500nm X-Y resolution
- Time to acquire: 8h



EXPANDING CODEX •

Christian Schuerch
Darci Philipps

TISSUE ATLAS



FFPE



'Studying Malignancies, One Cell at a Time'

- *ADVANCED PROTEOMIC TECHNOLOGIES* -

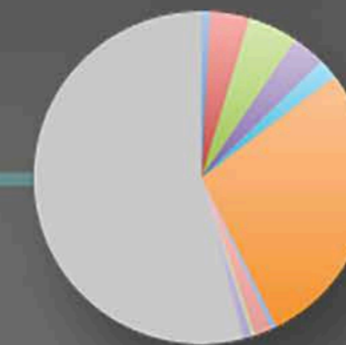
VARIATION • Frequencies of Immune Cell Subsets

- B Cells
- CD4 T Cells
- CD8 T Cells
- NK Cells
- Myeloid DC
- Macrophages
- Monocytes
- Non Classical Monocytes
- Basophils
- Granulocytes
- Unassigned Cells

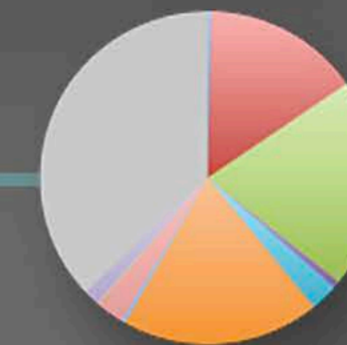
CD45+
3-18%



R1118



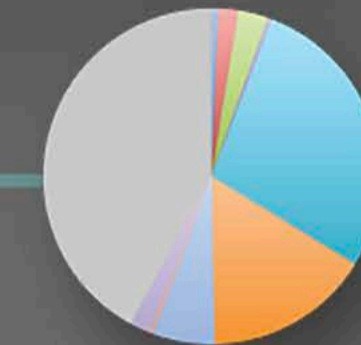
X2643



X2619

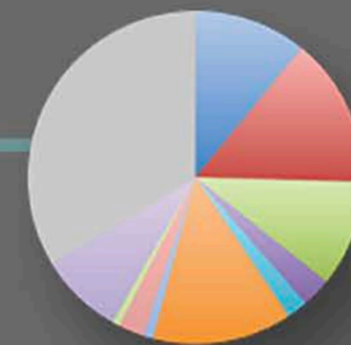


Z378

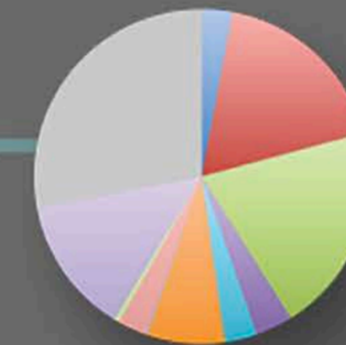


R1116

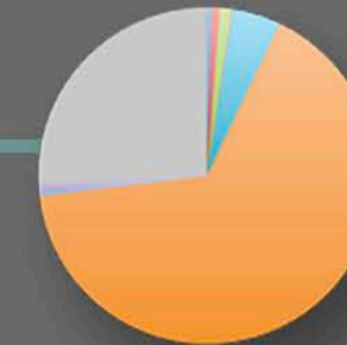
CD45+
20-60%



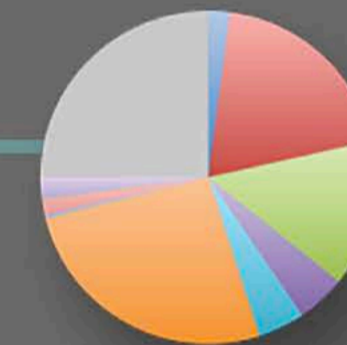
Z500



Z536

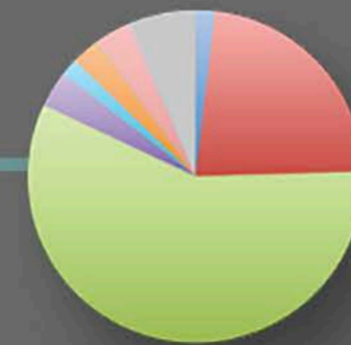


X2661



X2638

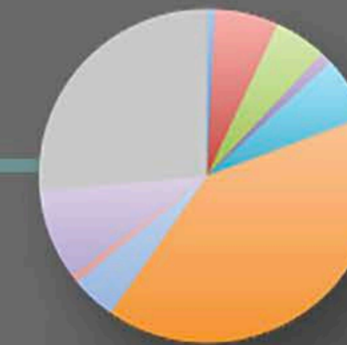
CD45+
60-75%



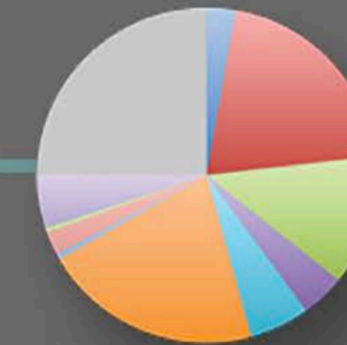
X2667



Z289

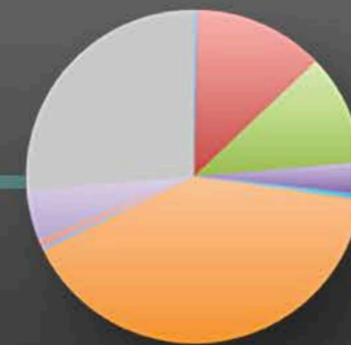


X2617



X2648

CD45+
80-95%



Z403



Z467



Z377



R1153

High Parameter

IMMUNE PROFILING • Host Responses to Infection

From Blood to Tissues

David McIlwain, PhD

NIH | Center for Human Immunology
National Institute of Allergy & Infectious Disease



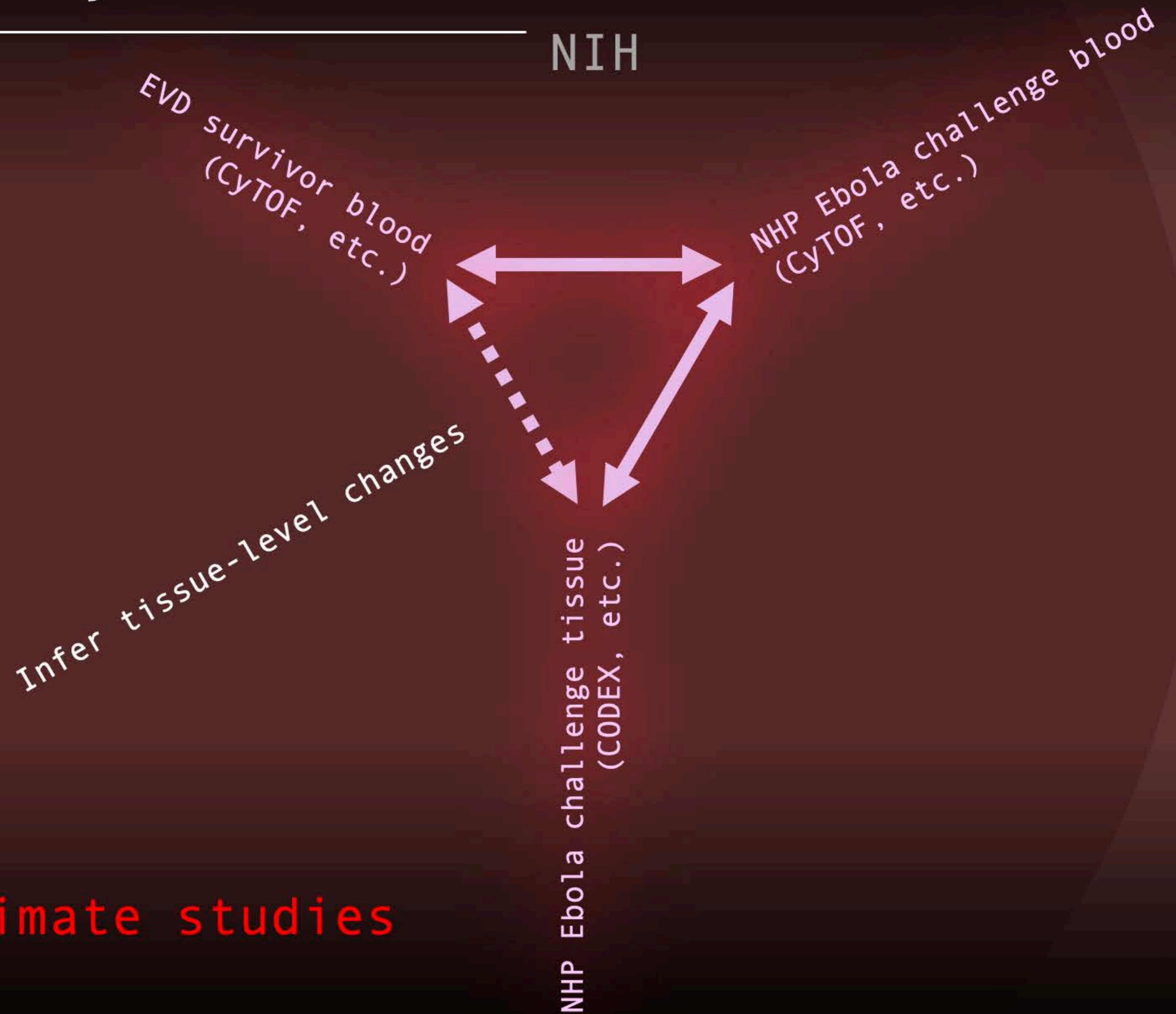
STANFORD MEDICINE



Influenza

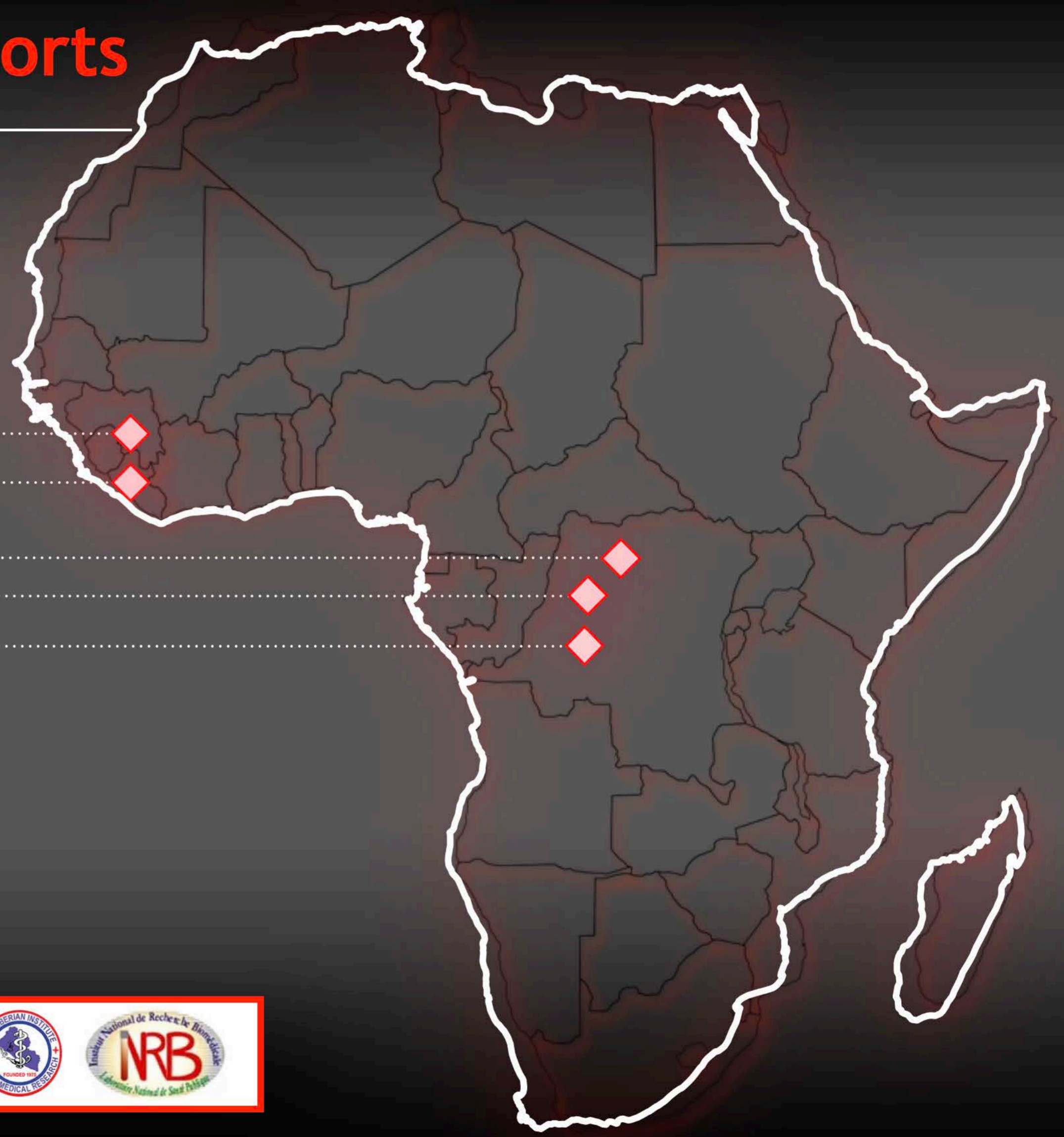
- 5-15% of the world's population affected annually
- 250,000-500,000 deaths/yr globally
- Economic costs \$100 billion/yr for U.S. alone
- Pandemic potential: 50 million deaths in 1918

