

BICCN – two review papers

NEWS & VIEWS FORUM | 06 October 2021

A census of cell types in the brain's motor cortex

An atlas of the cell types found in the motor cortex of the brain has been built using various types of data. Two neuroscientists explain the technological feats involved in the project, as well as the utility of the resource for future research.

Johan Winnubst  & Silvia Arber 

Article | Open Access | Published: 06 October 2021

A multimodal cell census and atlas of the mammalian primary motor cortex

[BRAIN Initiative Cell Census Network \(BICCN\)](#)

[Nature](#) 598, 86–102 (2021) | [Cite this article](#)

[Metrics](#)

BICCN – Transcriptomic and epigenomic atlas

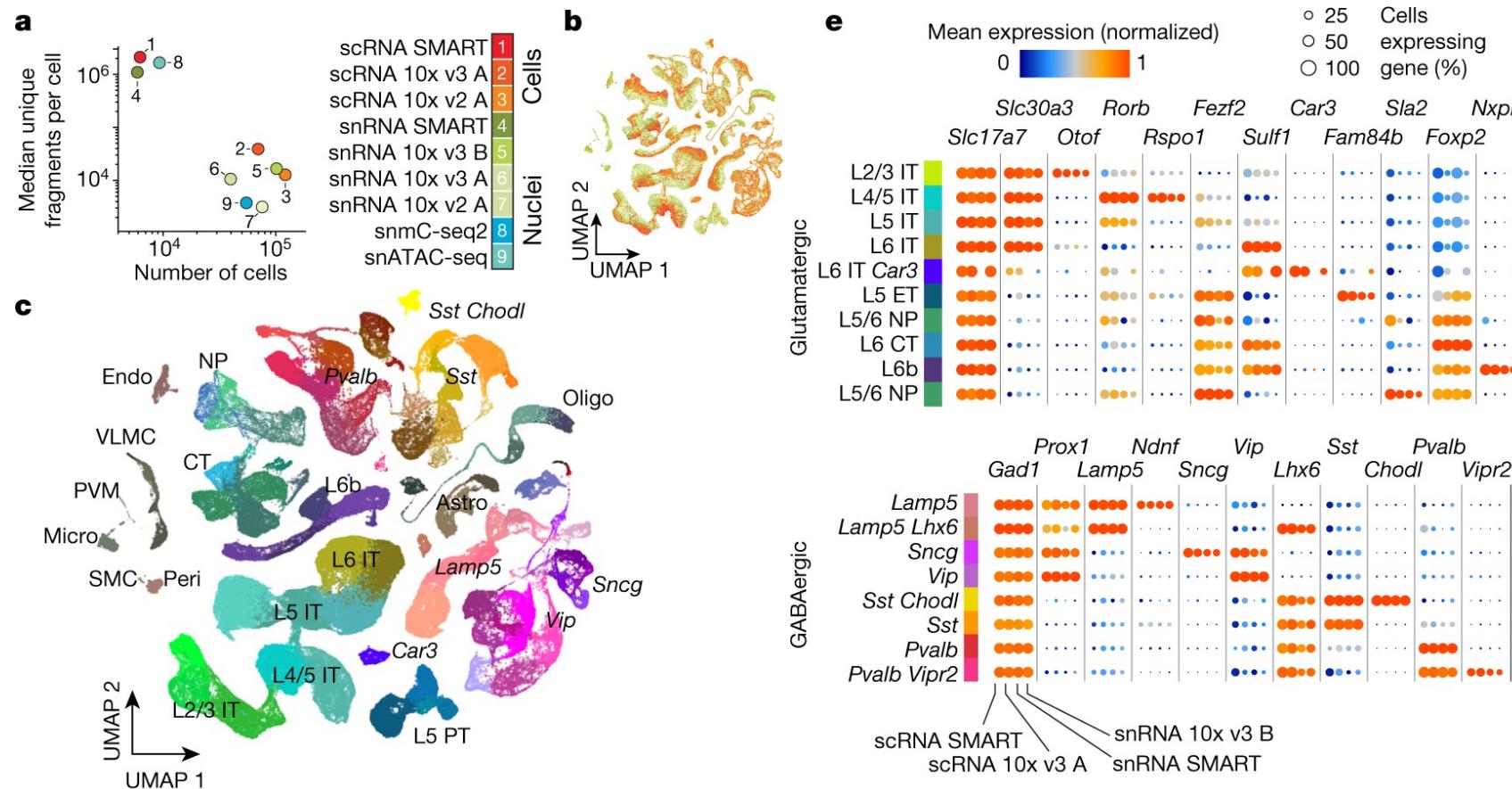
Article | Open Access | Published: 06 October 2021

A transcriptomic and epigenomic cell atlas of the mouse primary motor cortex

Zizhen Yao, Hanqing Liu, [...] Eran A. Mukamel

[Nature](#) 598, 103–110 (2021) | [Cite this article](#)

