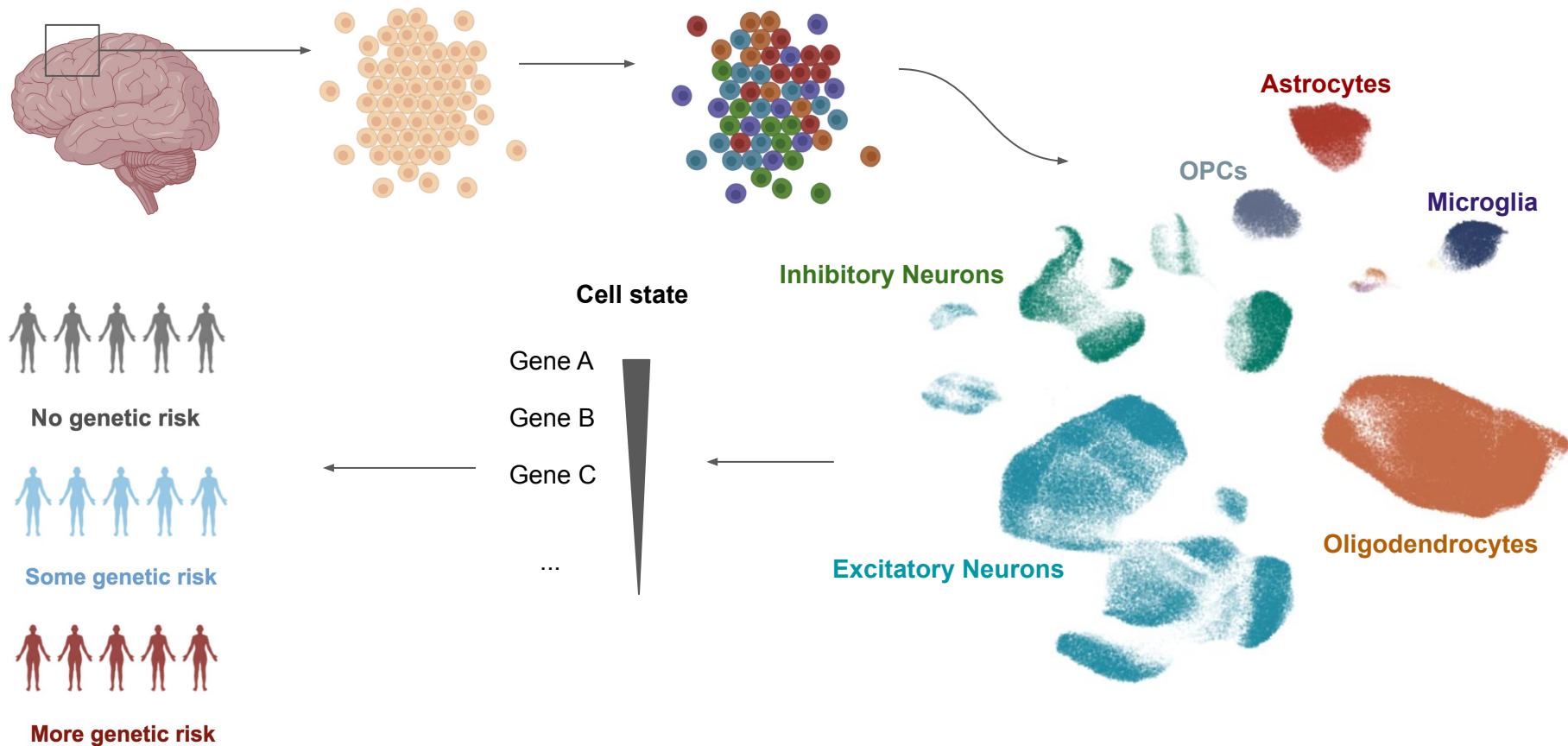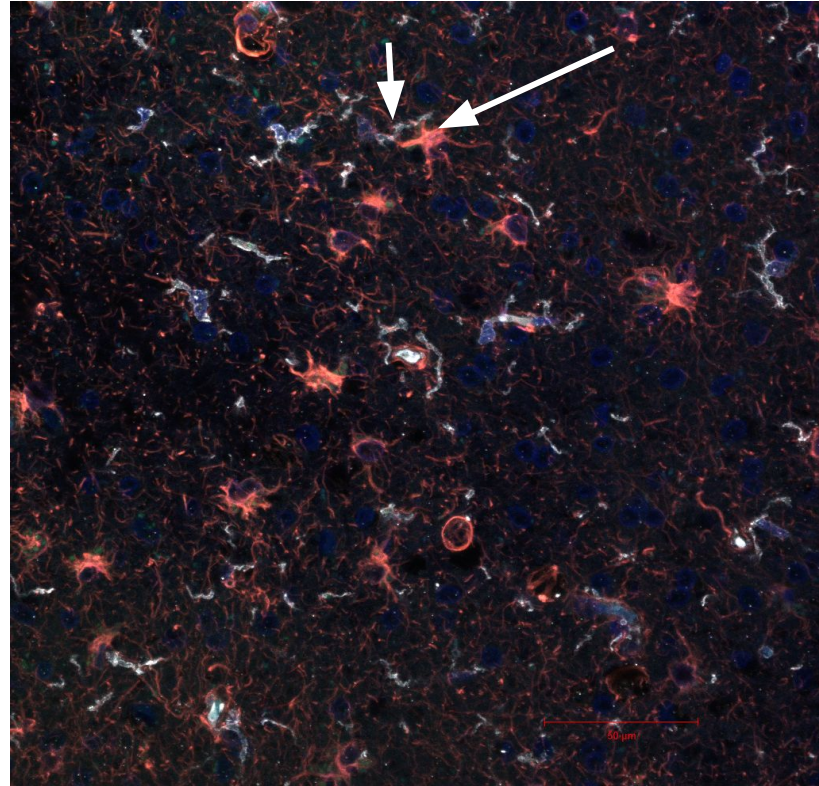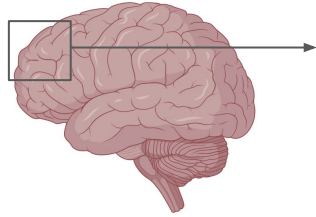