STUDY • Cross Species CyTOF Panel



STUDY • EVD Survivor Cohorts

19
55



Guinea - 2014

Liberia - 2014

D.R. Congo (Yambuku) - 1976

D.R. Congo (Boende) - 2014

D.R. Congo (Kikwit) - 1995

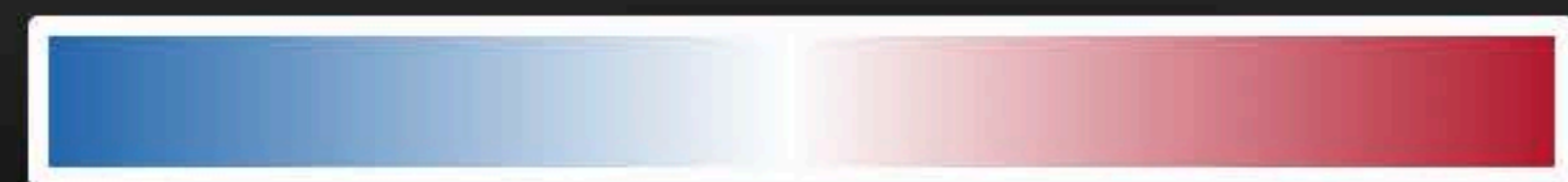
Collaborators:



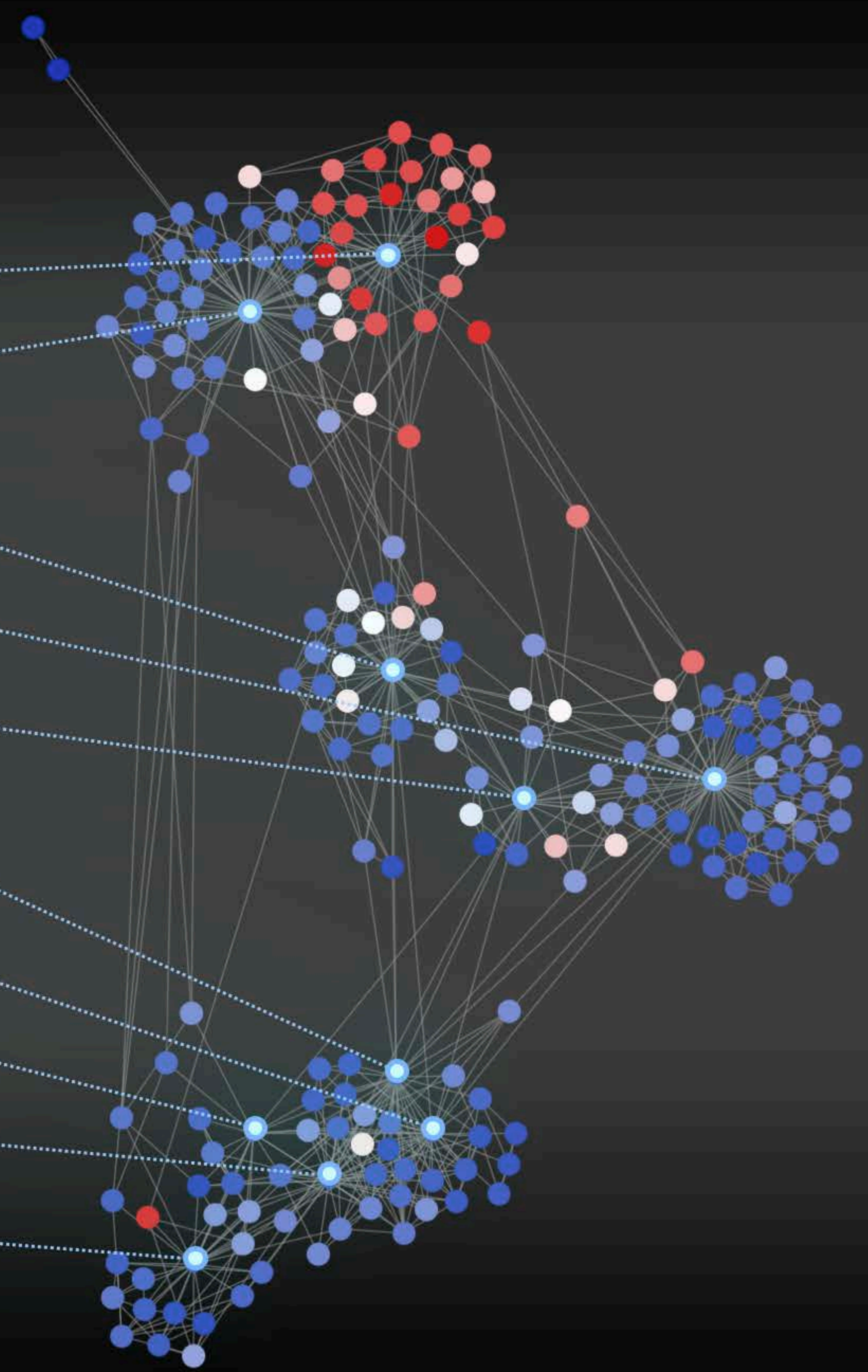
SCAFFOLD MAP • CD8

385 samples mapped
to 200 clusters

- CD8+ Tcells
- CD4+ Tcells
- NK cells
- Granulocytes
- Basophils
- ncMCs
- cMCs
- pDCs
- mDCs
- B cells

