

A fluorescence microscopy image of a neuronal network. The neurons are stained with a white marker, showing their cell bodies and extensive dendritic and axonal branching. The background is filled with a dense network of red-stained fibers, likely representing another cell type or a specific protein expression. The overall appearance is a complex, interconnected web of biological structures.

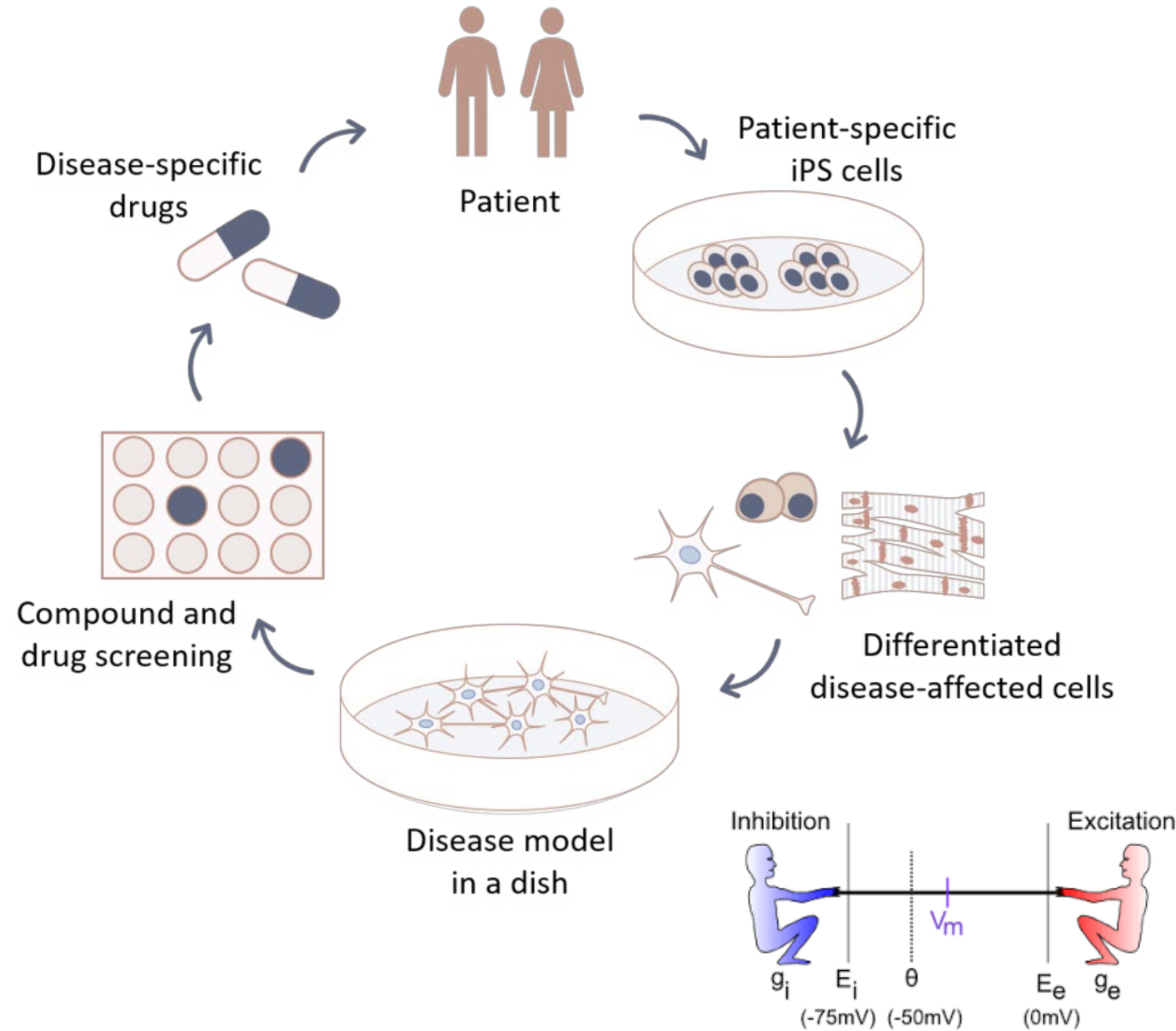
Understanding the cell-type specific function of EHMT1 in neuronal network function