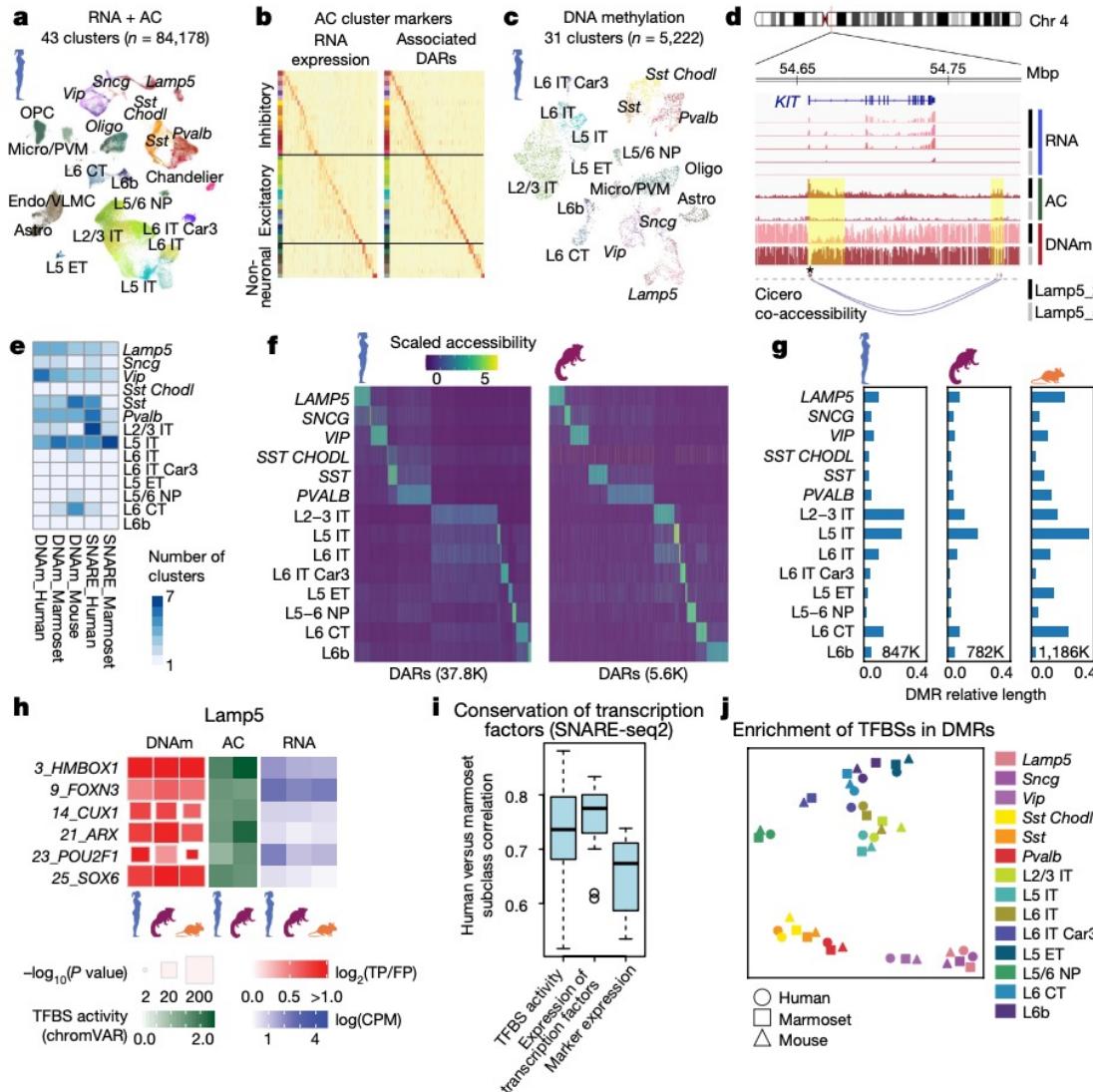
Metrics



- 526 k RNA-seq
 - 10k snmC-seq
 - 81k ATAC-seq

Comparative cellular analysis of motor cortex in human, marmoset and mouse

Trygve E. Bakken Nikolas L. Jorstad, [...] Ed S. Lein

Nature 598, 111–119 (2021) | [Cite this article](#)41 Altmetric | [Metrics](#)

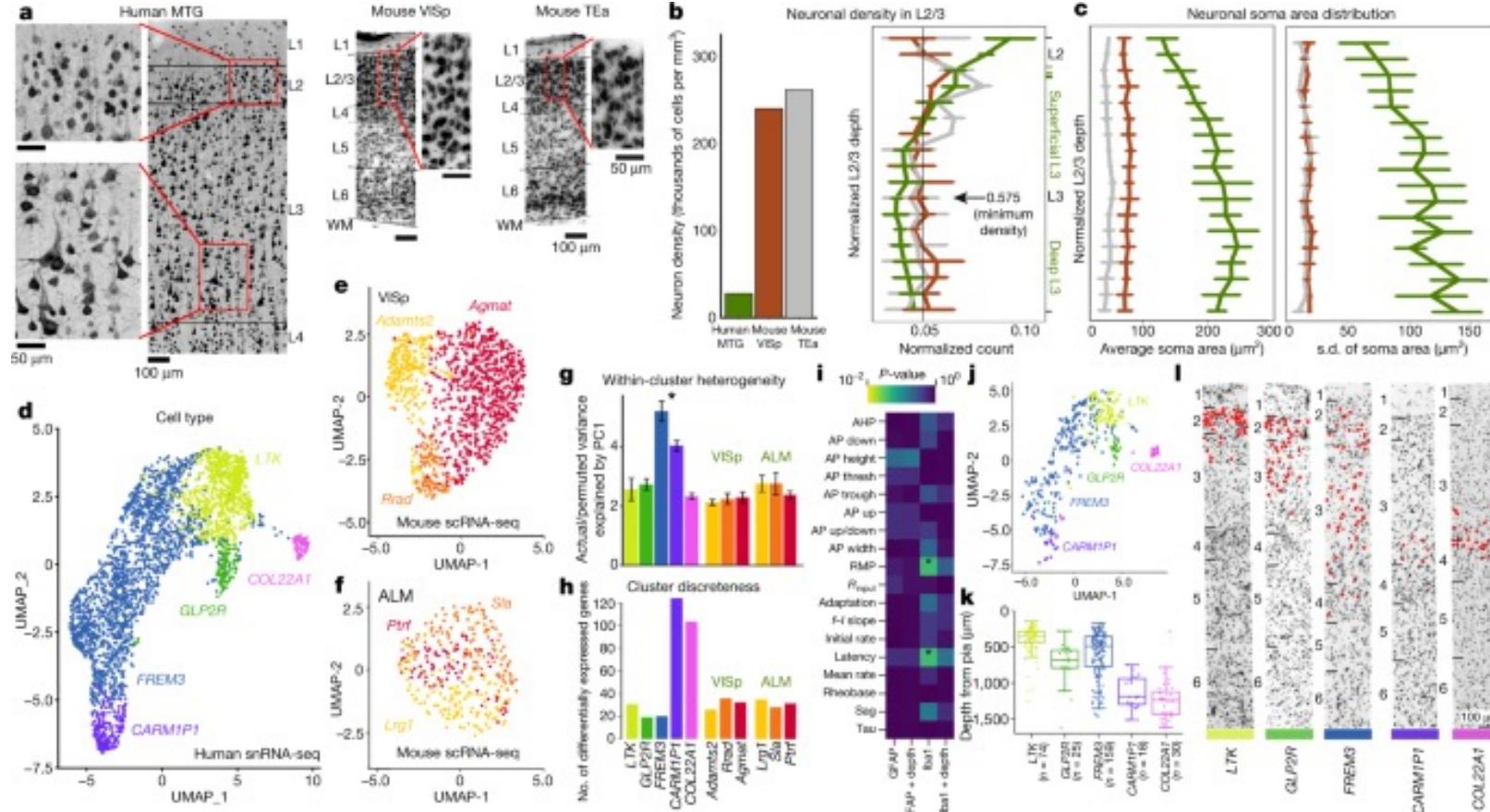
- SMART-seq V4
- Droplet Chromium V3
- snmC-seq2
- SNARE-seq2

Human neocortical expansion involves glutamatergic neuron diversification

Jim Berg, Staci A. Sorensen, [...] Ed S. Lein 

Nature 598, 151–158 (2021) | [Cite this article](#)

[Metrics](#)



- Patch-seq
(electrophysiological + molecular)