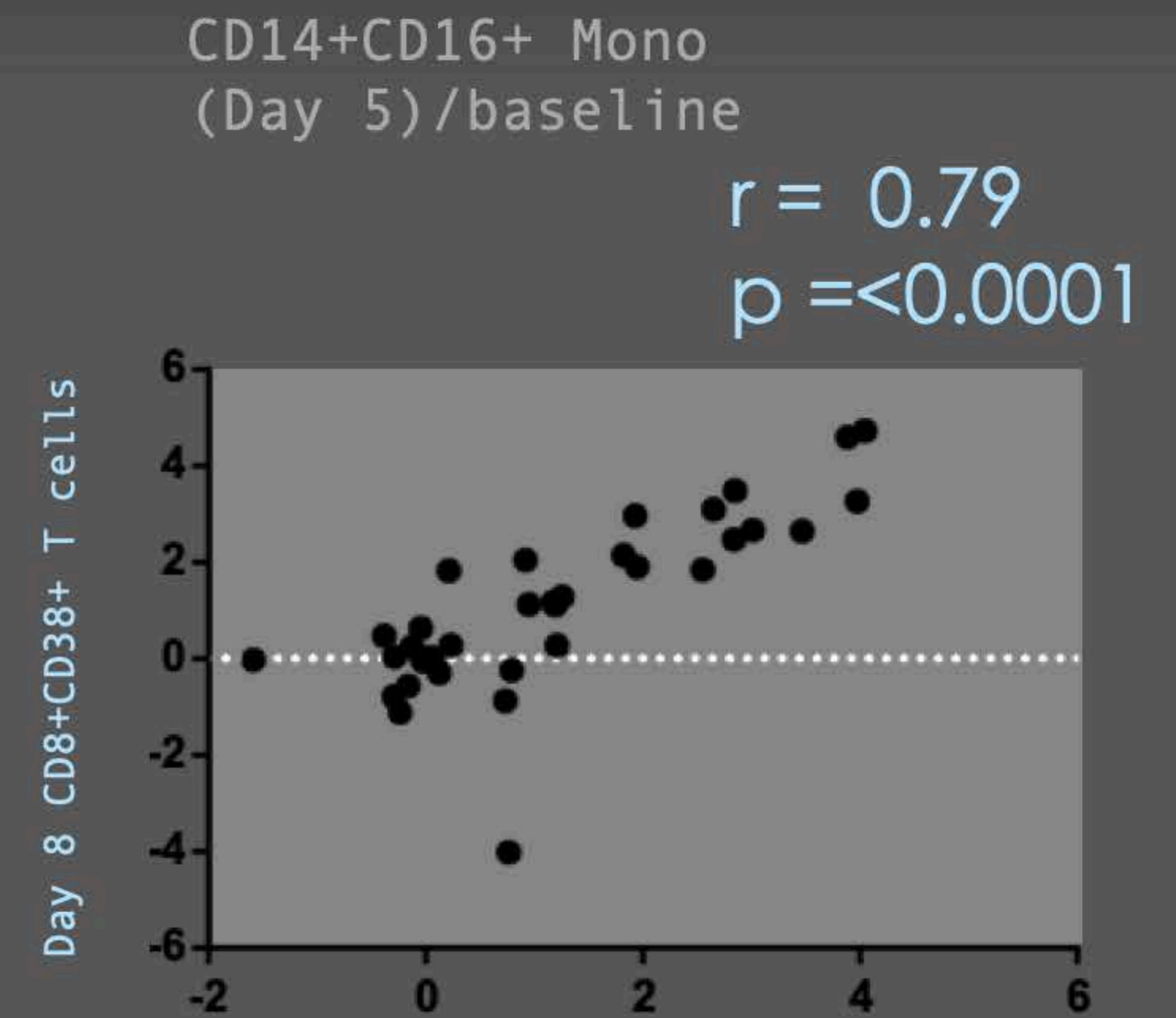
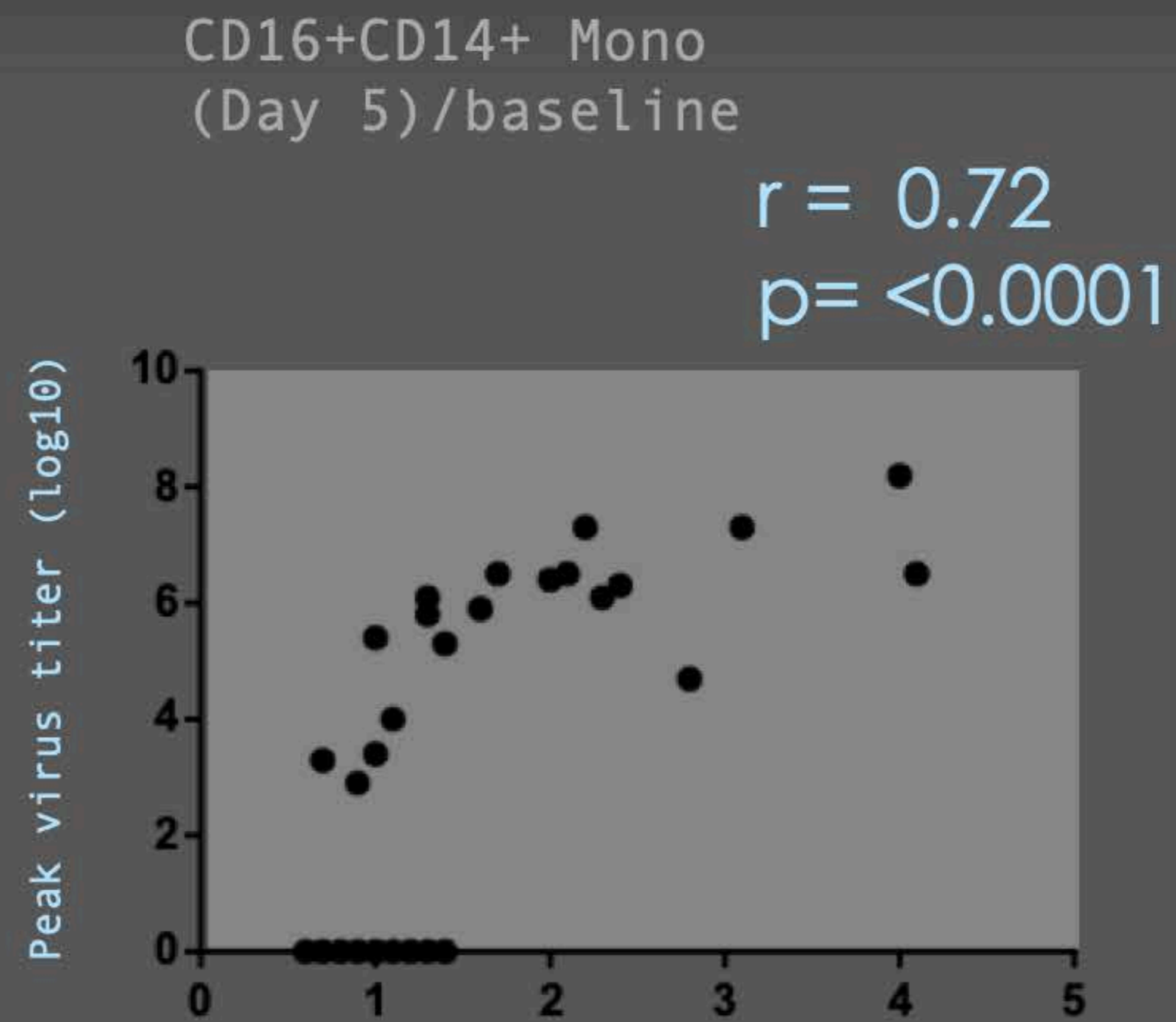
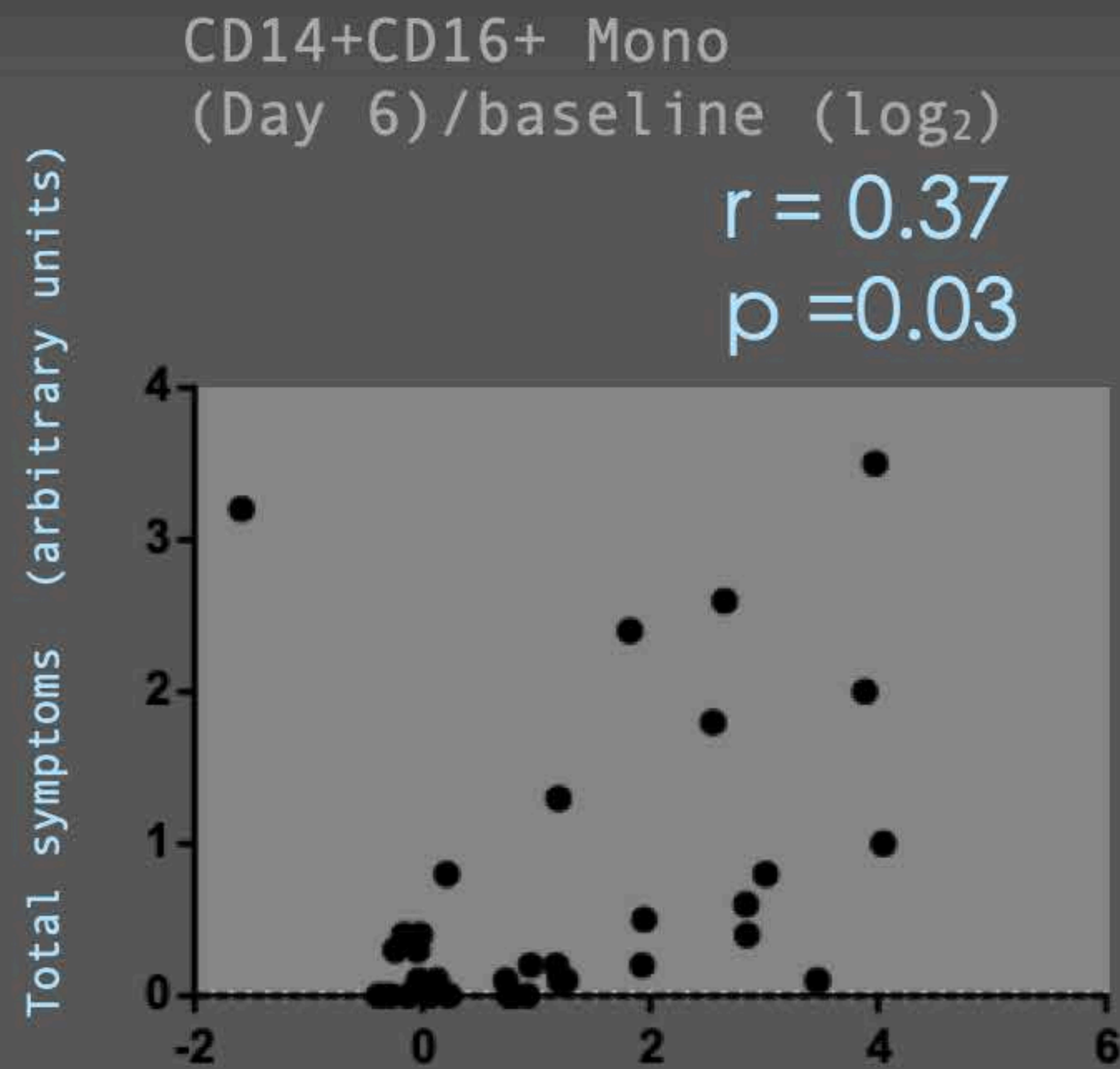


Fold vs. baseline
(Log₂)



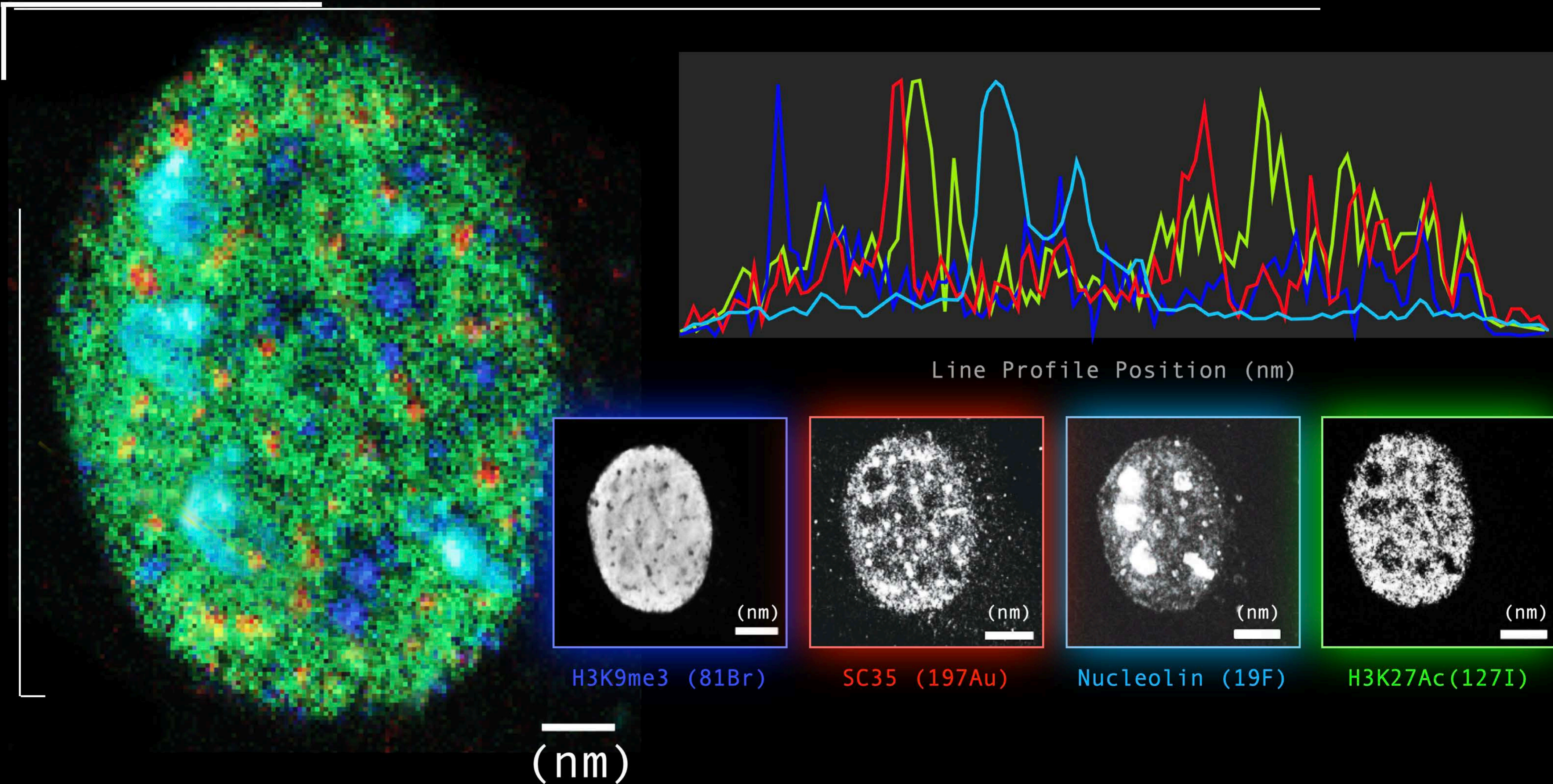
CORRELATIONS • H1N1

preliminary data



CD14+CD16+ Intermediate monocytes correlate with symptoms, virus titer, and T cell activation