Nael Nadif Kasri, PhD
Donders Institute, Radboudumc Nijmegen

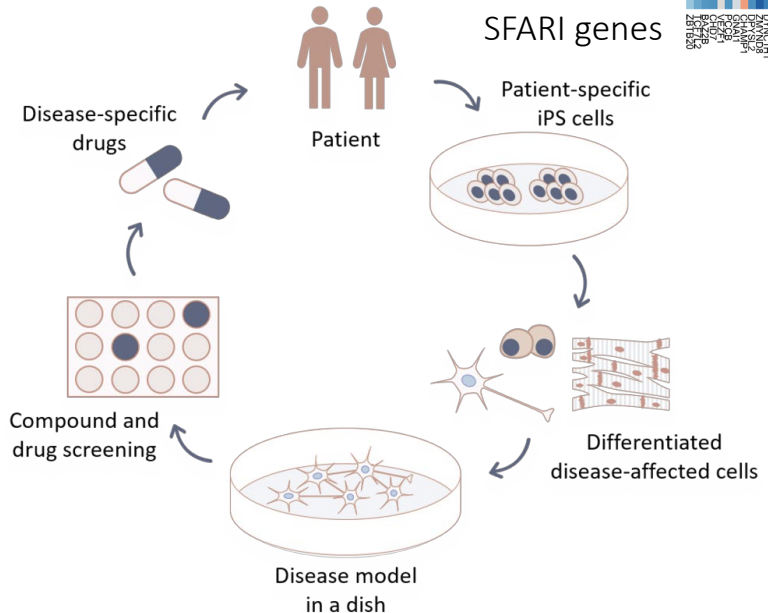
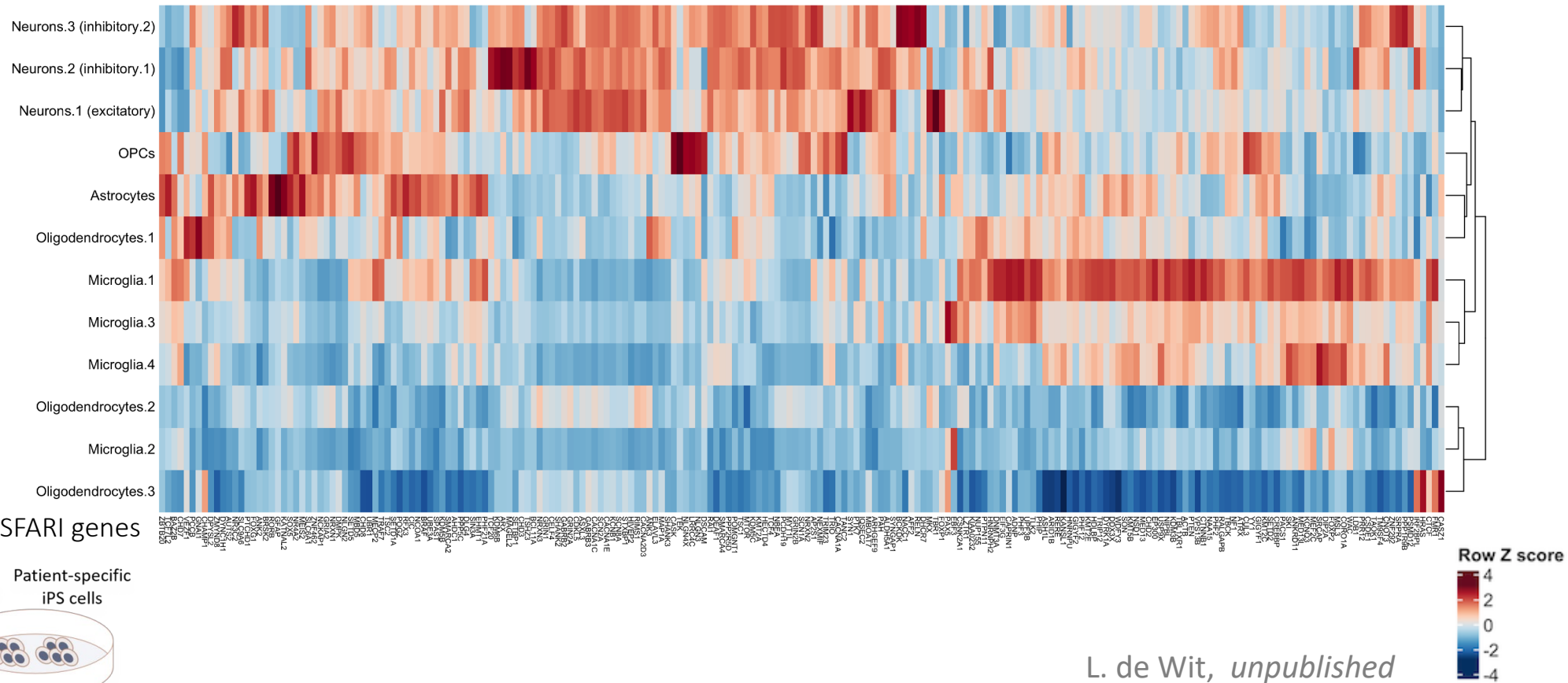
Kleefstra syndrome Conference, Ljubljana