# Measuring single-cell transcriptomic states in the human brain
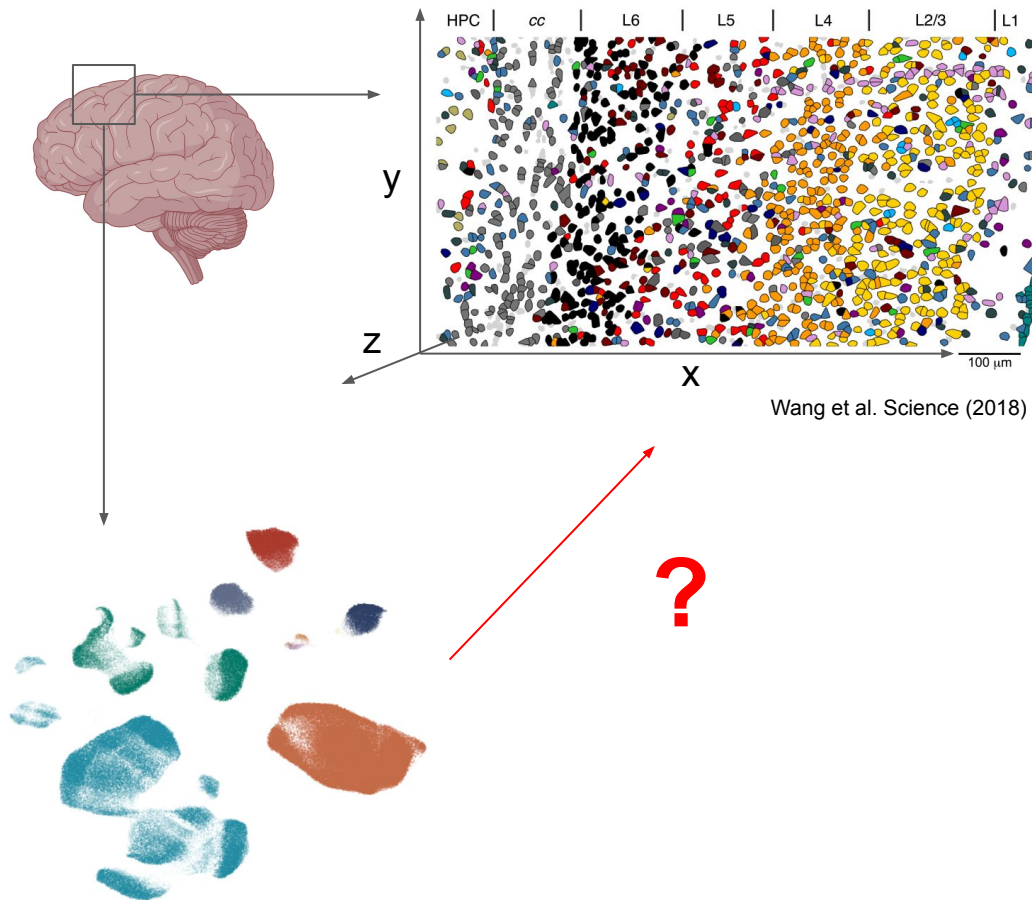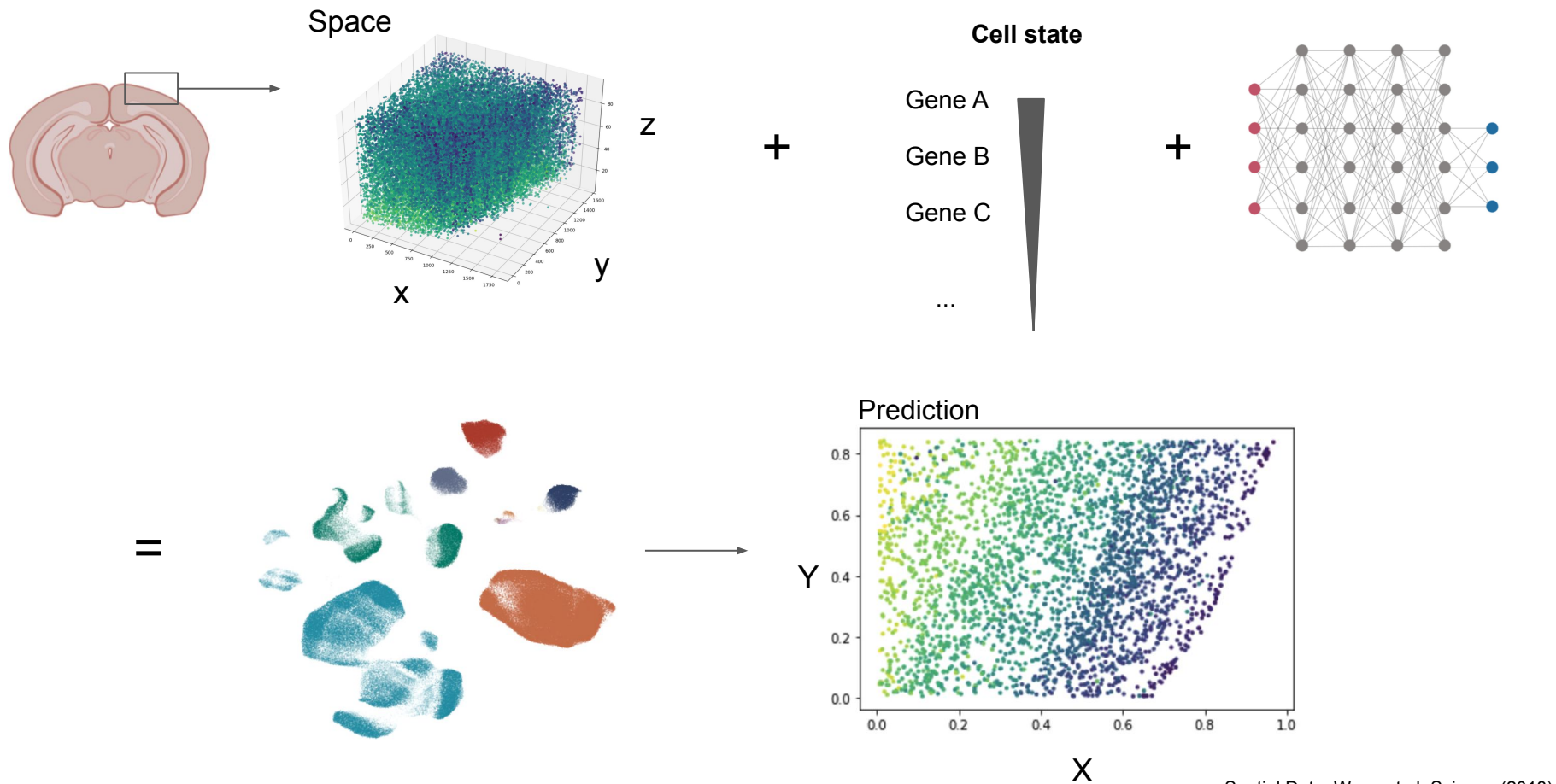
Djuna von Maydell
Tsai & Kellis labs