3D MIBI • Determining Molecular Distribution with Precision



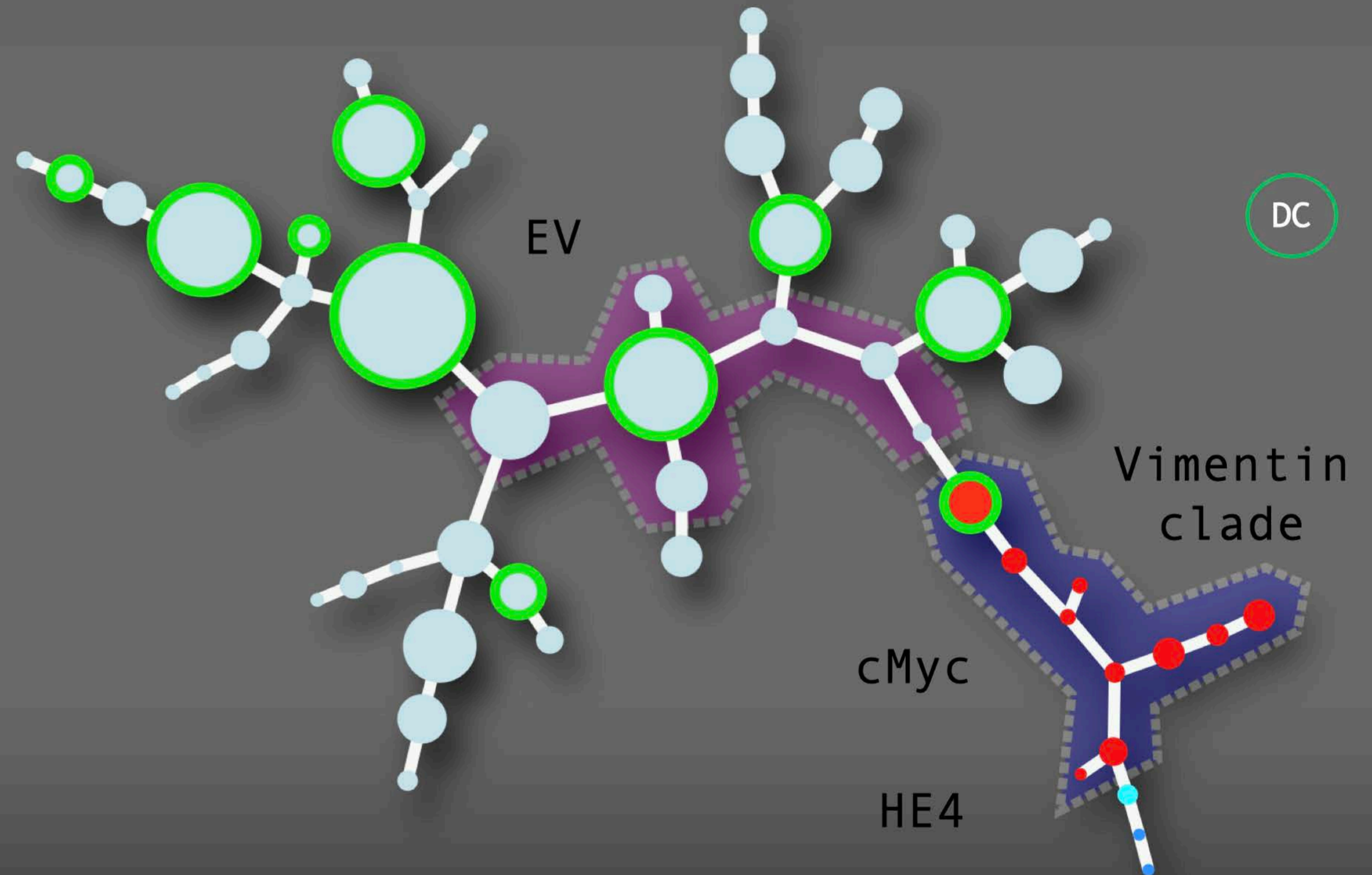
THERAPEUTIC OPPORTUNITIES •

Three new tumor cell types recur across samples

- EV co-expressing cells
- Dominant clusters
- V/cMyc/HE4 cell subsets

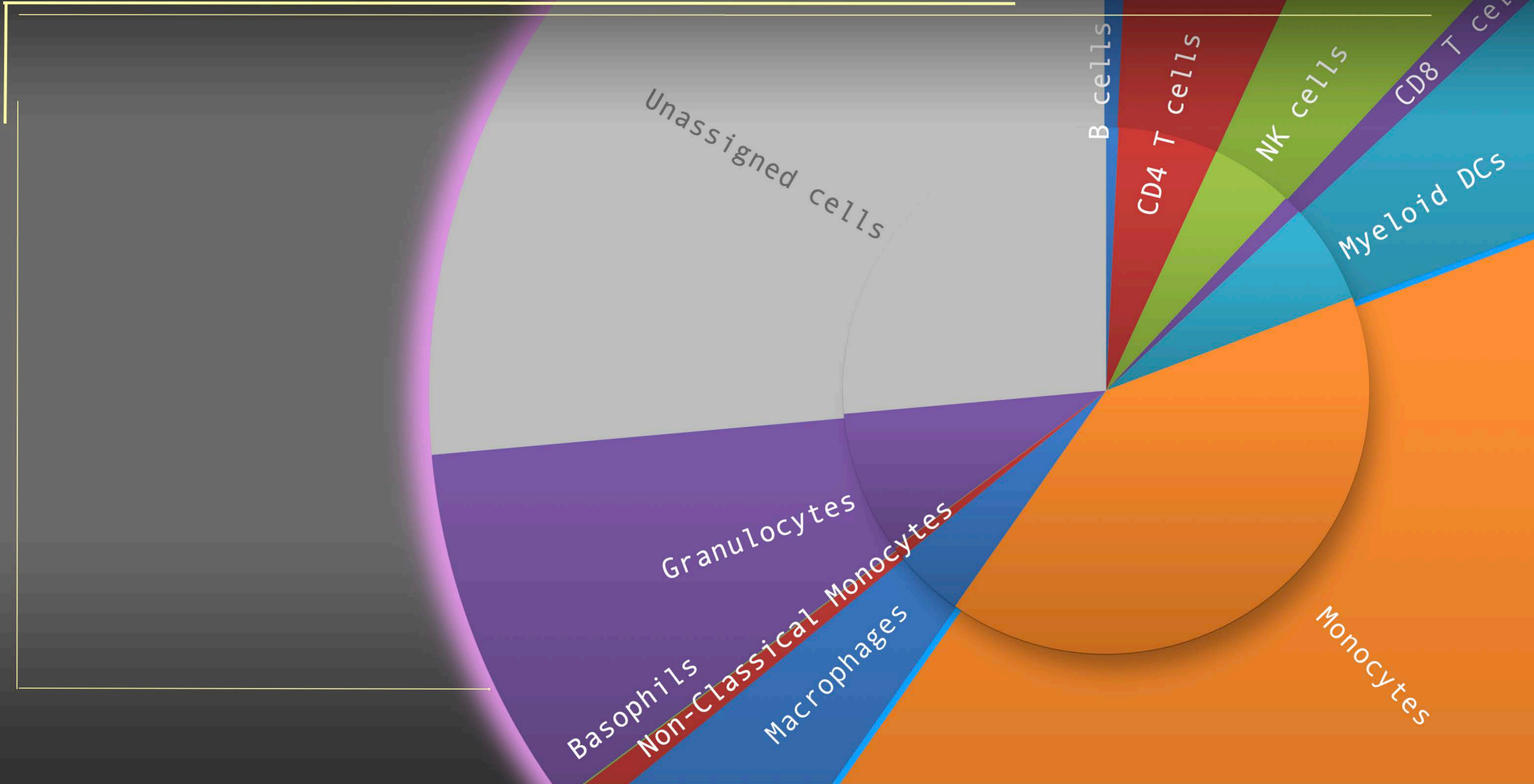
Clinical relevance

- New targets
- Diagnostic test
- Time to relapse
- Extent of surgical debulking



IMMUNE CELL FREQUENCIES

• Manual Gating



Garry P. Nolan
PhD



Christopher Green
MD, PhD

Rachford & Carlota Harris Professor
Dept. of Microbiology & Immunology
Stanford School of Medicine
Baxter Lab for Stem Cell Biology

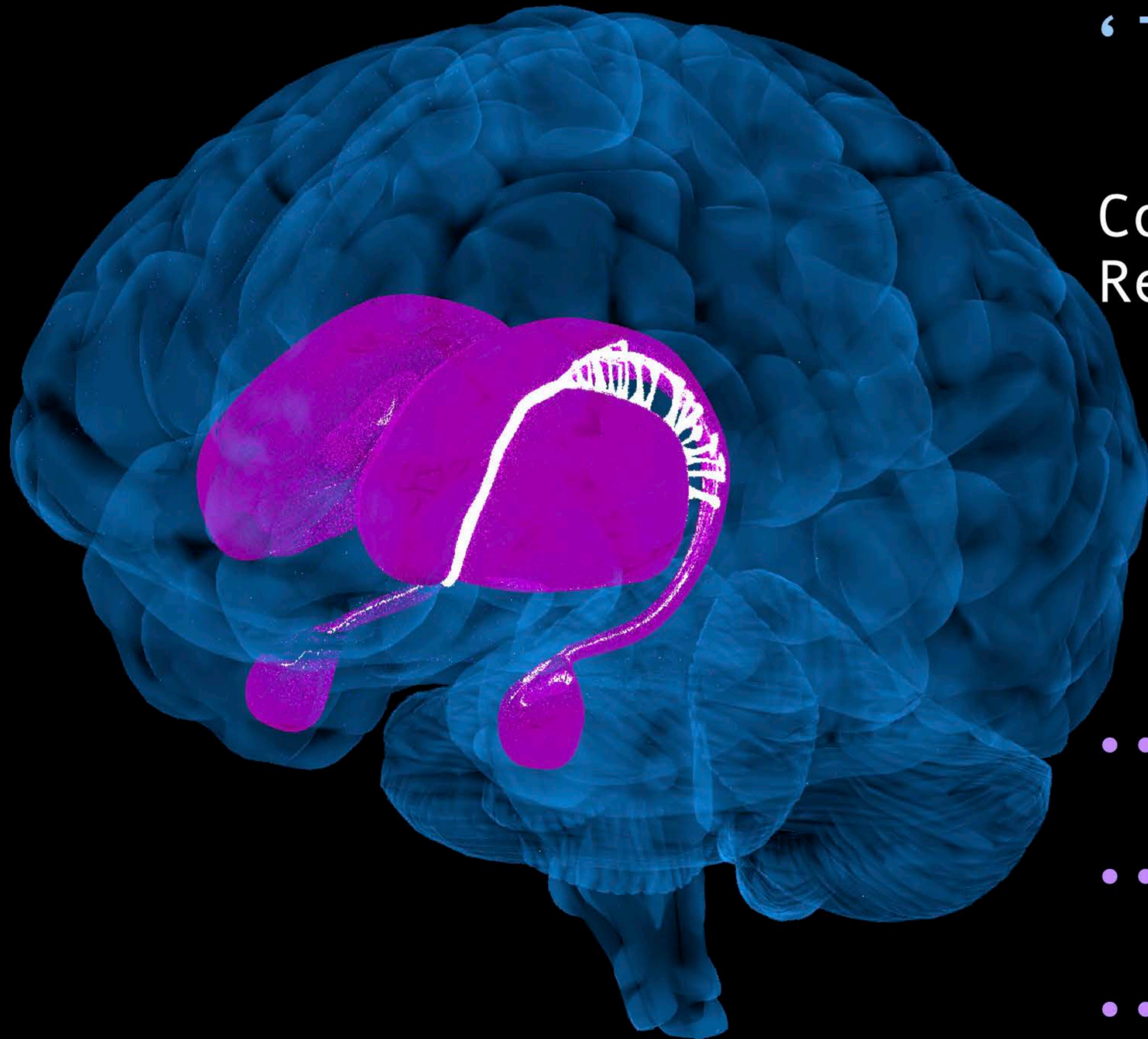
Forensic Neurology
Dept. of Diagnostic Radiology
Detroit Med. Cntr &
Wayne State School of Med.
Med:For, Inc. Chairman

CAUDATE \leftrightarrow PUTAMEN • An Introduction

Together -

'THE DORSAL STRIATUM

Cognitive, Executive, Decision-Making
Relay-Center for High-Functioning



- ... 'The Oldest'
- ... 'The Most Evolutionarily Developed'
- ... 'The Most Advanced'

HYPOTHESIS • Caudate Overdevelopment

CASE CRITERIA

Pick first 8 [of 24] in random series of IntFxSndx Diagnosed Experiencers

Common Syndrome Sub-Set Features

Orbs

High Strangeness events

Fear

Family witnessed / followed

FUO

Multiple witnesses

NOT Required (but may have)