June 1-2 2023

iPSC-derived models for translational studies: Opportunities and challenges

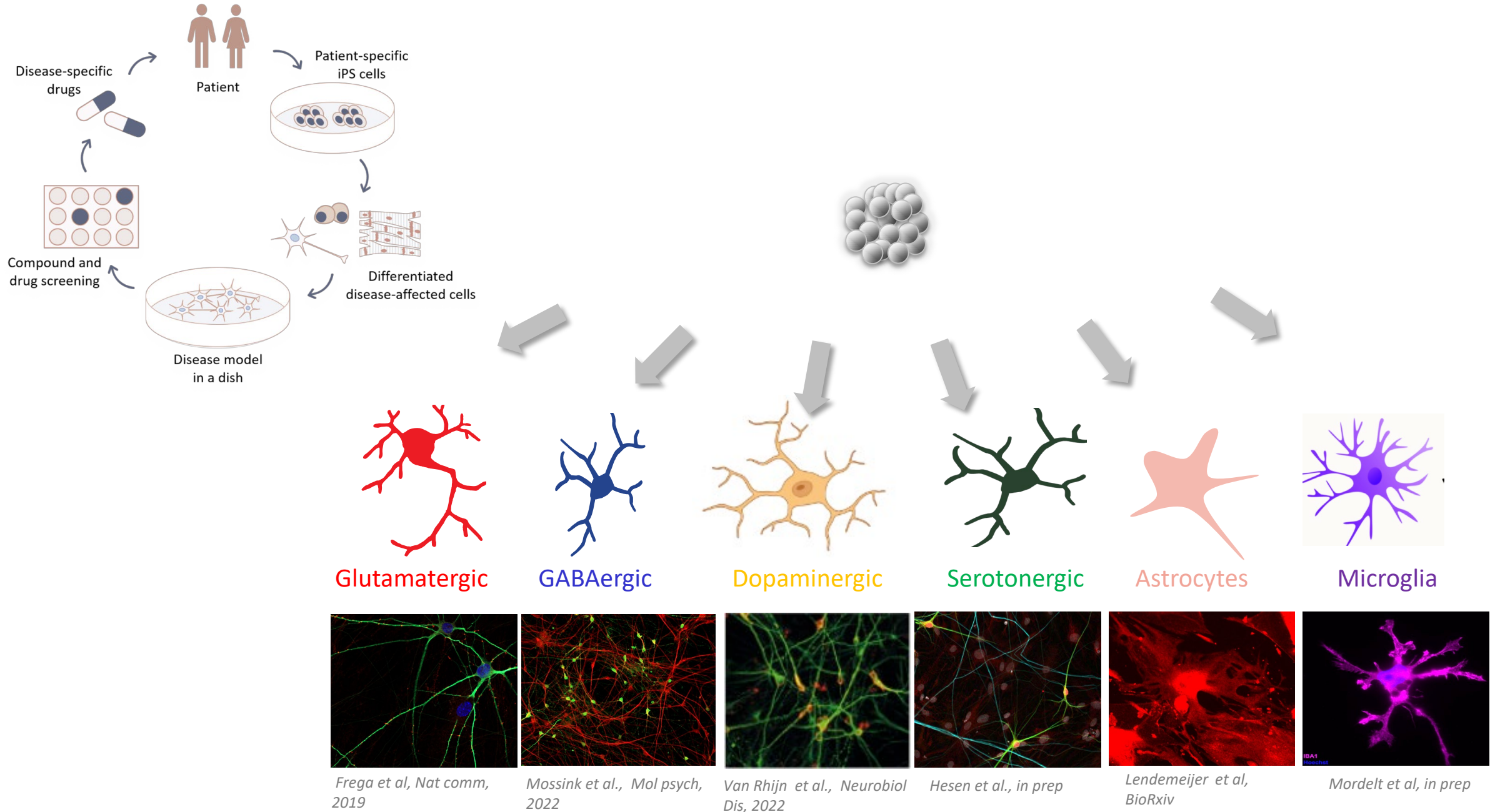


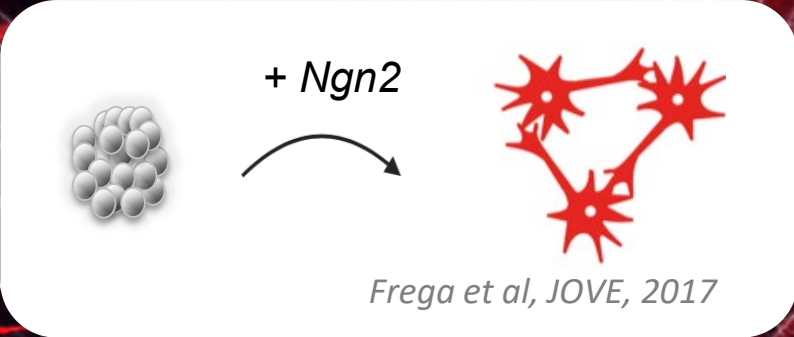
iPSC-derived models for translational studies: Opportunities and challenges



L. de Wit, unpublished

iPSC-derived models for translational studies: directed differentiation





Fast protocol
± 3 weeks

High efficiency

Mature neurons

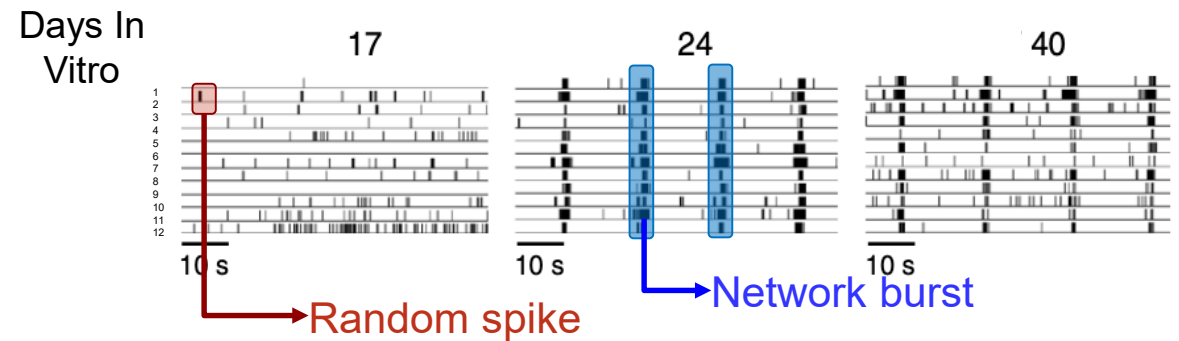
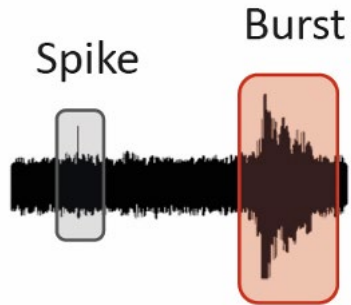
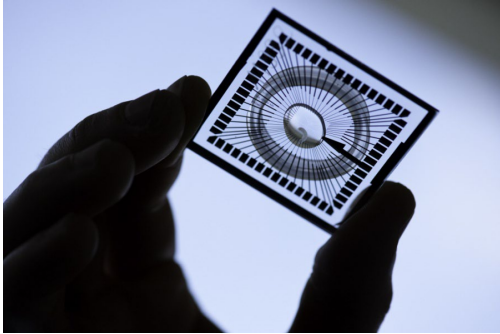
Homogeneous
population