BICCN – DNA-methylation atlas

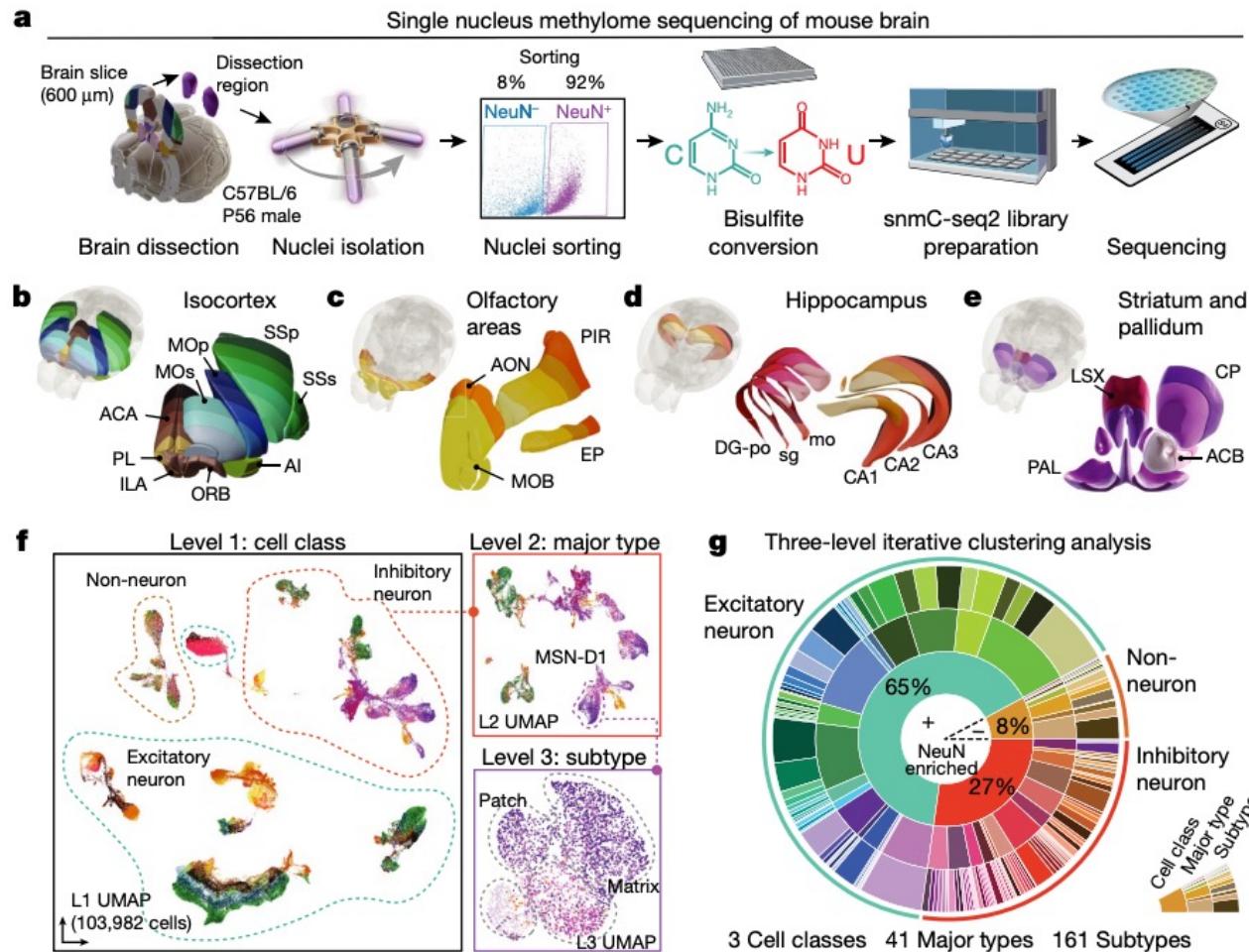
Article | Open Access | Published: 06 October 2021

DNA methylation atlas of the mouse brain at single-cell resolution

Hanqing Liu, Jingtian Zhou, [...] Joseph R. Ecker [✉](#)

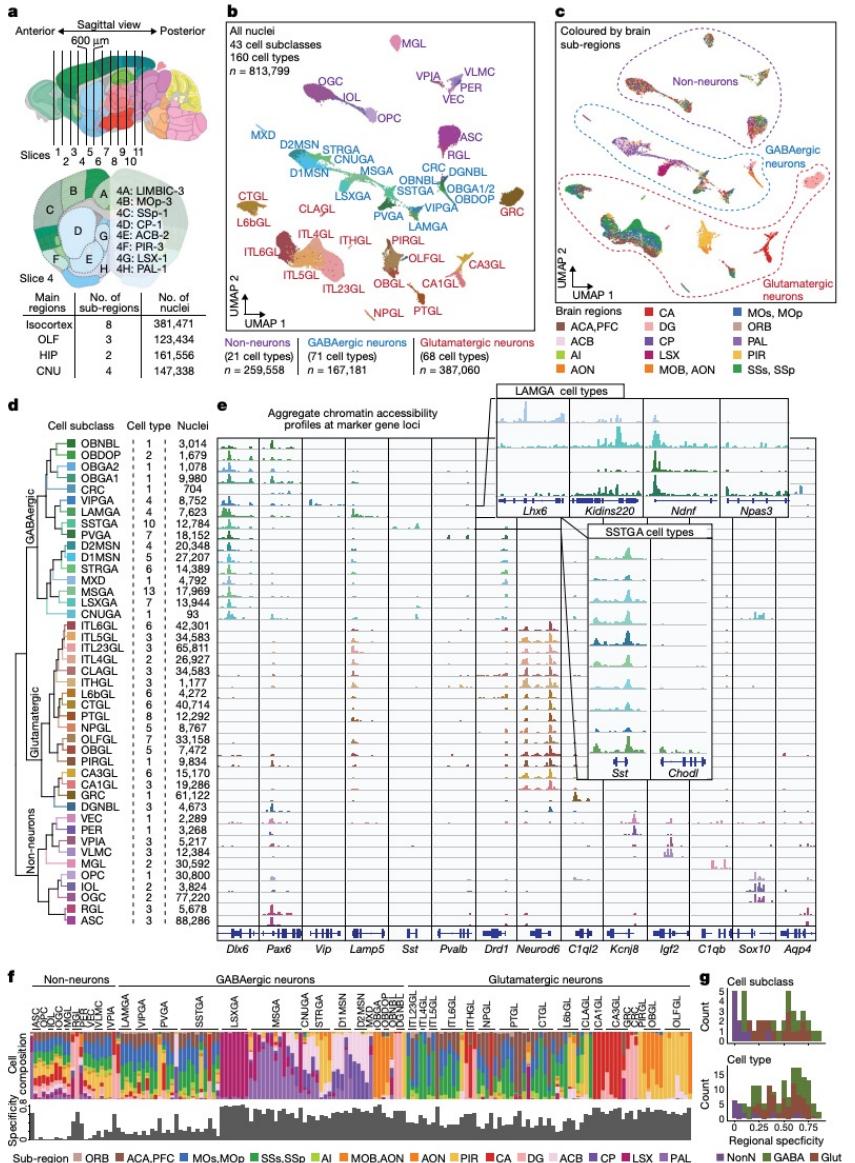
Nature 598, 120–128 (2021) | [Cite this article](#)

Metrics



- 104K snmC-seq2

BICCN – chromatin accessibility atlas



Article | Open Access | Published: 06 October 2021

An atlas of gene regulatory elements in adult mouse cerebrum

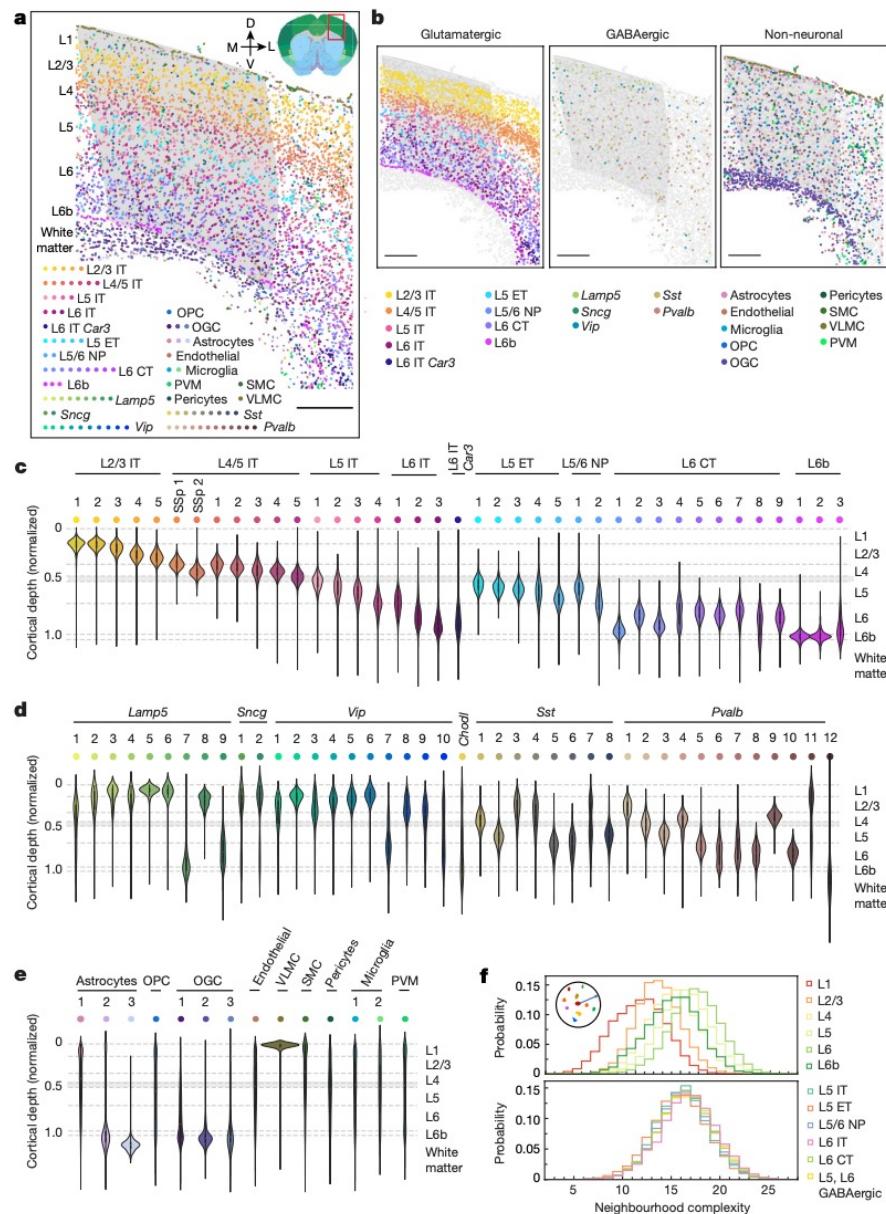
Yang Eric Li, Sebastian Preissl, [...] Bing Ren