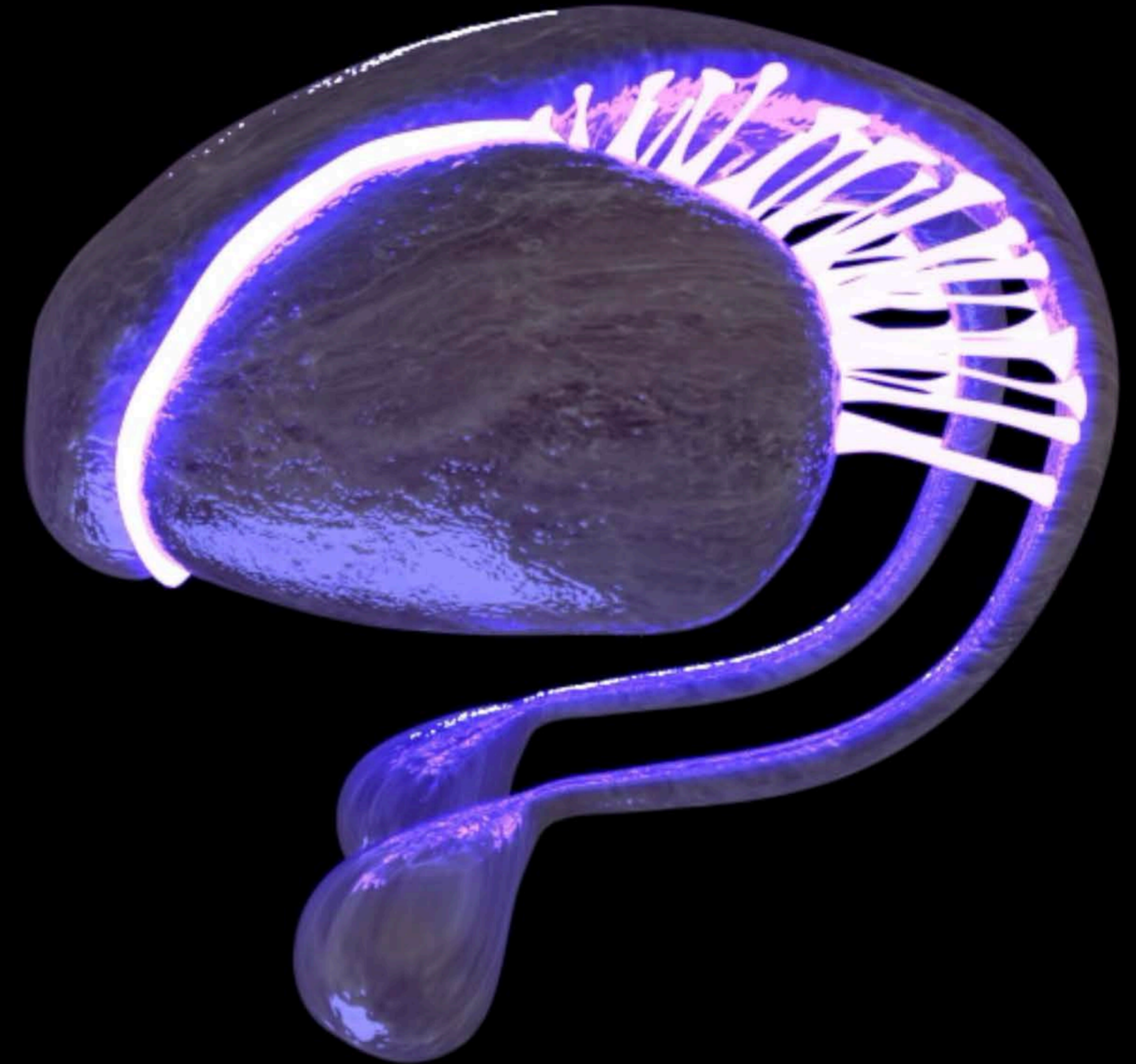
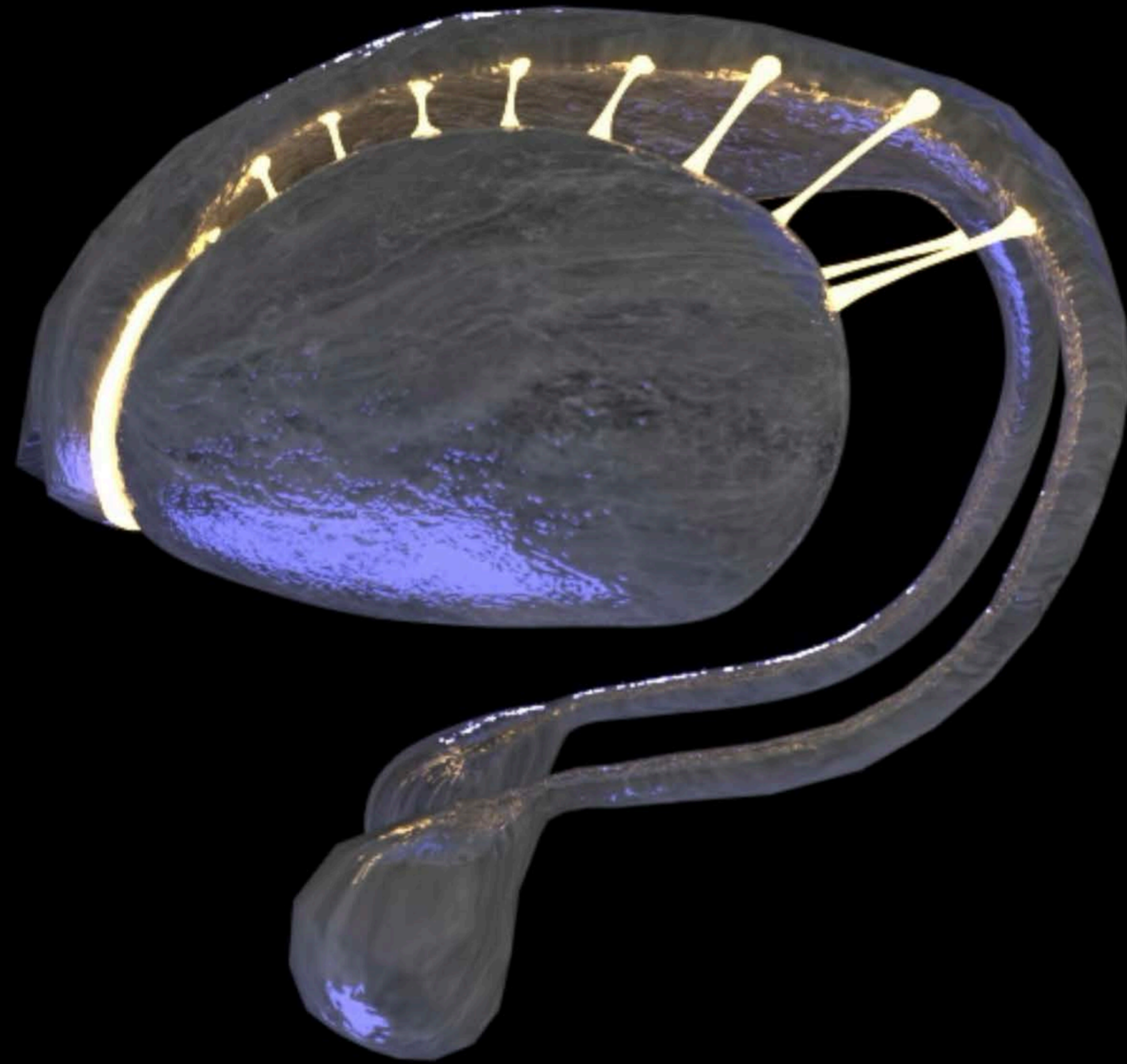
Marks

Endocrinopathy

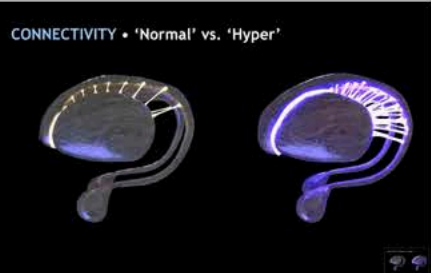
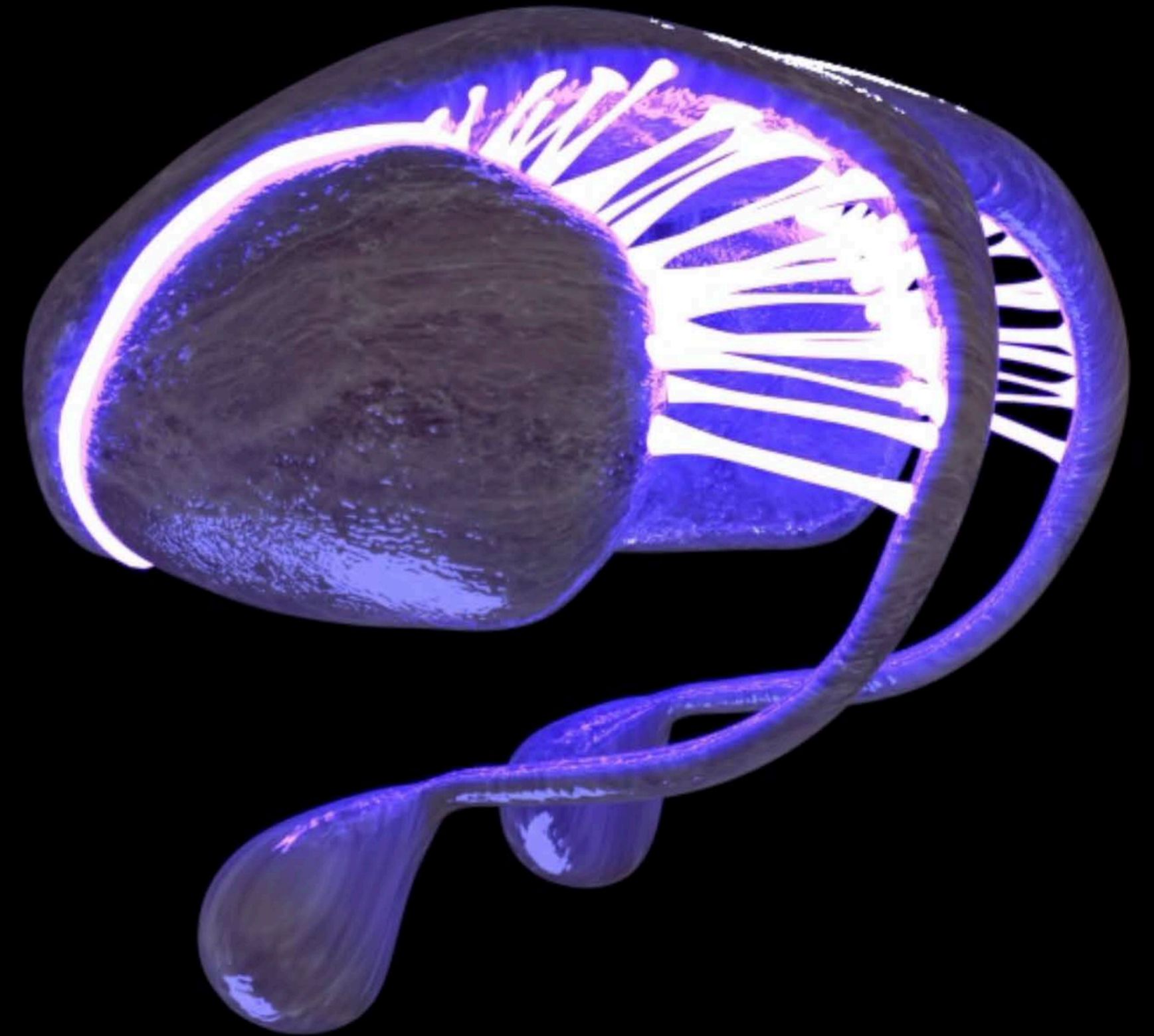
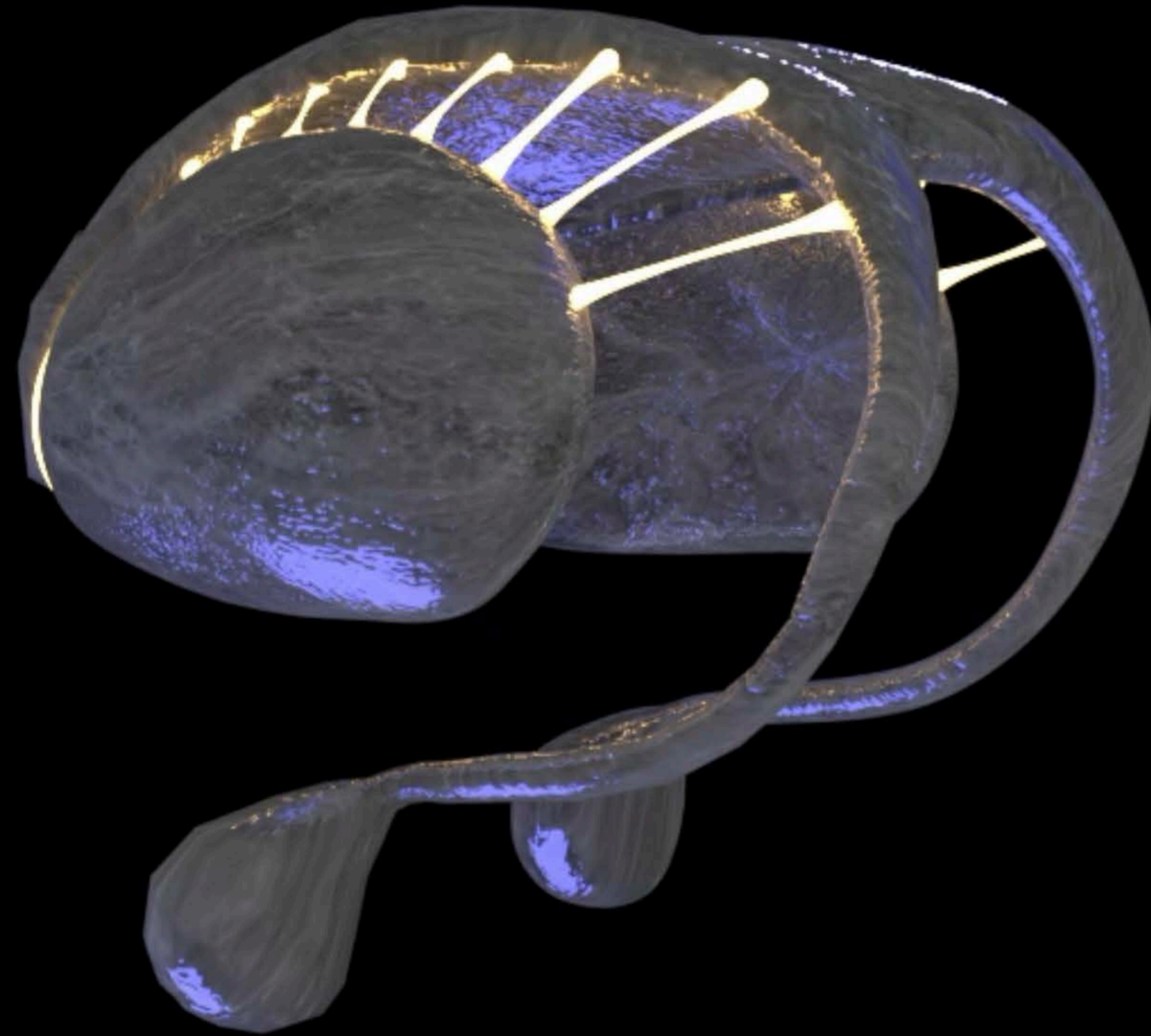
Neurological findings w/ MRI positives