Glia dependent

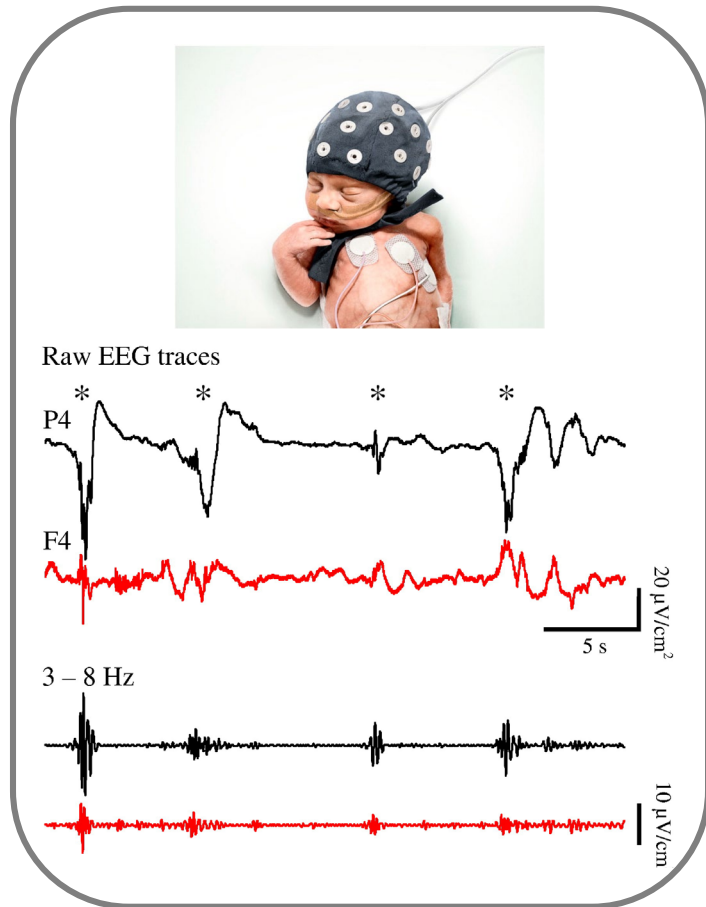


Measuring neural network activity of iPSC-derived networks

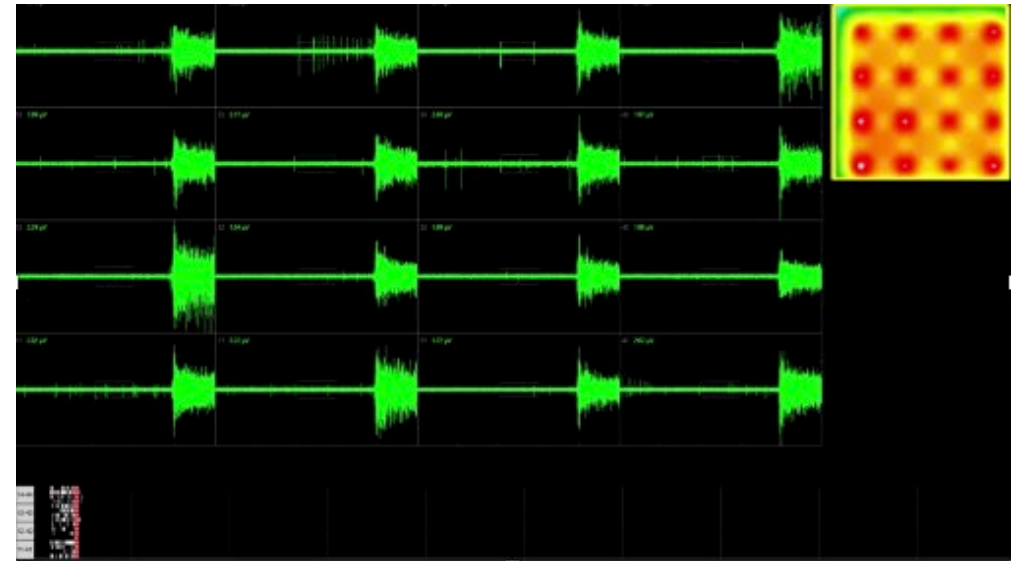
Micro-electrode arrays



Measuring neural network activity of iPSC-derived networks

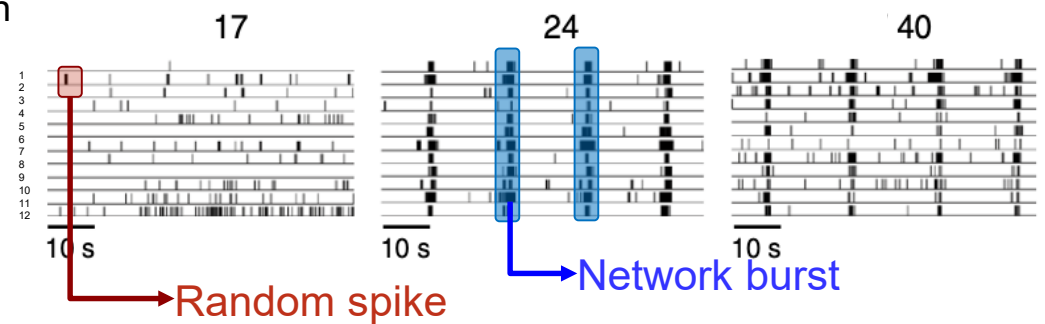


Synchronized Bursts are observed in human brain

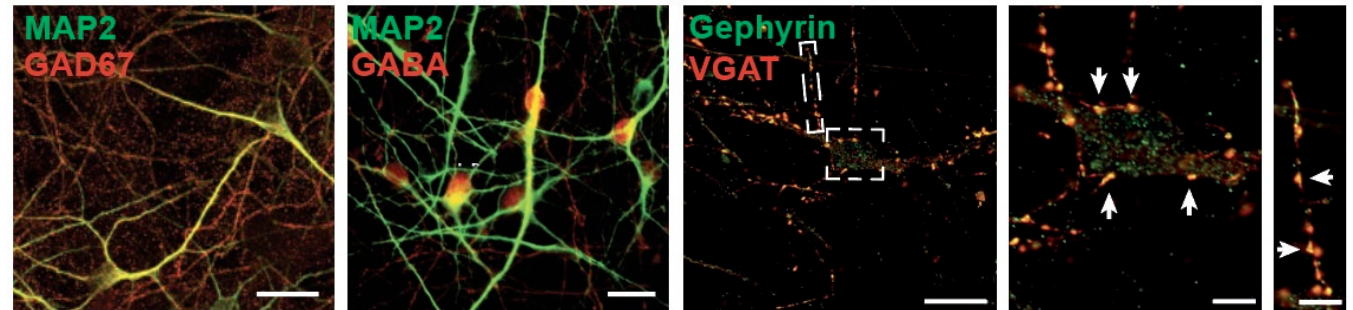
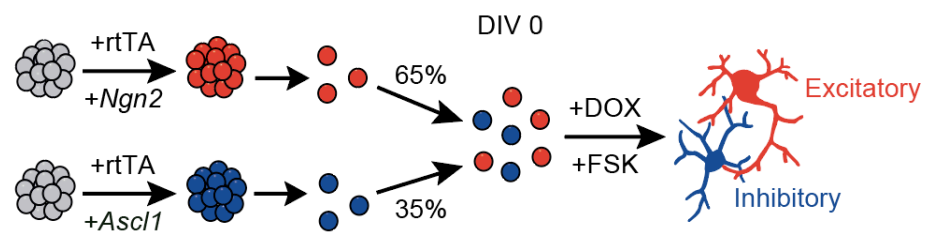
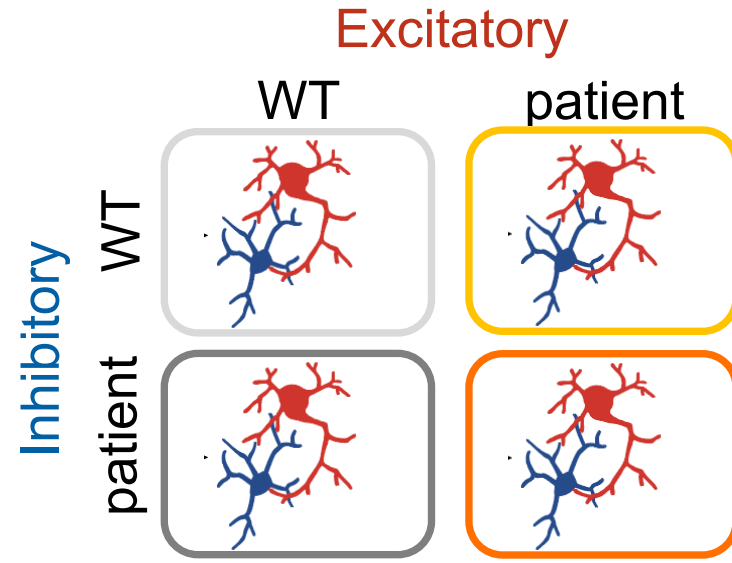
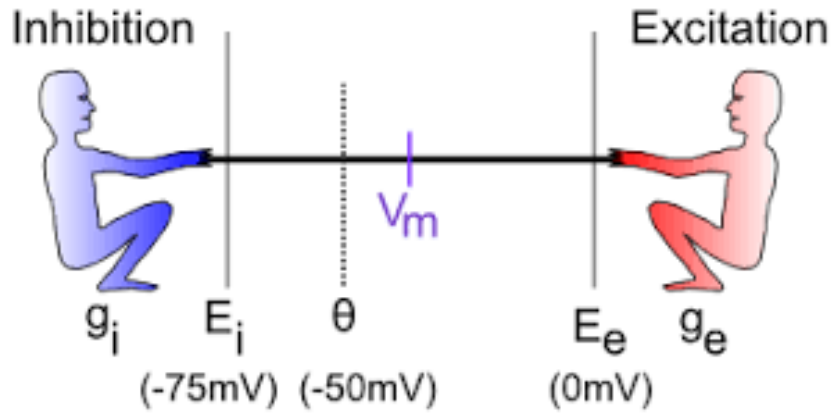