Nature 598, 129–136 (2021) | Cite this article

Metrics

- 814K snATAC-seq

BICCN – spatial atlas



Article | Open Access | Published: 06 October 2021

Spatially resolved cell atlas of the mouse primary motor cortex by MERFISH

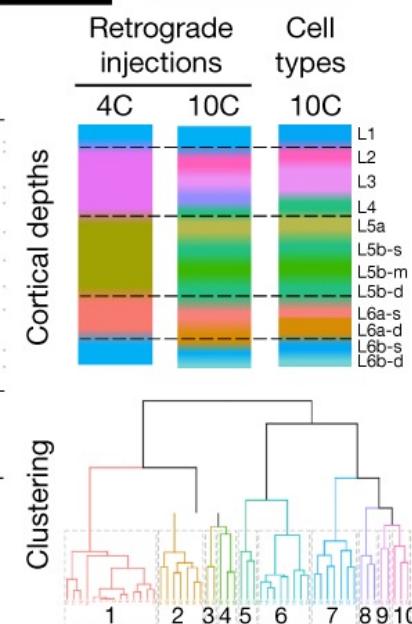
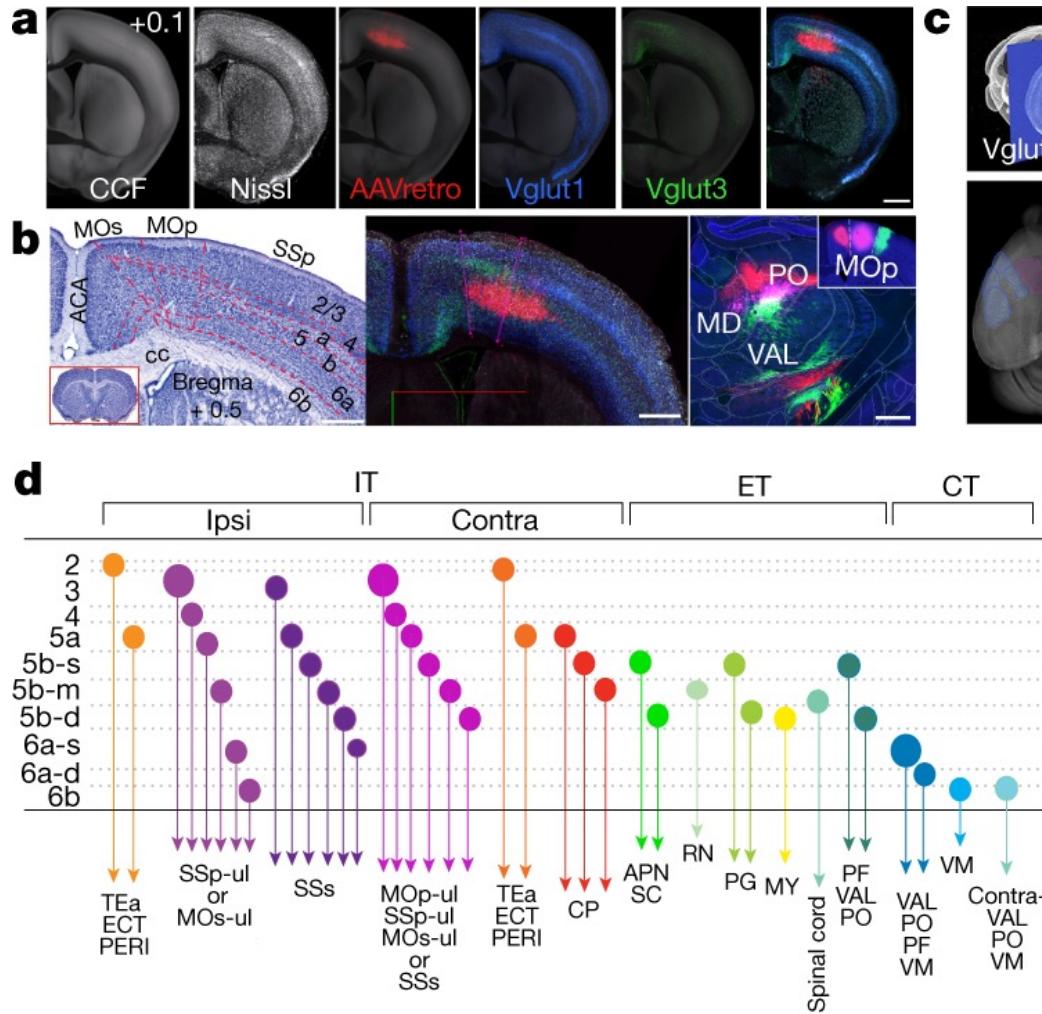
Meng Zhang, Stephen W. Eichhorn, Brian Zingg, Zichen Yao, Kaelan Cotter, Hongkui Zeng, Hongwei Dong & Xiaowei Zhuang

Nature 598, 137–143 (2021) | Cite this article

5 Altmetric | Metrics

- 300K MERFISH

BICCN – 3D anatomy structure



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Cellular anatomy of the mouse primary motor cortex

Rodrigo Muñoz-Castañeda, Brian Zingg, [...] Hong-Wei Dong

Nature 598, 159–166 (2021) | [Cite this article](#)

Metrics

- Genetic and viral labelling,
- Barcoded anatomy by sequencing
- Whole-brain imaging
-