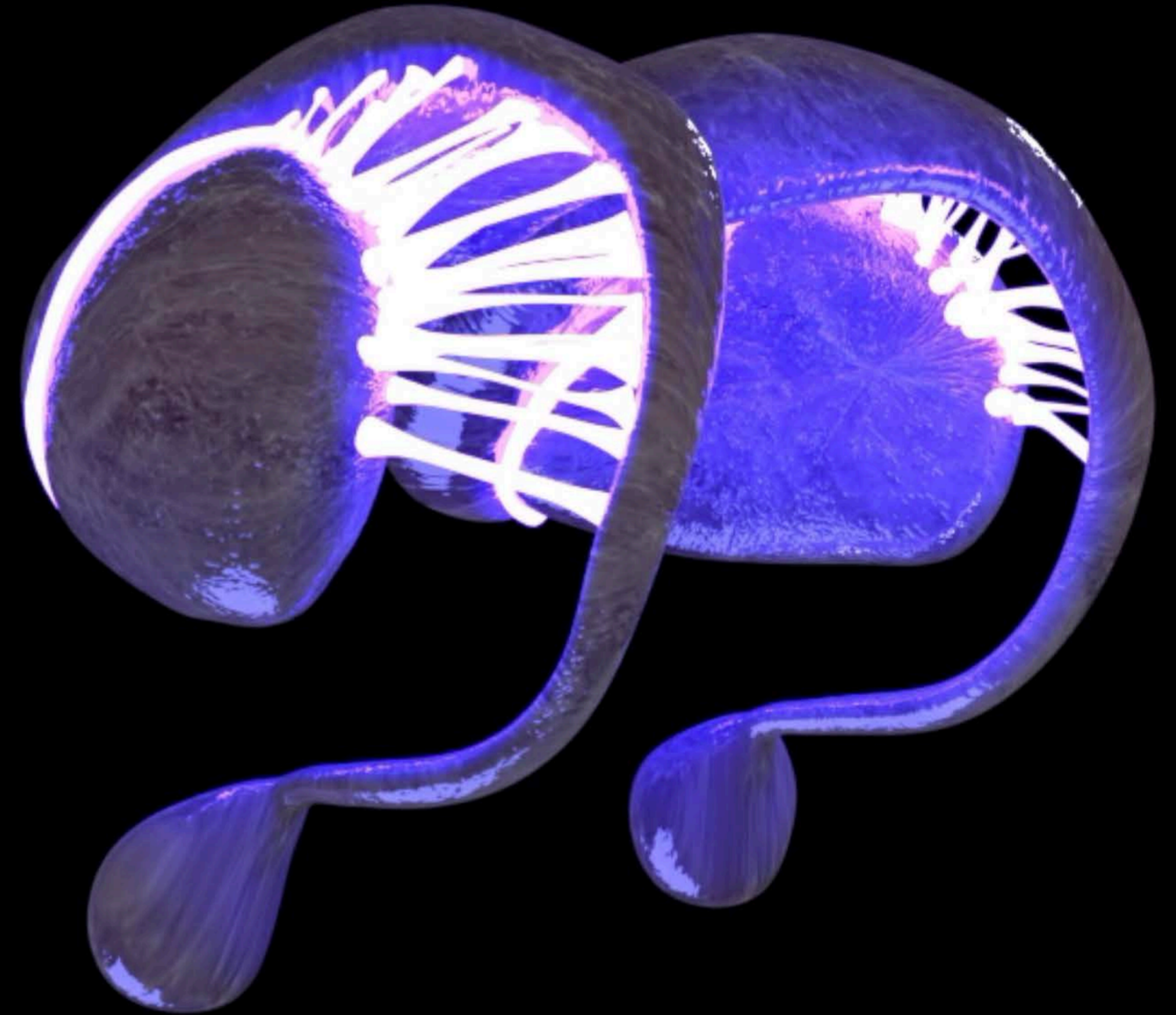
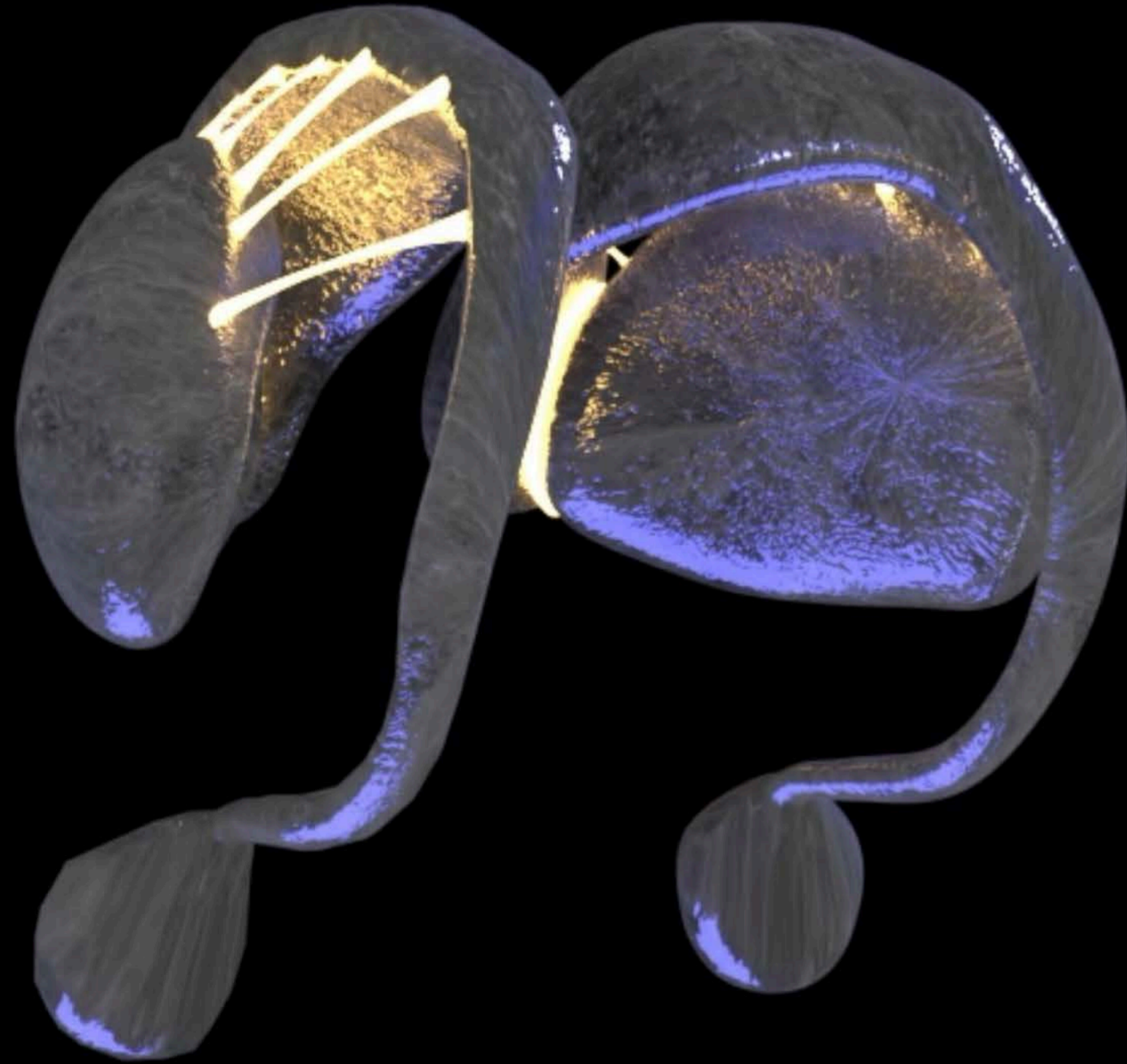
CONNECTIVITY • 'Normal' vs. 'Hyper'