Development

Days In Vitro

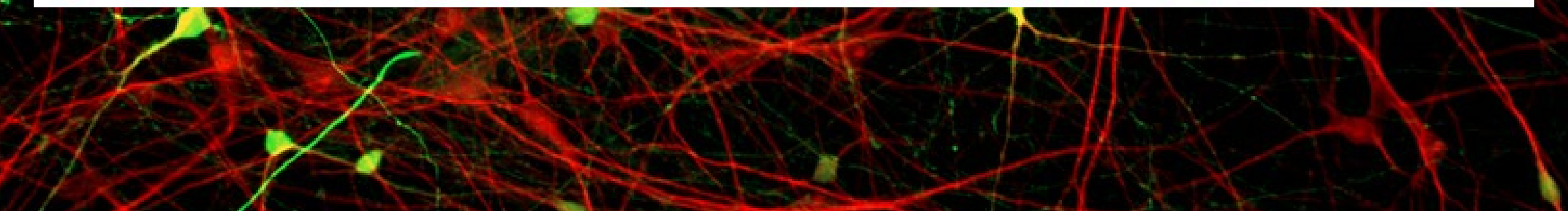
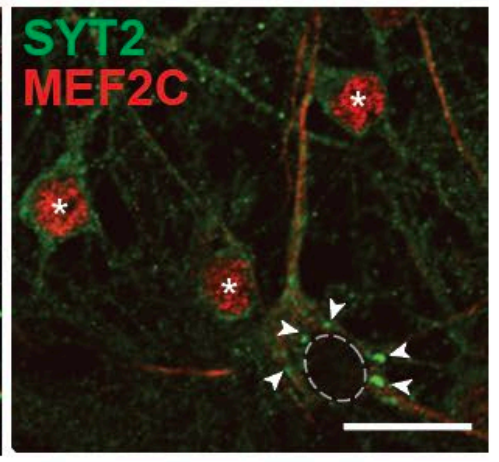
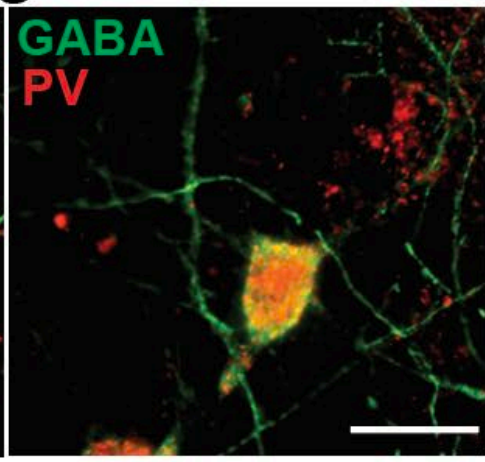
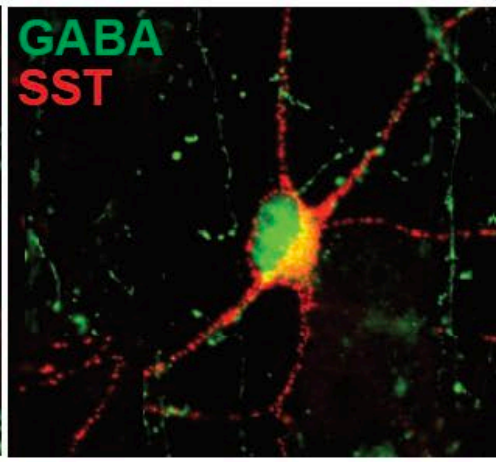
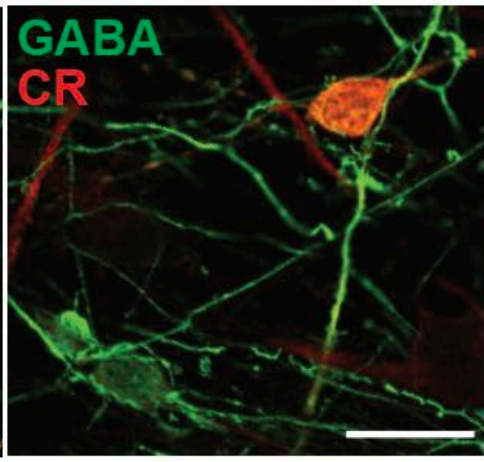
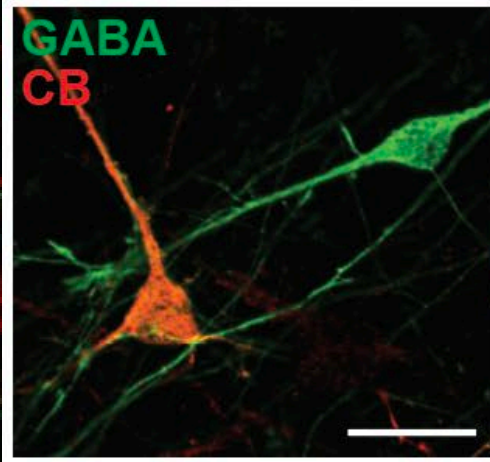
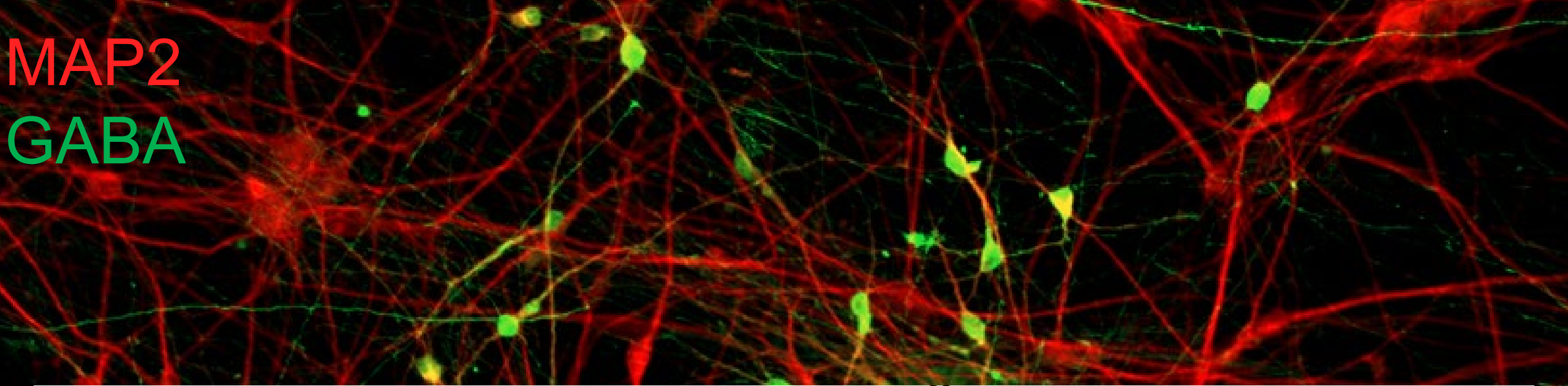