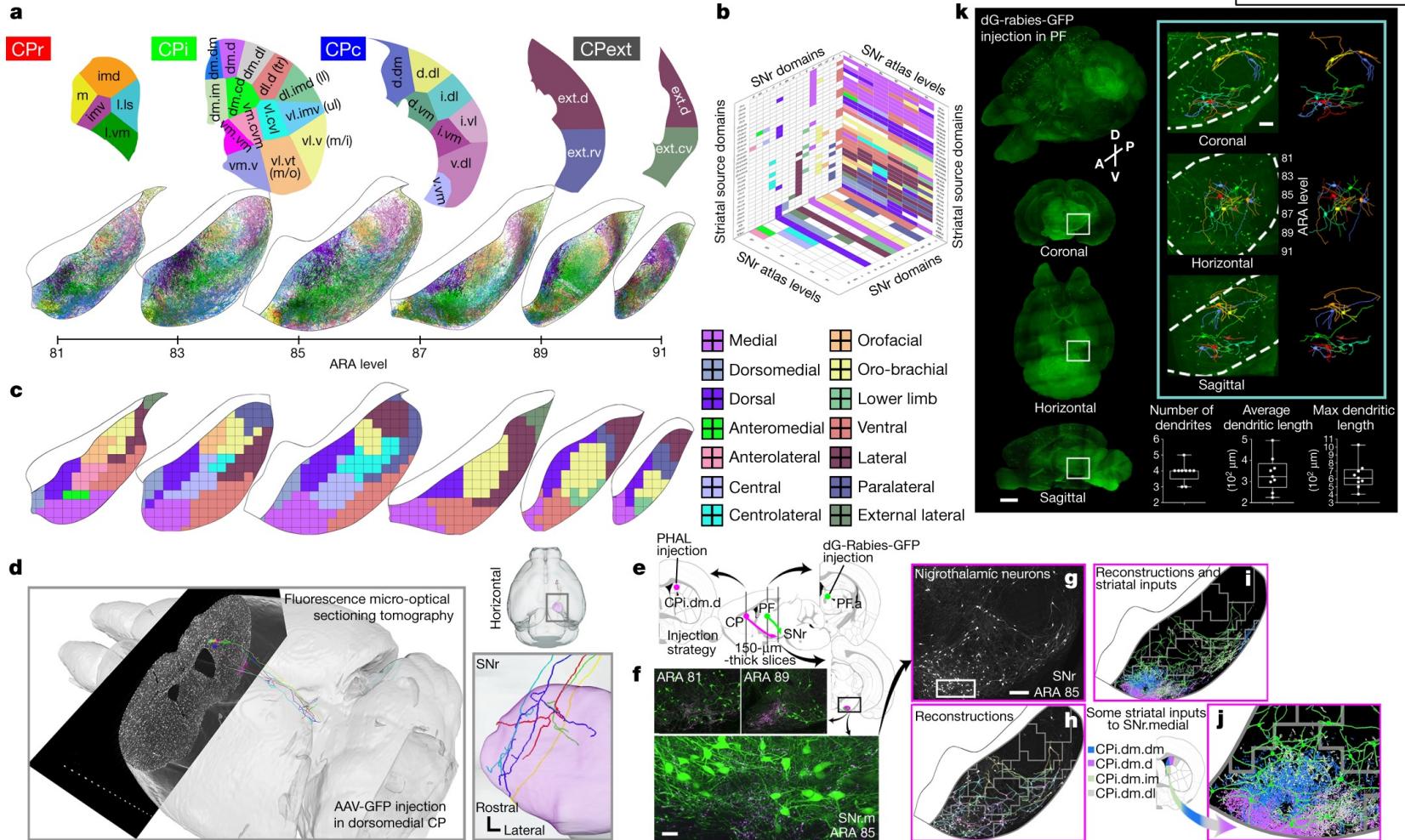
The mouse cortico–basal ganglia–thalamic network

Nicholas N. Foster Joshua Barry, [...] Hong-Wei Dong

Nature 598, 188–194 (2021) | Cite this article

[Metrics](#)

BICCN – network



BICCN – retrograde labelling with DNA-methylation sequencing

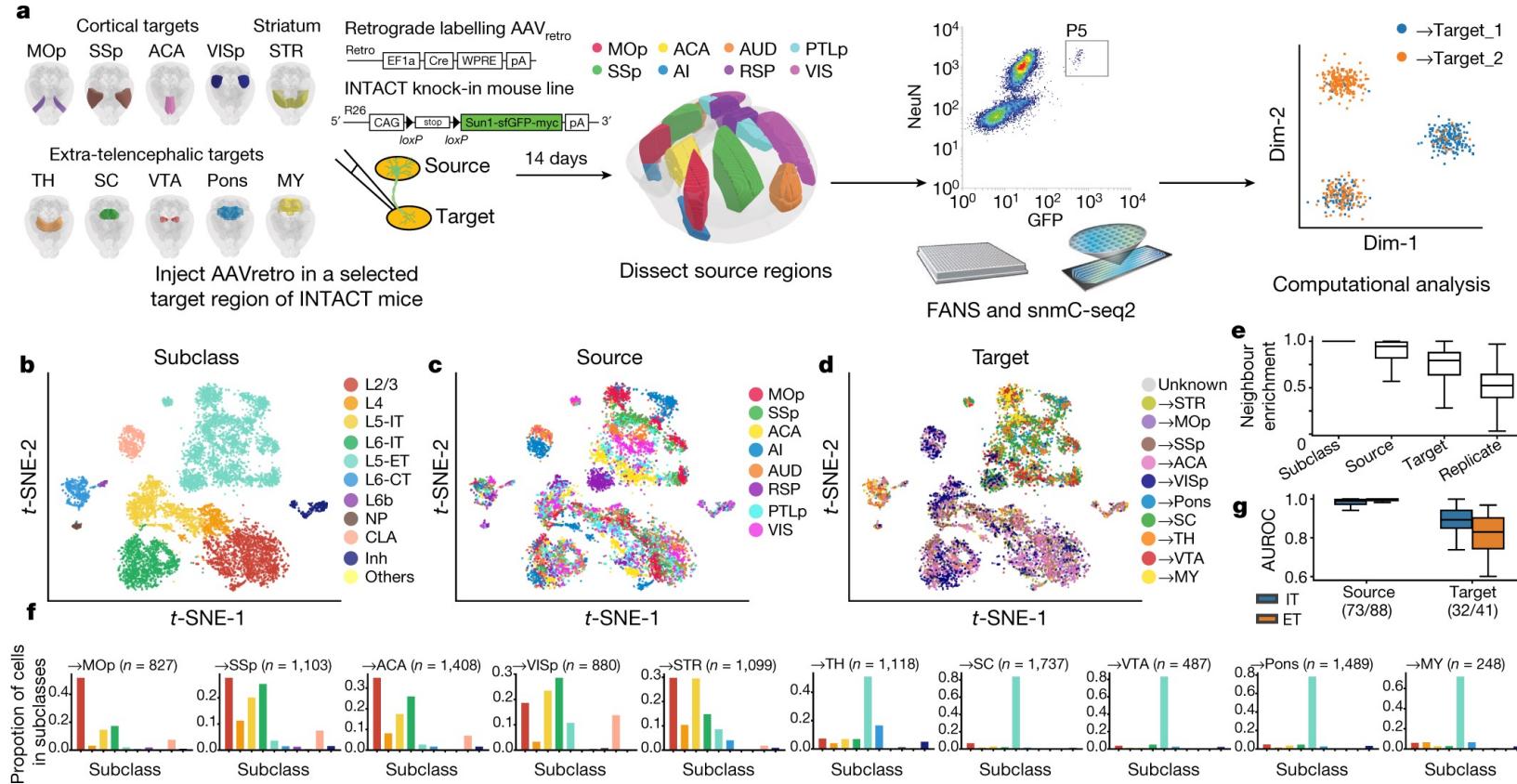
Article | Open Access | Published: 06 October 2021

Epigenomic diversity of cortical projection neurons in the mouse brain

Zhuzhu Zhang, Jingtian Zhou, [...] Edward M. Callaway [✉](#)