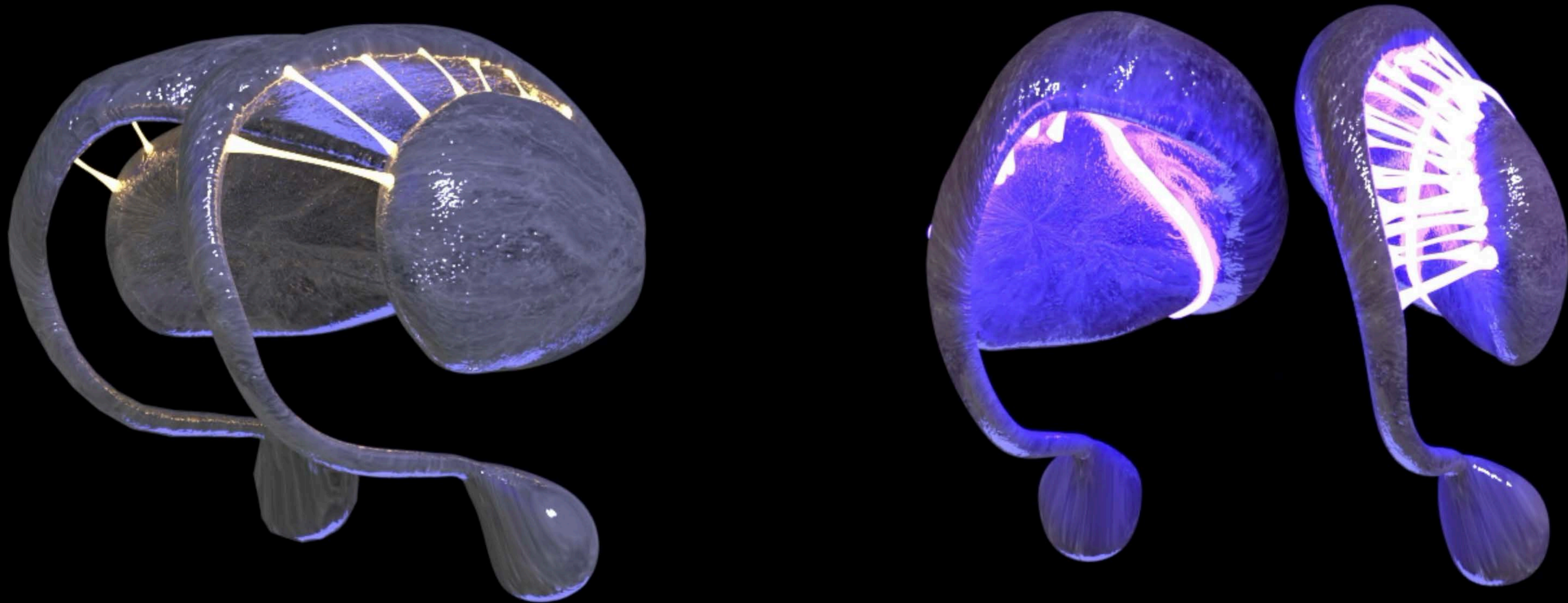
CONNECTIVITY • 'Normal' vs. 'Hyper'



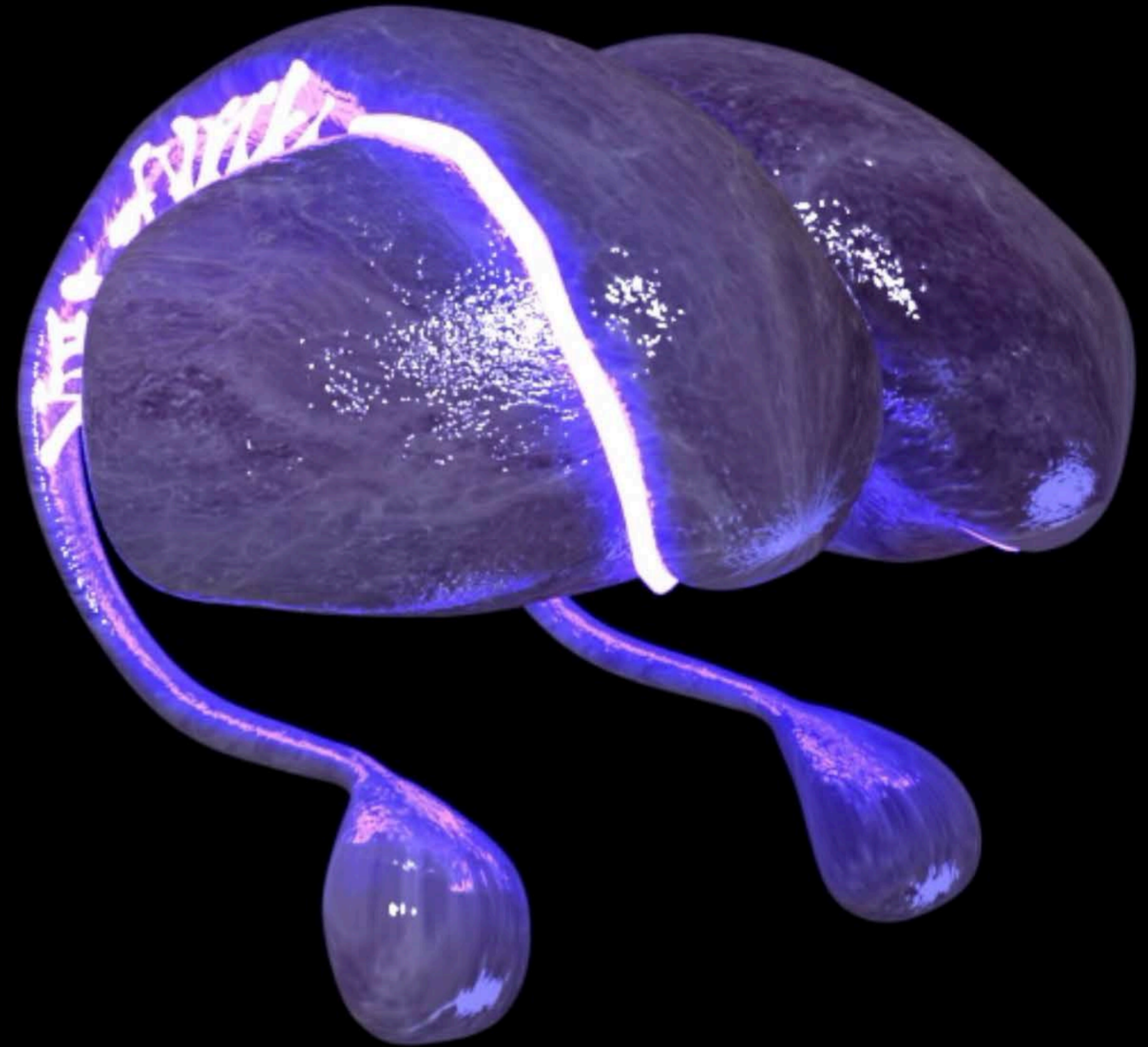
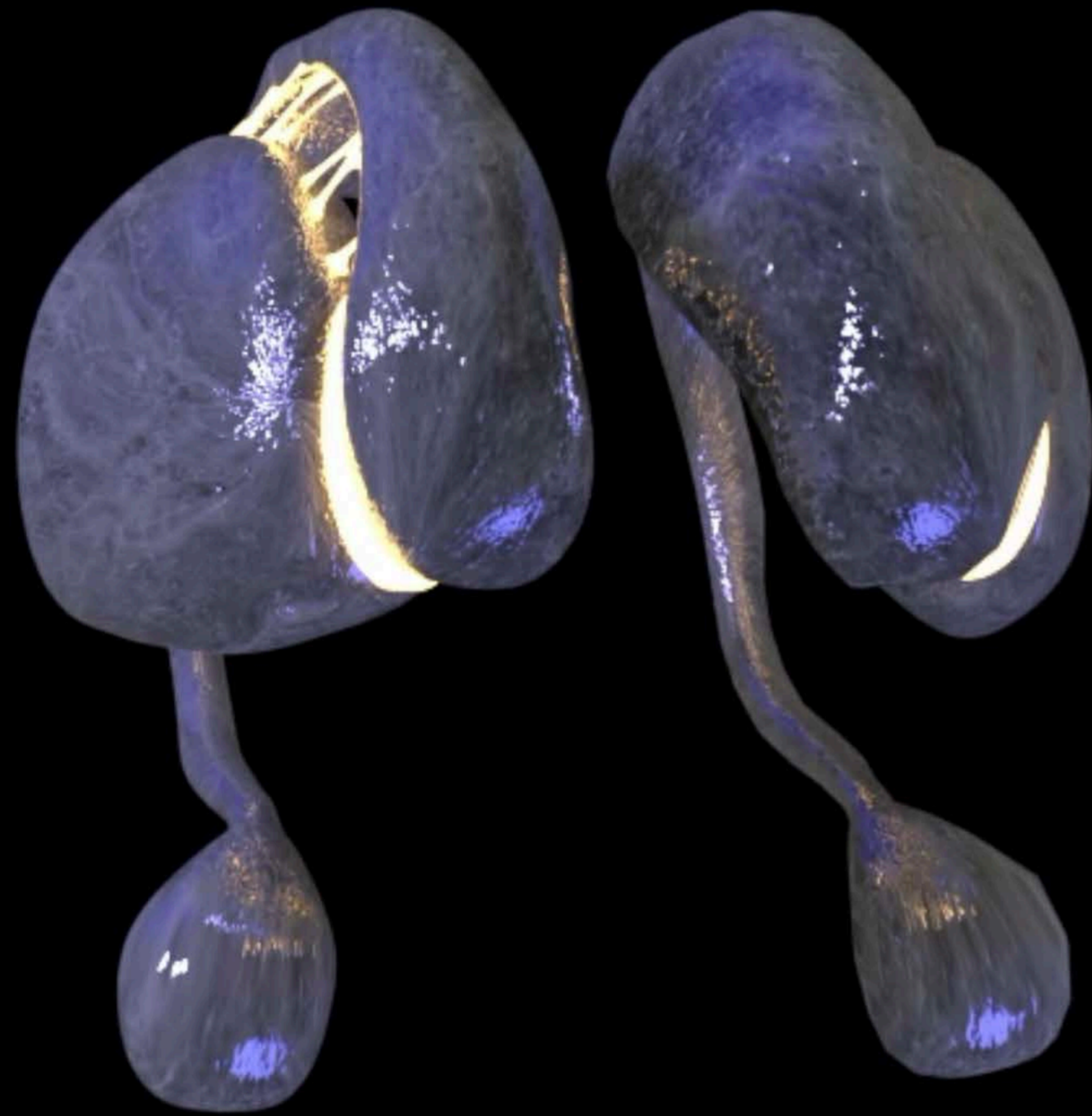
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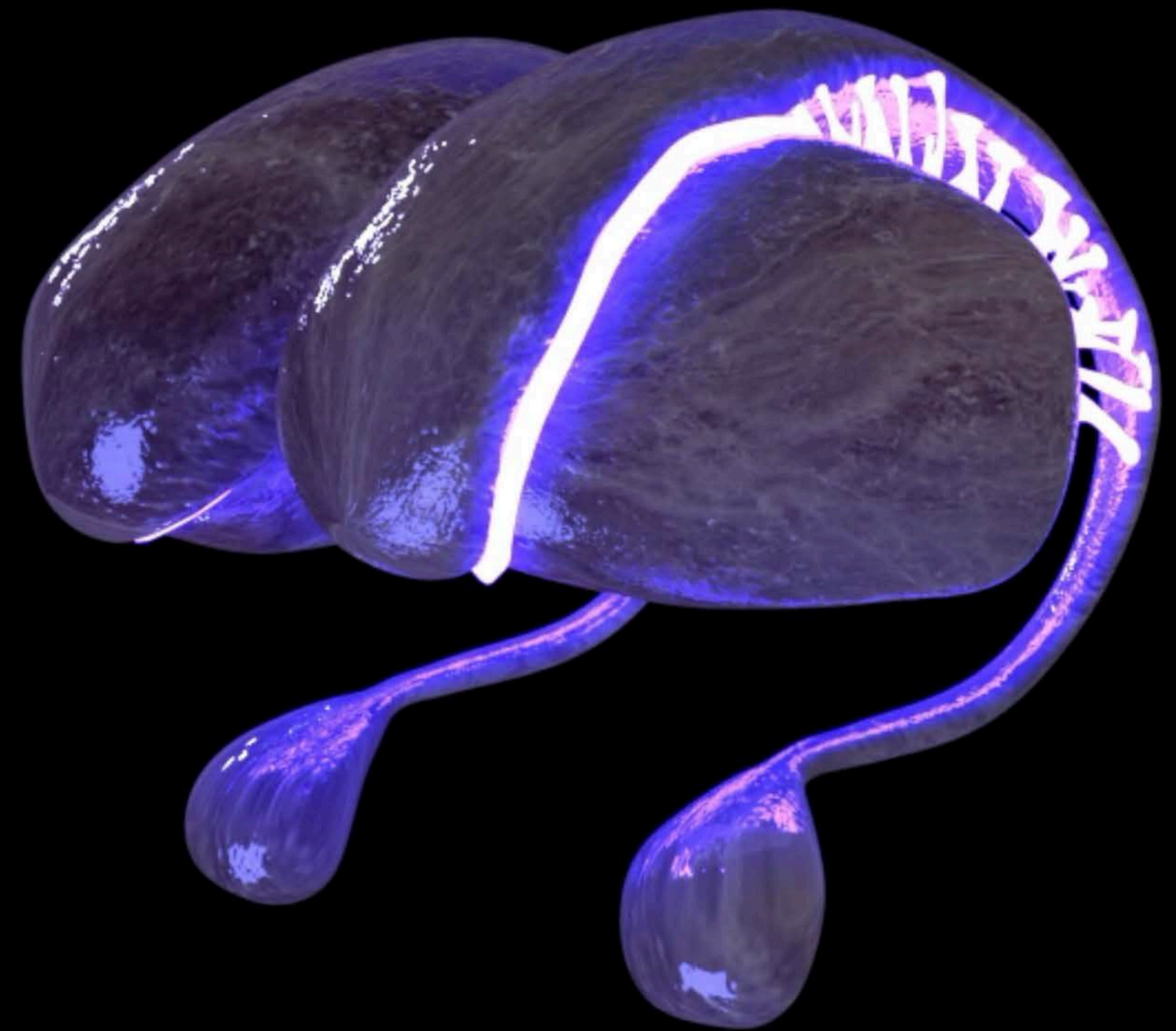
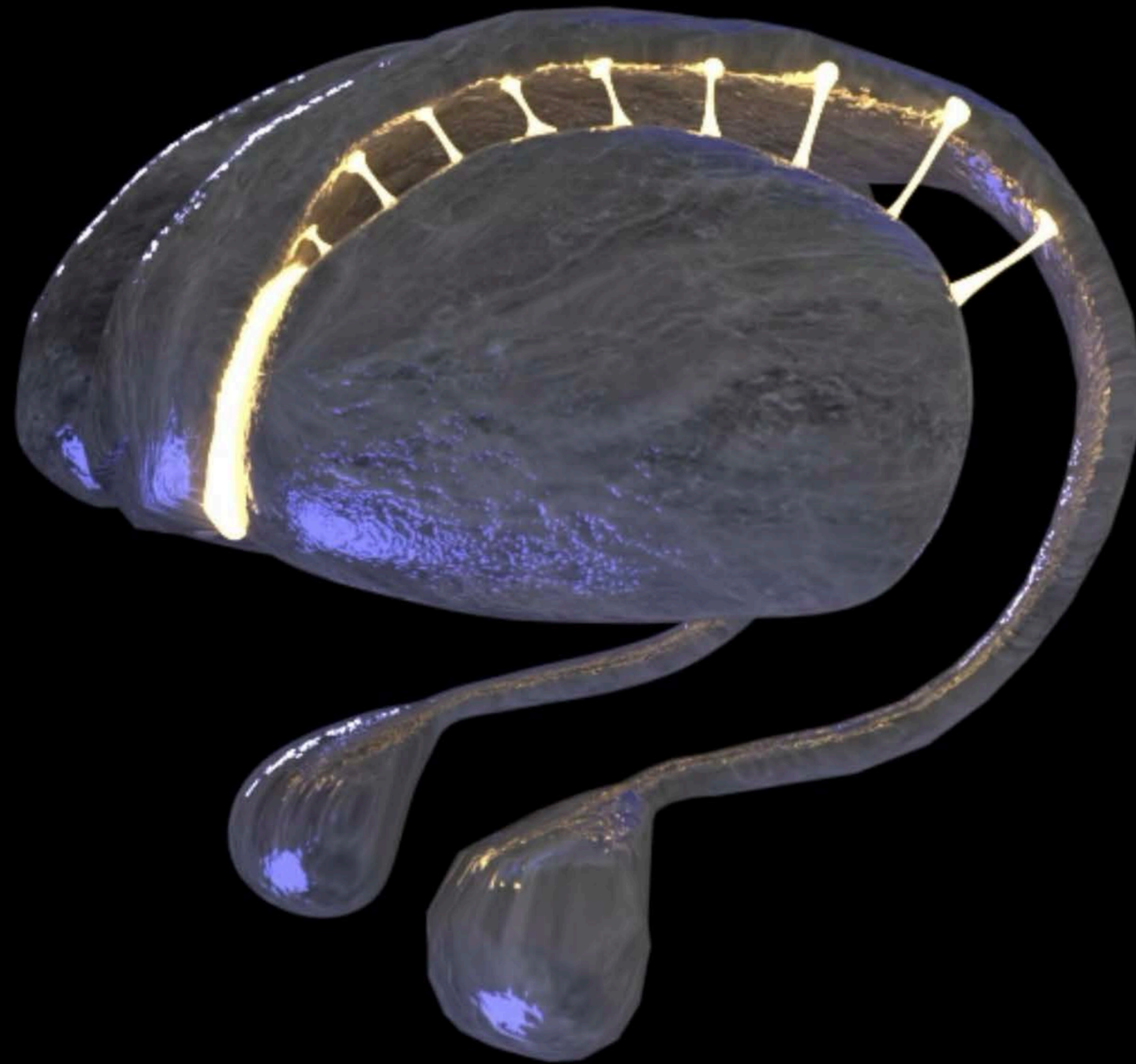
CONNECTIVITY • 'Normal' vs. 'Hyper'