Probing for E/I balance in human pluripotent stem cell-derived neuronal networks

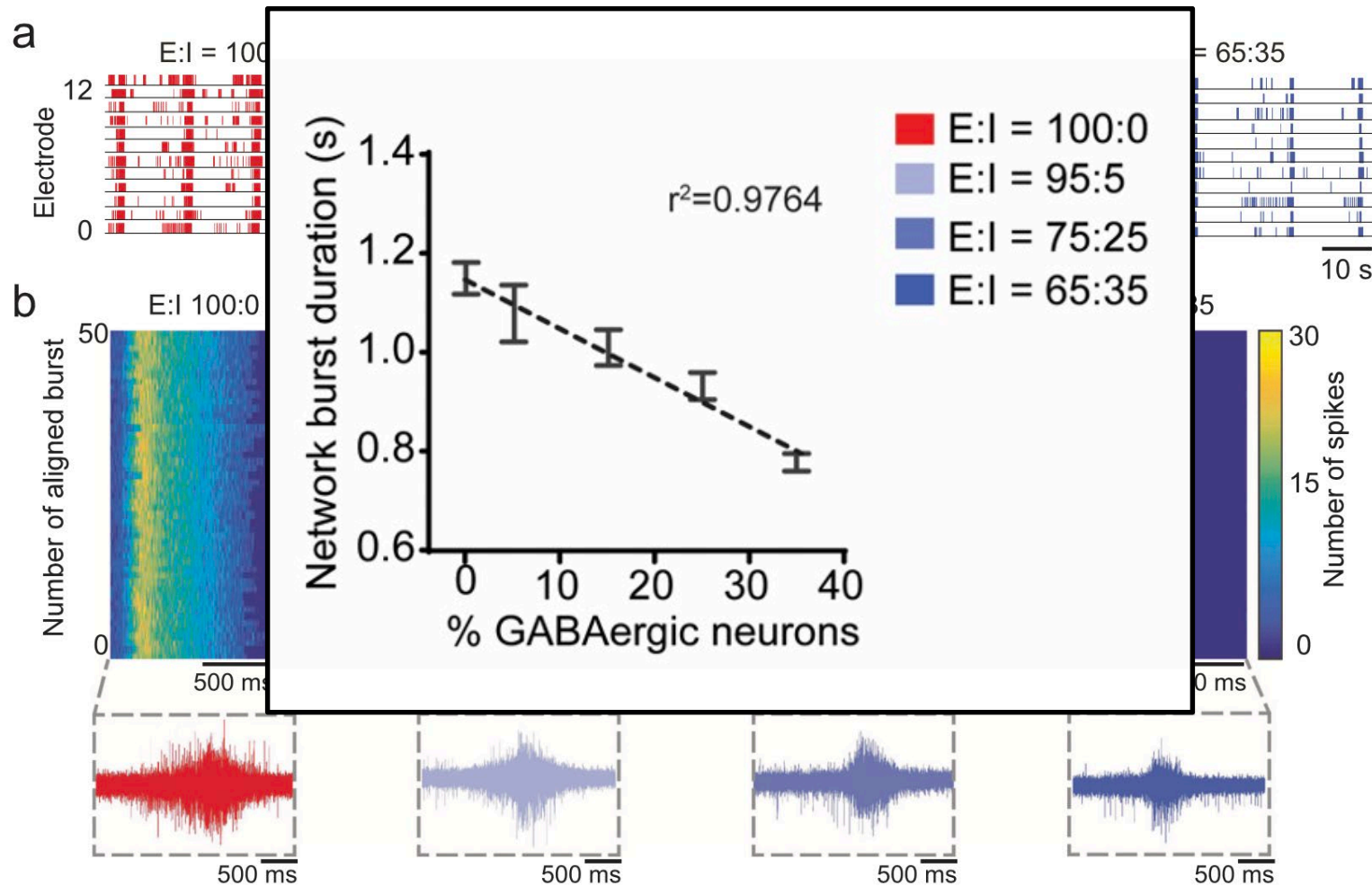
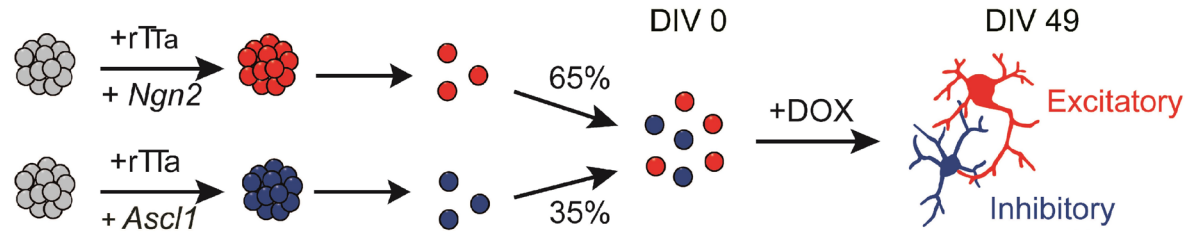