Nature 598, 167–173 (2021) | [Cite this article](#)

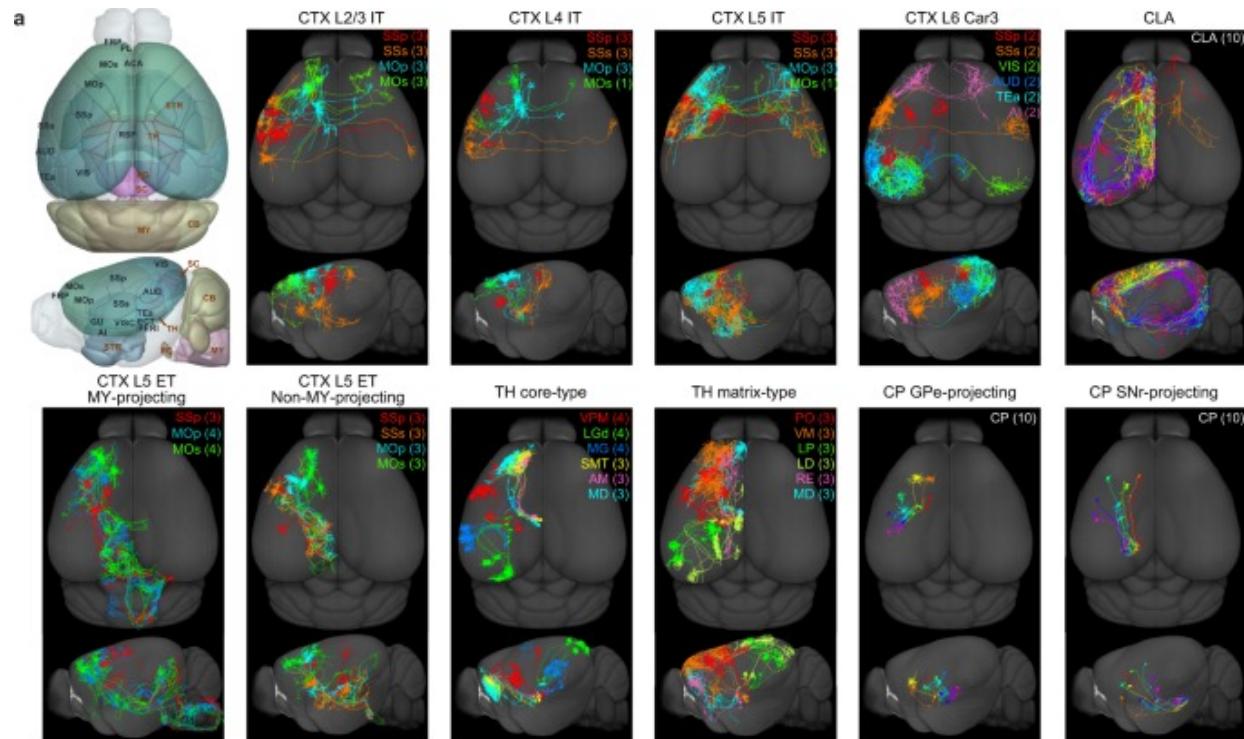
[Metrics](#)



- 12k Epi-retro-seq

“To address these questions, we developed epi-retro-seq, which applies single-nucleus methylome sequencing (snmC-seq)⁸ to neurons dissected from cortical source regions that were labelled on the basis of their long-distance projections to specific cortical and subcortical targets (Fig. 1a)’

BICCN – morphological + molecular reconstruction



b

Morphological features	Brain structures		Cortex				Thalamus		Striatum	
	Driver lines used		Cux2	Pixnd1	Gnb4	Fezf2, Pvalb	Trnif1, Vipr2	Trnif1, Vipr2, Pixnd1		
	Transcriptomic subclasses		L2/3 IT	L4 IT	L5 IT	Car3	L5 ET	Prkcd, Grin2c	Drd1	Drd2
	Projection classes			Corticocortical (intratelencephalic)			Corticofugal (extratelencephalic)		Thalamocortical	
	Projection types		L2/3 IT	L4 IT	L5 IT	L6 Car3	CLA	MY-projecting	Non-MY-projecting	Striatofugal
	Source regions analyzed in this study		MO _p , MO _s , SS _p , SS _s	MO _p , MO _s , SS _p , SS _s	MO _p , MO _s , SS _p , SS _s	Lateral cortex (ORBI, MO _p , SS _p , SS _s , AI, GU, VIS, AUD, TEa, etc.)	CLA	MO _p , MO _s , SS _p , SS _s	MO _p , MO _s , SS _p , SS _s	VPM, VPL, VPMpc, VPLpc, LGd, MG, VAL, AM, SMT, MD, PO, LP, LD, MD, IAD, VM, CM, (RE, PVT)
	Regional difference		Yes	Yes	Yes	Yes	-	Yes	Yes	Yes
	Projection patterns		Divergent	Divergent	Divergent	Divergent	Divergent	Divergent	Parallel	Divergent
	Average number of targets		5.4	2.4	11.3	14.3	20.9	16.2	8.9	2.9
	Topographic distribution		-	-	-	Yes	Yes	-	Yes	7.2
	Projection target variability among cells within a region		High	High	High	High	High	High	Low	3.0
									Medium	2.0
									Low	Yes
									Low	Yes

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Morphological diversity of single neurons in molecularly defined cell types

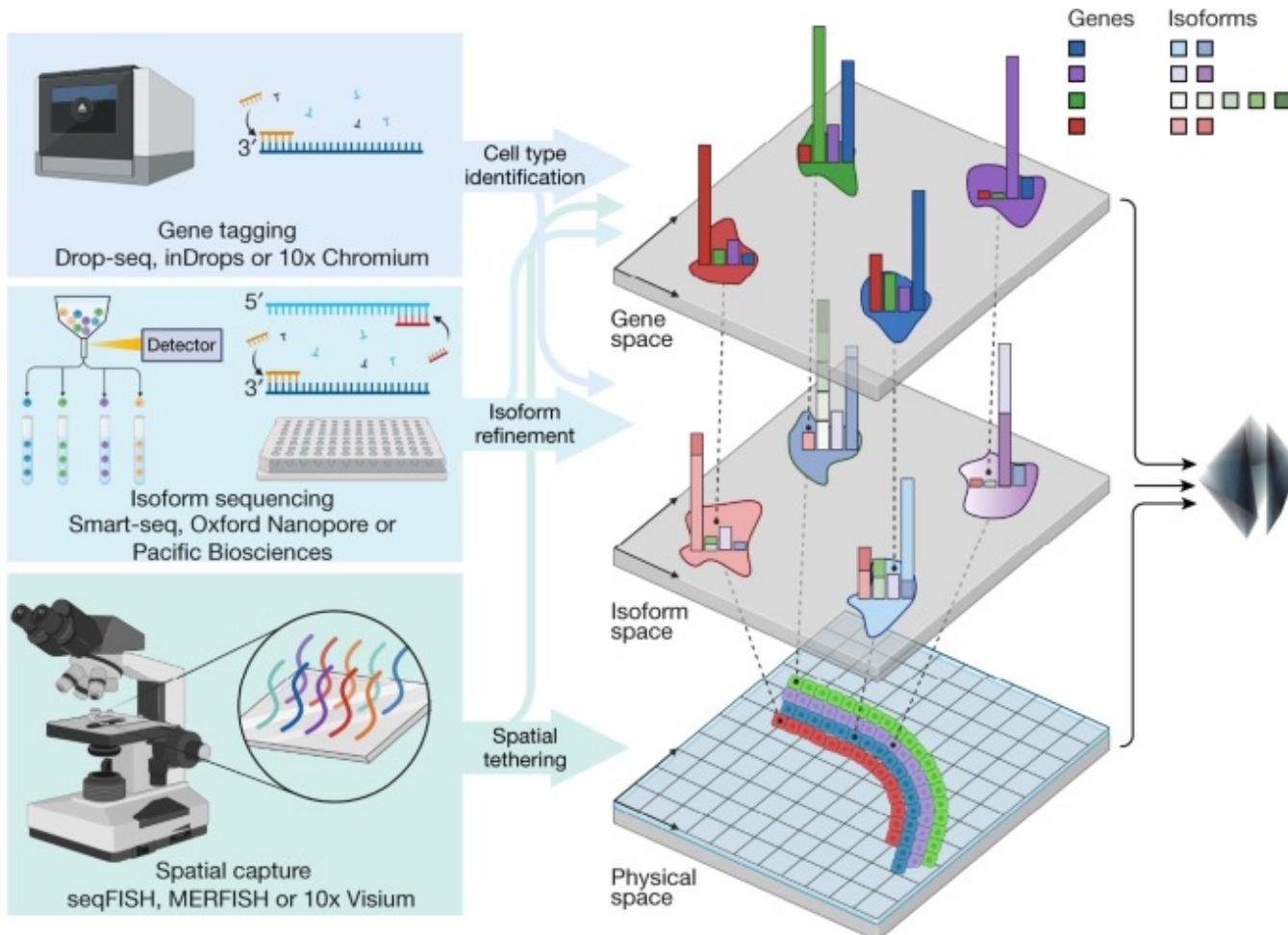
Hanchuan Peng , Peng Xie, [...] Hongkui Zeng