Human prefrontal cortex at single-cell resolution

Learning spatial relationships from single-cell transcriptomics

HPC | cc | L6 | L5 | L4 | L2/3 | L1

y

z

x

100 µm

Wang et al. Science (2018)

?

Learning spatial relationships from single-cell transcriptomics

Spatial Data: Wang et al. Science (2018)

We think this prediction will help us interpret cell states more accurately.
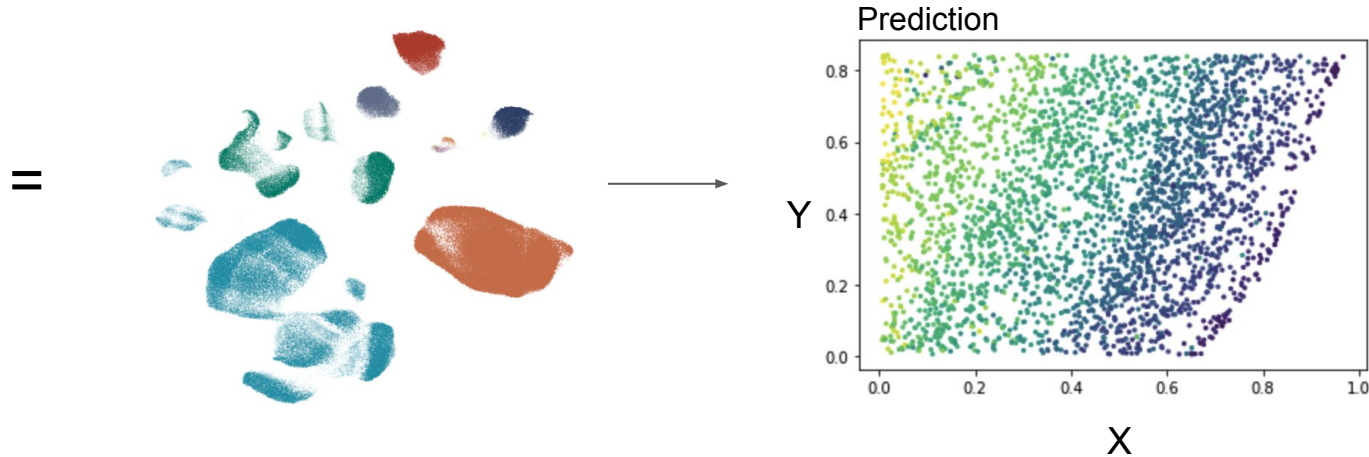