- Adapted from Yang et al., 2017
- Adapted from Shi, Z., et al., 2016

MAP2
GABA

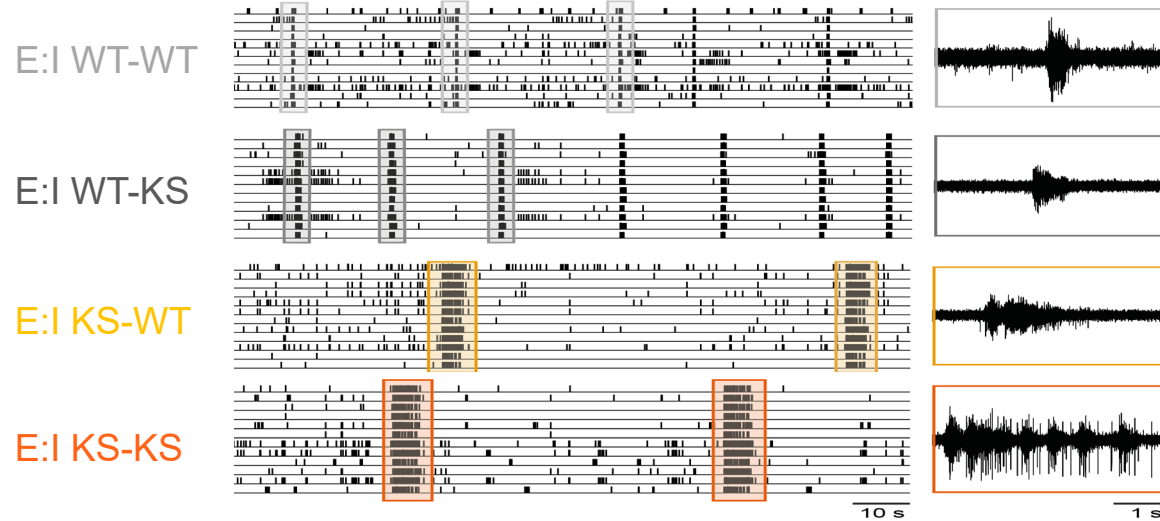
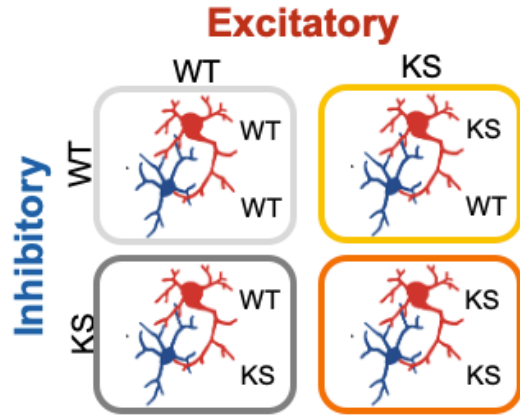


iGABAergic neurons exhibit scalable functional inhibitory control

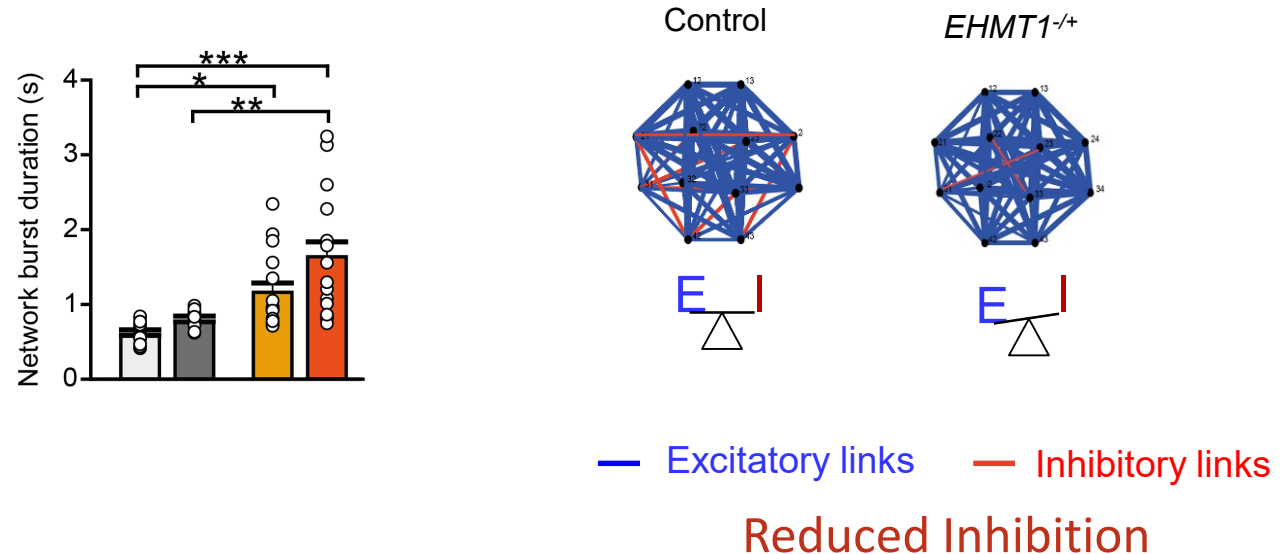
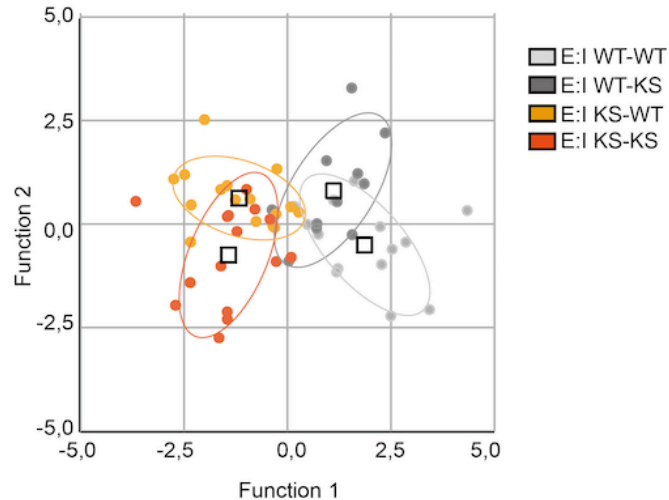