Nature 598, 174–181 (2021) | Cite this article

Metrics

- sparse labelling, whole-brain imaging, reconstruction, registration and analysis.

BICCN – Isoform



Article | Open Access | Published: 06 October 2021

Isoform cell-type specificity in the mouse primary motor cortex

A. Sina Booeshaghi, Zizhen Yao, Cindy van Velthoven, Kimberly Smith, Bosiljka Tasic, Hongkui Zeng & Lior Pachter [✉](#)

Nature 598, 195–199 (2021) | [Cite this article](#)

[Metrics](#)

- 6k smart-seq
- 280k MERFISH
- 94K 10X

BICCN – Dynamic signatures

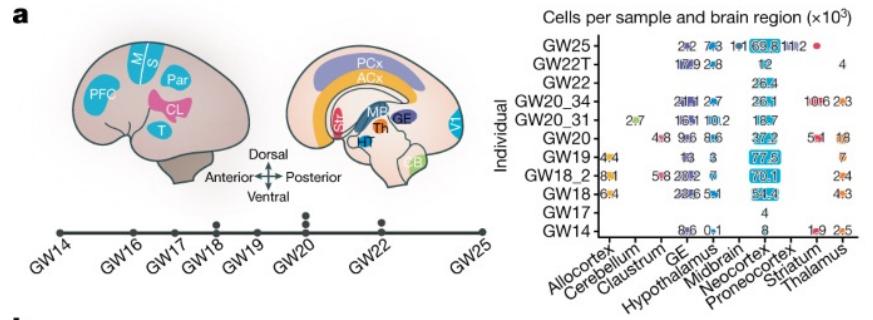
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An atlas of cortical arealization identifies dynamic molecular signatures

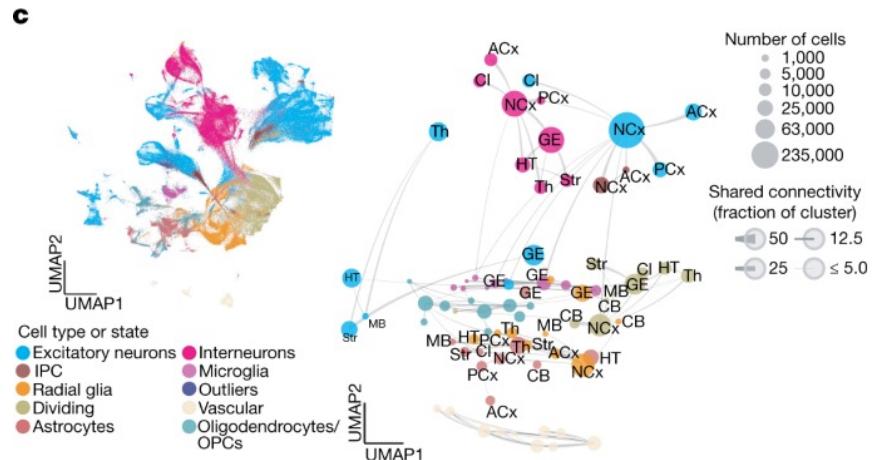
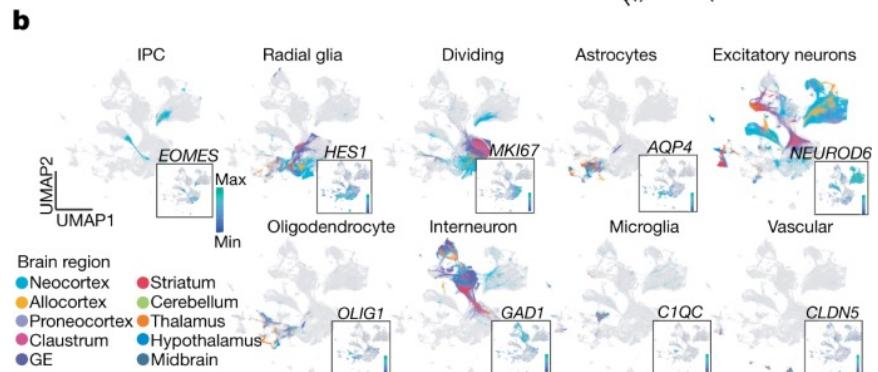
Aparna Bhaduri , Carmen Sandoval-Espinosa, Marcos Otero-Garcia, Irene Oh, Raymund Yin, Ugomma C. Eze, Tomasz J. Nowakowski & Arnold R. Kriegstein 

Nature 598, 200–204 (2021) | [Cite this article](#)

