# The regulatory landscape of the developing fly brain

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The **Drosophila brain** is the workhorse in neuroscience.

Positioned at a **sweet spot** of modest brain size versus complex behavior, it has been subject to connectomics, transcriptomics and genetic studies.

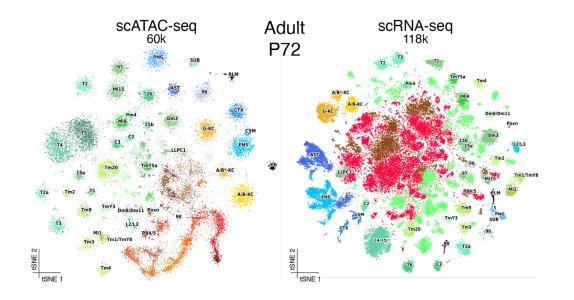
To fully understand the neuronal diversity we created an **epigenomic dataset of 240k cells**, and through integration with transcriptomics inferred **enhancer-GRNs**.

Finally, we study **neurogenesis and synaptog**enesis and reveal key regulators.

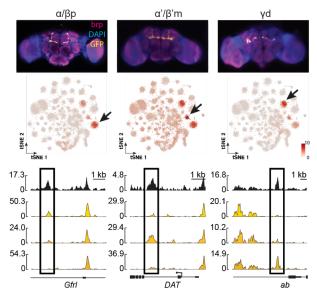
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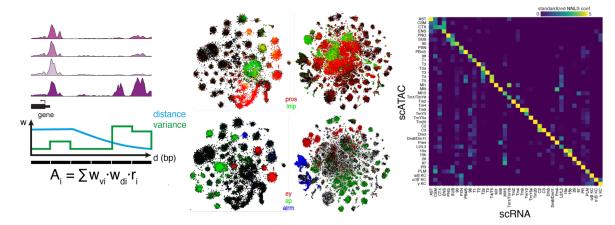
### Adult brain atlas