CONNECTIVITY • 'Normal' vs. 'Hyper'



CONNECTIVITY • 'Normal' vs. 'Hyper'



ÅTOMNAUT

A.M.P.

Operating Principles

AMP combines 3 microscope technologies:

FIM field-ion microscopy

- structure: surface imaging using gas ions

APT atom probe tomography

- chemistry/color: element identification via TOF-MS

FIB focussed ion beam

- in-situ sample prep: mills needle-shaped specimens

Patented laser-decontamination silences interference produced by the technology combination; e.g. FIM requires an imaging gas, APT requires gas-free, a difference in vacuum pressure requirements of 1,000,000x.

AMP solves the two biggest problems with predecessor techniques:

1. Specimen preparation: 50x faster. Months into days, makes AMP 'routine'.
2. Complete atomic resolution: 100x improvement. Every atom, 3D position & identity.

Pharma

\$ 1,100,000,000,000

Trillion

Markets

Protein Crystallography

\$ 1,300,000,000

Billion

7% CAGR

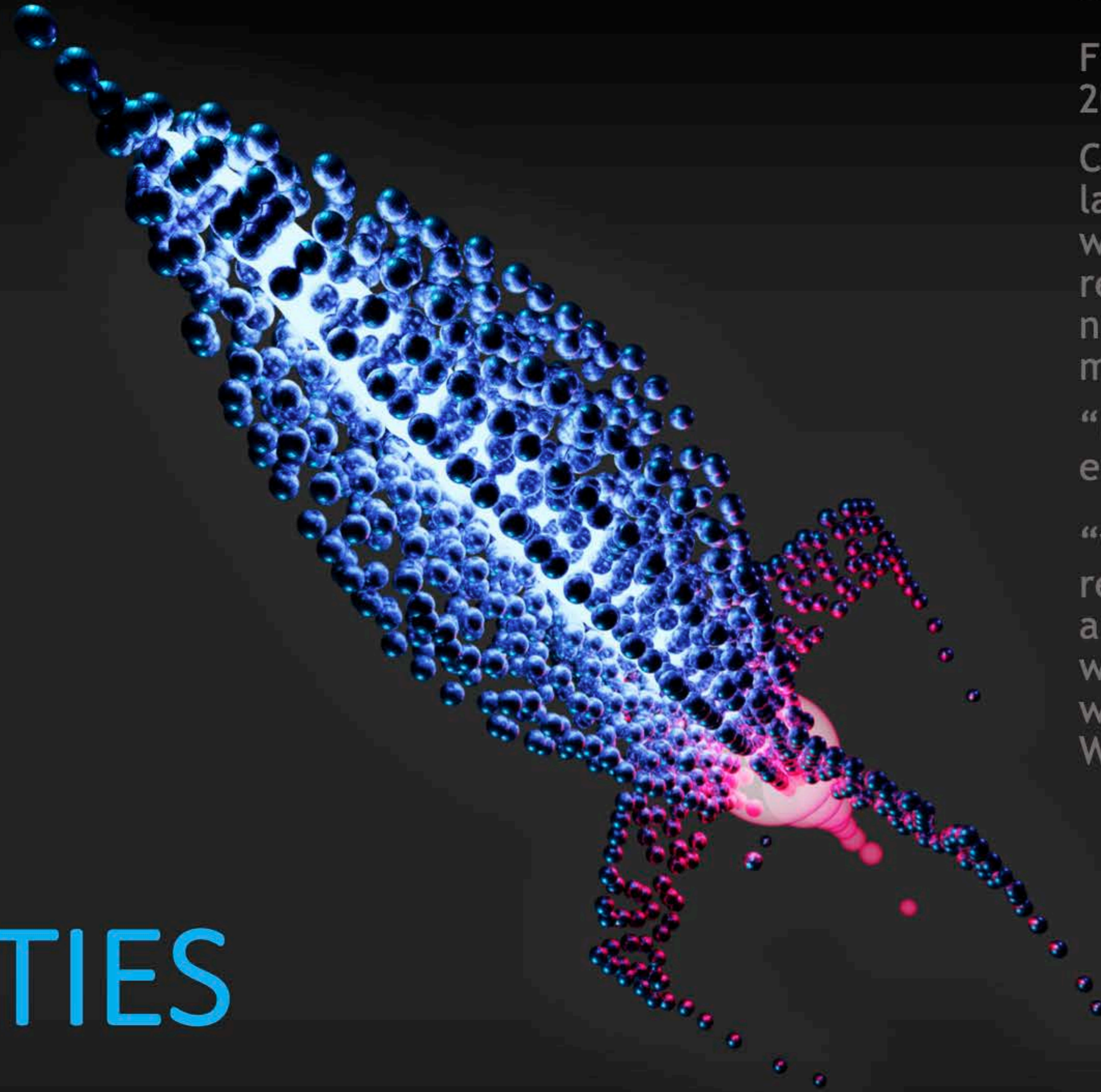
R&D

\$ 182,000,000,000

Billion

From DNA strands to extended chromatin, AMP can provide full field-of-view 3D atom maps.

Not as a competitor with high-speed DNA sequencing, but as a unique 3D spatial & chemical mapping tool of DNA, AMP will reveal previously hidden structure-function relationships.



Comparative technology

FISSEQ by ReadCoor
2016 Series A, \$25M

Compared to AMP, FISSEQ views a larger region of data (μm scale) but with poorer spatial and chemical resolution - AMP's opportunity is in nm-scale spatial sequencing measurements.

“Locating the products of genes is everything”

“will give neurologists, cancer researchers, clinicians, geneticists, and others the 3D coordinates of working genes that will inform their work in new ways.” (Terry, 2016, Wyss Institute)

OPPORTUNITIES

DNA / RNA

Atomic Scale Spatial Sequencing