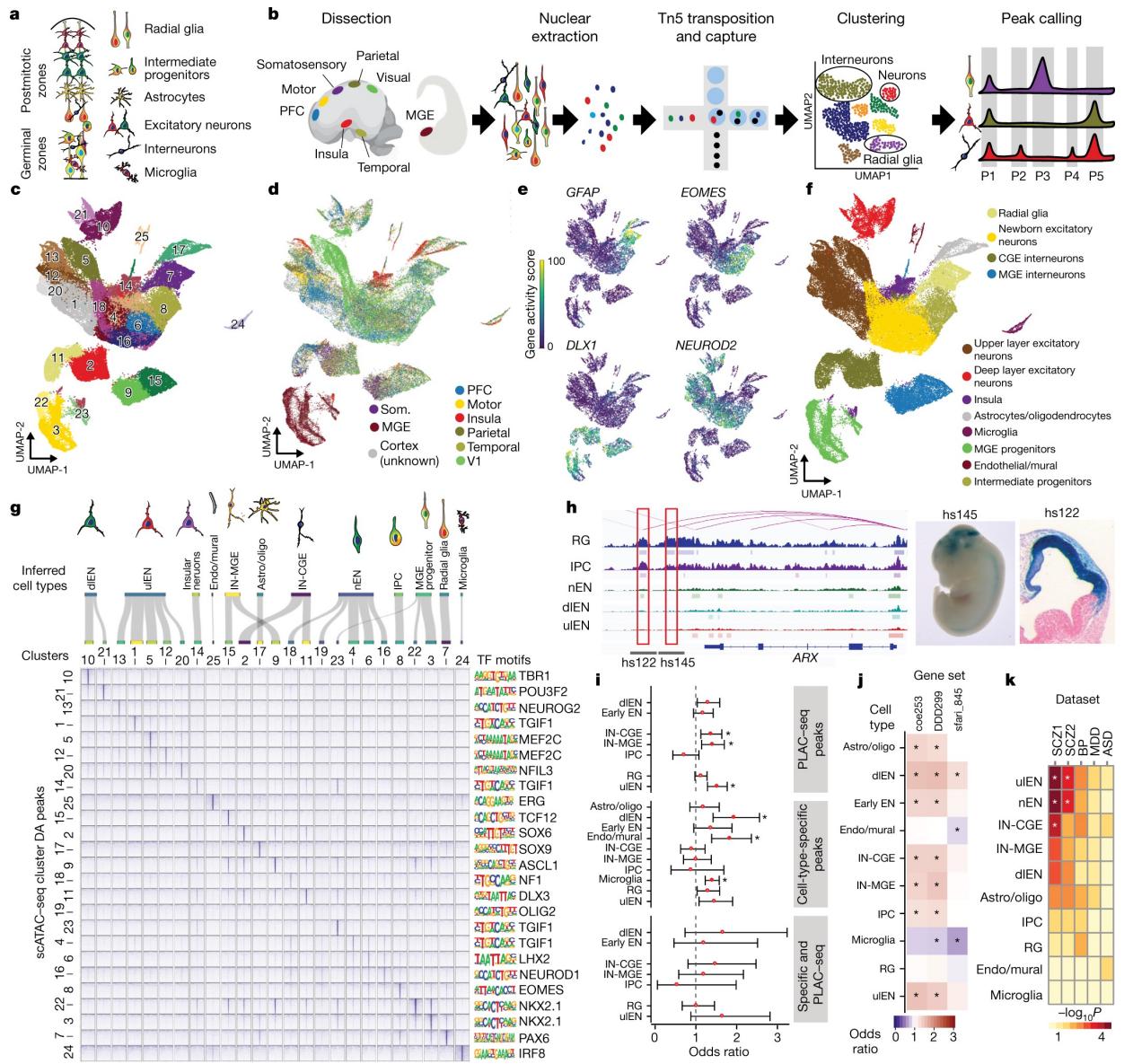
12 Altmetric | [Metrics](#)



- 700K RNA-Seq



BICCN – ATAC-seq



Article | Open Access | Published: 06 October 2021

Single-cell epigenomics reveals mechanisms of human cortical development

Ryan S. Ziffra, Chang N. Kim, Jayden M. Ross, Amy Wilfert, Tychele N. Turner, Maximilian Haeussler, Alex M. Casella, Paweł F. Przytycki, Kathleen C. Keough, David Shin, Derek Bogdanoff, Anat Kreimer, Katherine S. Pollard, Seth A. Ament, Evan E. Eichler, Nadav Ahituv & Tomasz J. Nowakowski

Nature 598, 205–213 (2021) | Cite this article

Metrics

- 77K ATAC-seq

BICCN – RNA-seq across lobules

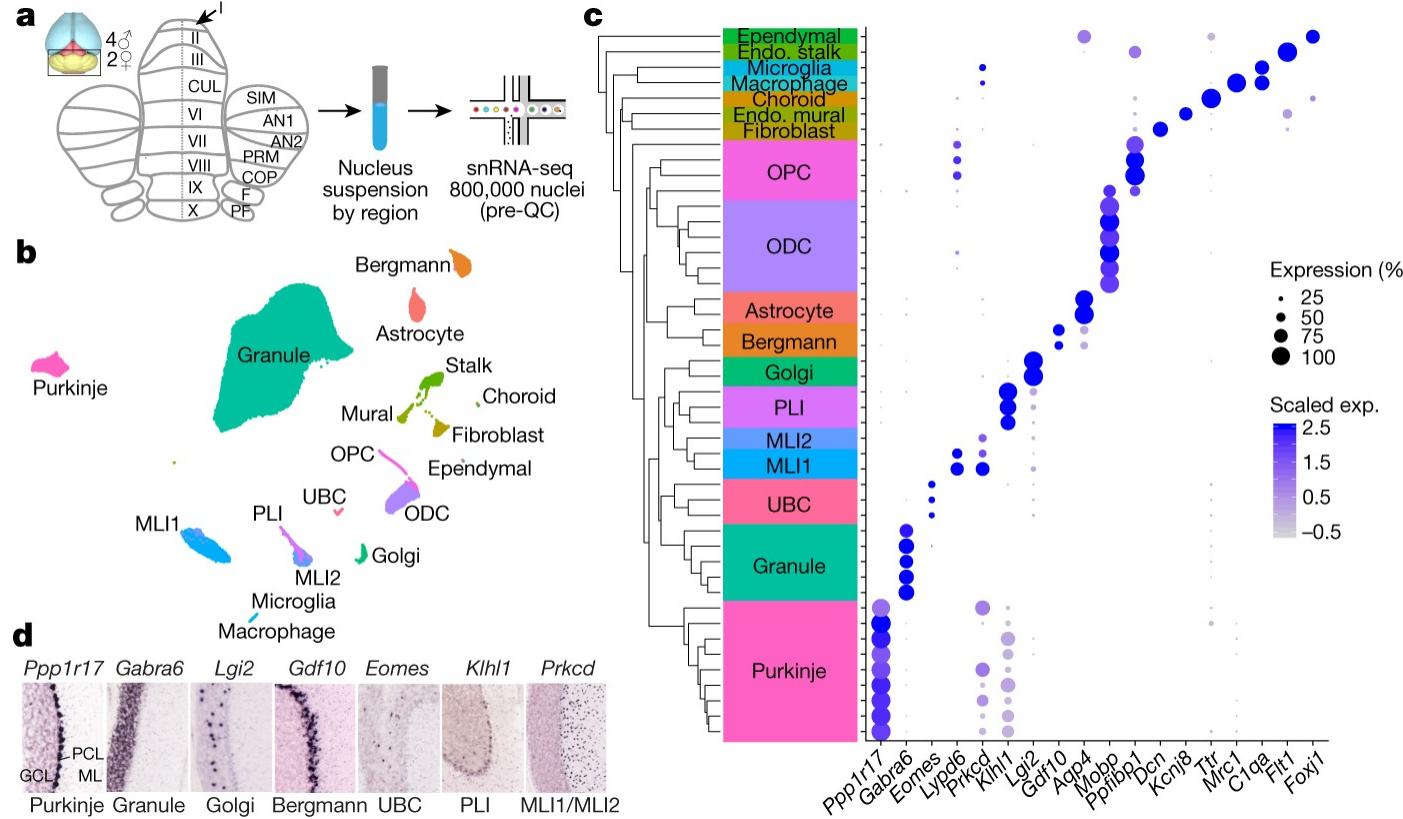
Article | Open Access | Published: 06 October 2021

A transcriptomic atlas of mouse cerebellar cortex comprehensively defines cell types

Velina Kozareva, Caroline Martin, Tomas Osorno, Stephanie Rudolph, Chong Guo, Charles Vanderburg, Naeem Nadaf, Aviv Regev, Wade G. Regehr & Evan Macosko [✉](#)

Nature 598, 214–219 (2021) | [Cite this article](#)

Metrics



- 611K snRNA-seq