## Integration with RNA-seq

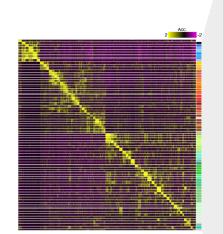


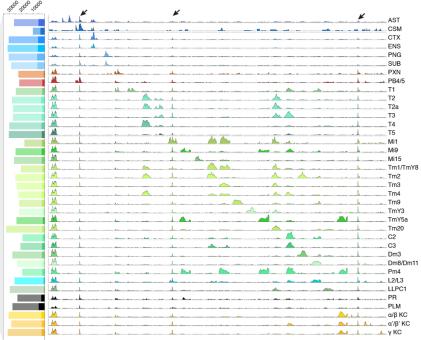
## Validation with FACS





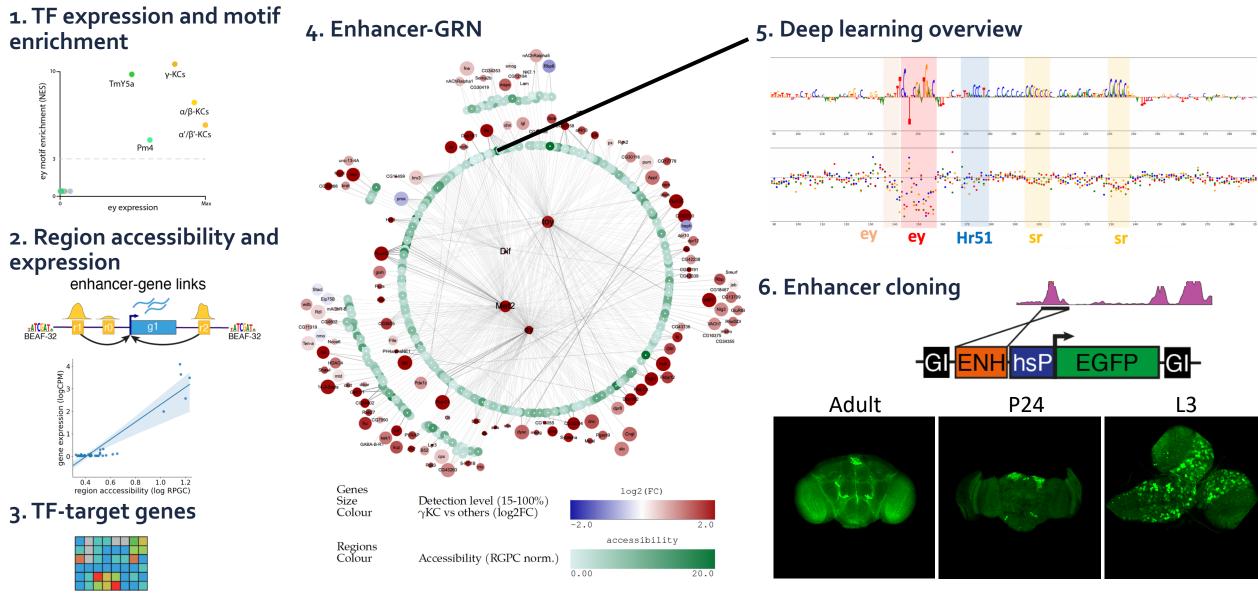
## Unique chromatin profiles





noe wun2 Optix wrapper indy Poxn Awh T/AP-2 acj6 bsh D Vmat Mef2 Pka-C3 vvl Vsx1 Lim3 ap kek3 nAchRr2 ort rgn ey GfrI DAT ab

## **Building enhancer-GRNs**



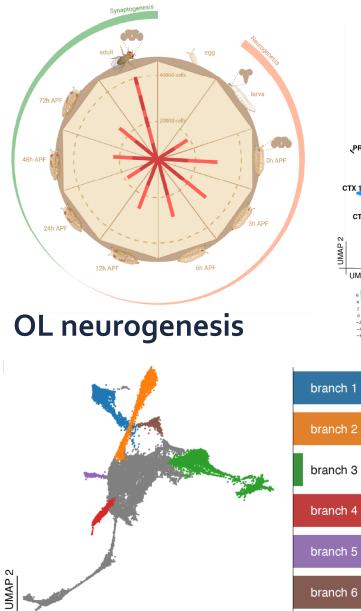
TF-gene coexpression

Region  $\rightarrow$  target gene links Width & transparency correlation score (GENIE3)

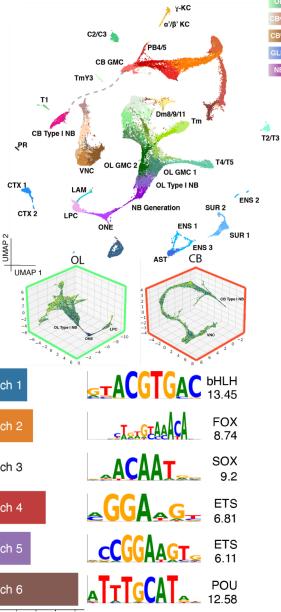
#### How it's made

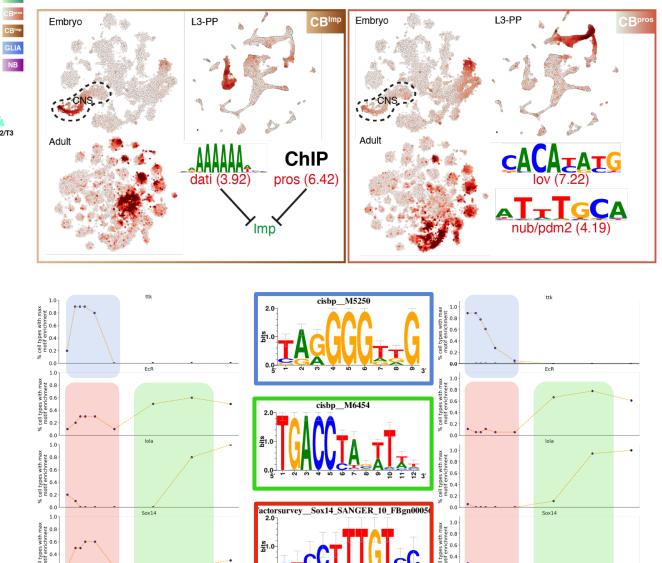
#### Larva-pupa atlas

## **CB** reprogramming



UMAP





40

0 100 200 300 400 500 600 nDARs