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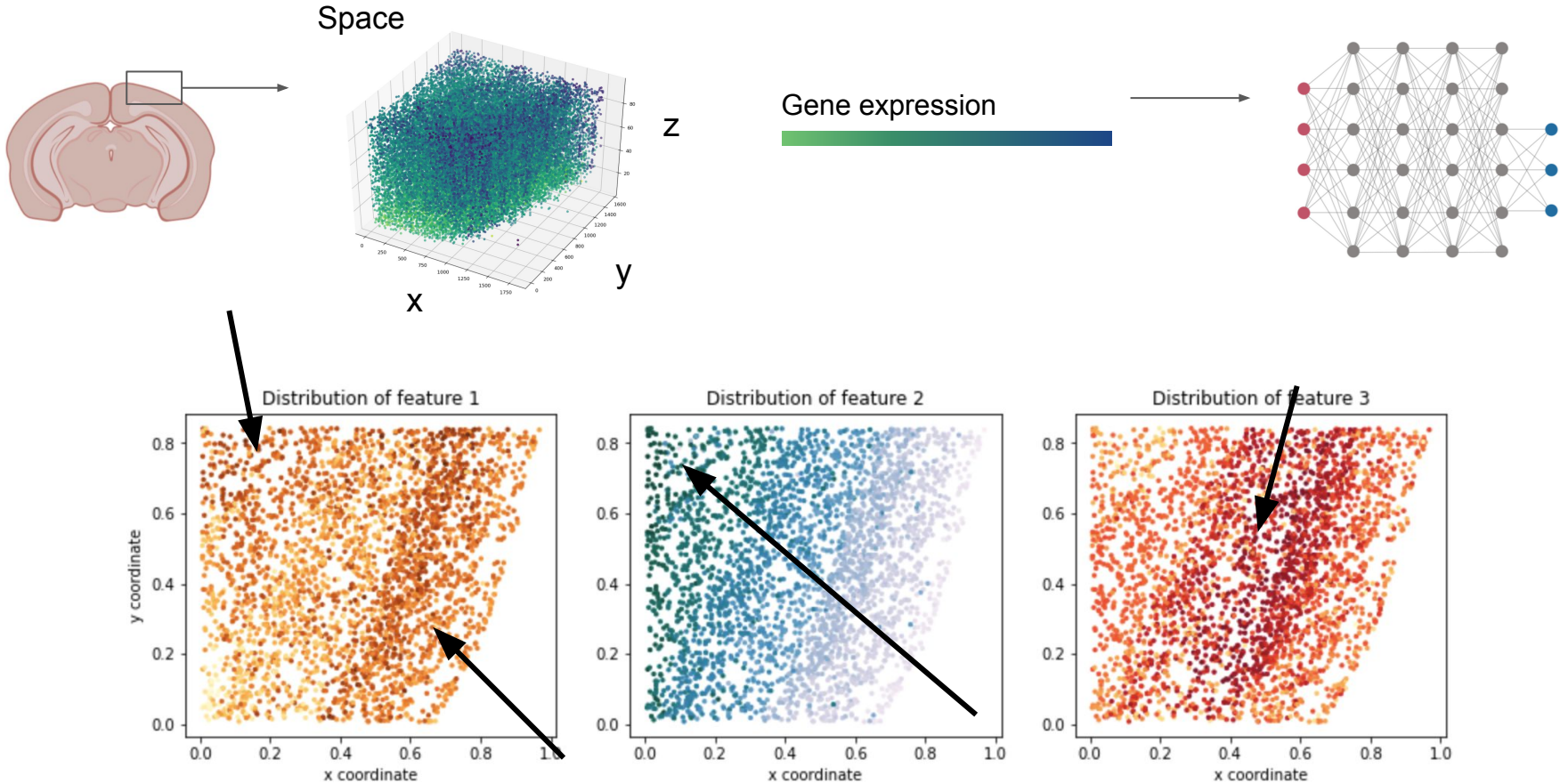
My Advisors:
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Everyone in the Tsai & Kellis labs!

Everyone in BCS!

Thank you for the Broshy Fellowship!

Learning spatial relationships from single-cell transcriptomics

Space

z

y

x

Gene expression

Distribution of feature 1

Distribution of feature 2

Distribution of feature 3

Effects of genetic risk variants on single-cell transcriptomic states