Contribution of excitation and inhibition to Kleefstra syndrome neuronal network phenotypes

KS: Haploinsufficiency of the *EHMT1* gene

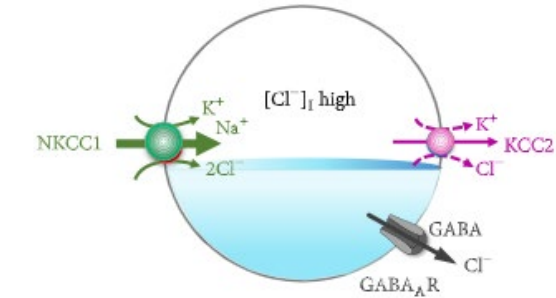


Discriminant analysis on all MEA parameters

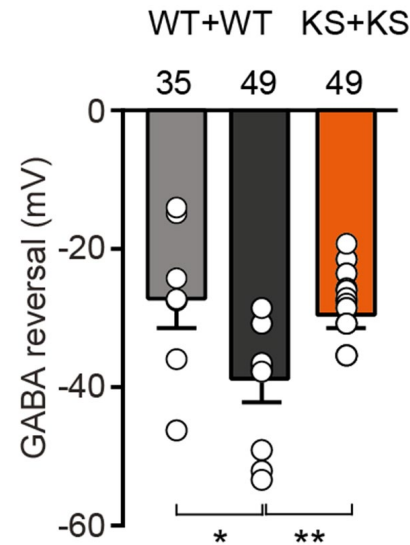
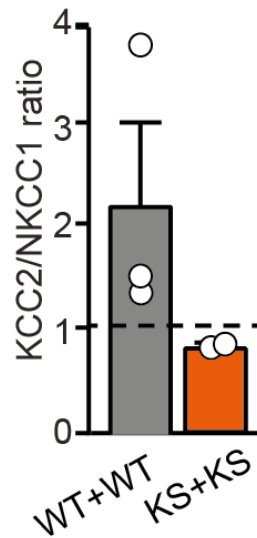
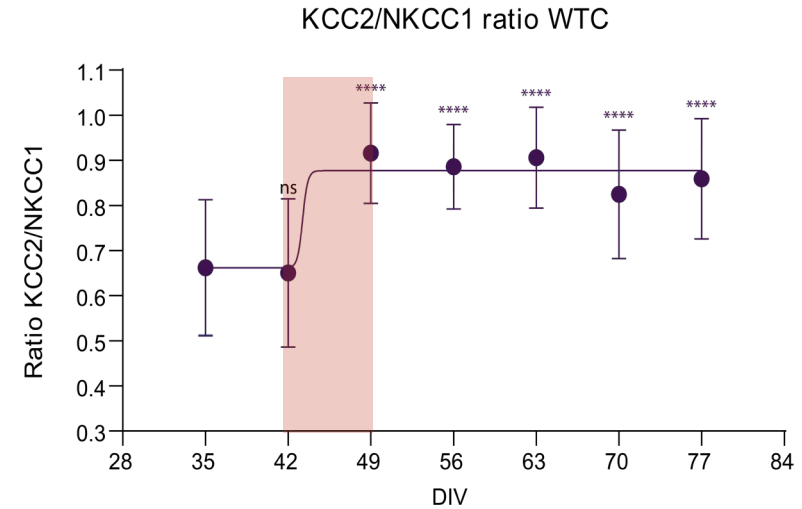
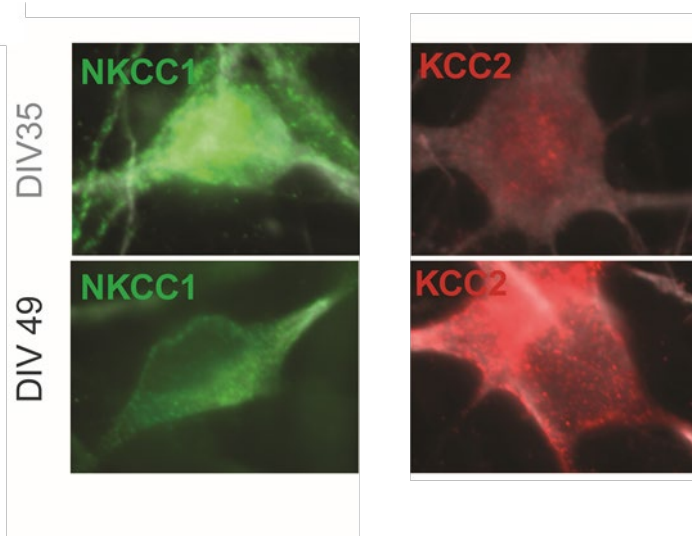
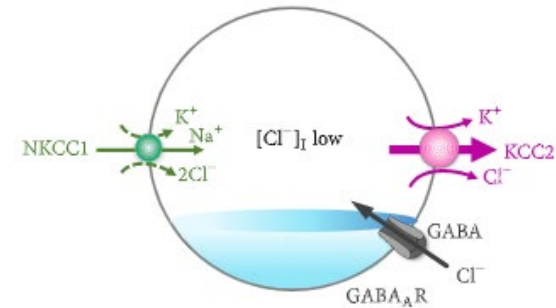


Delayed GABAergic maturation in Kleefstra neuronal networks

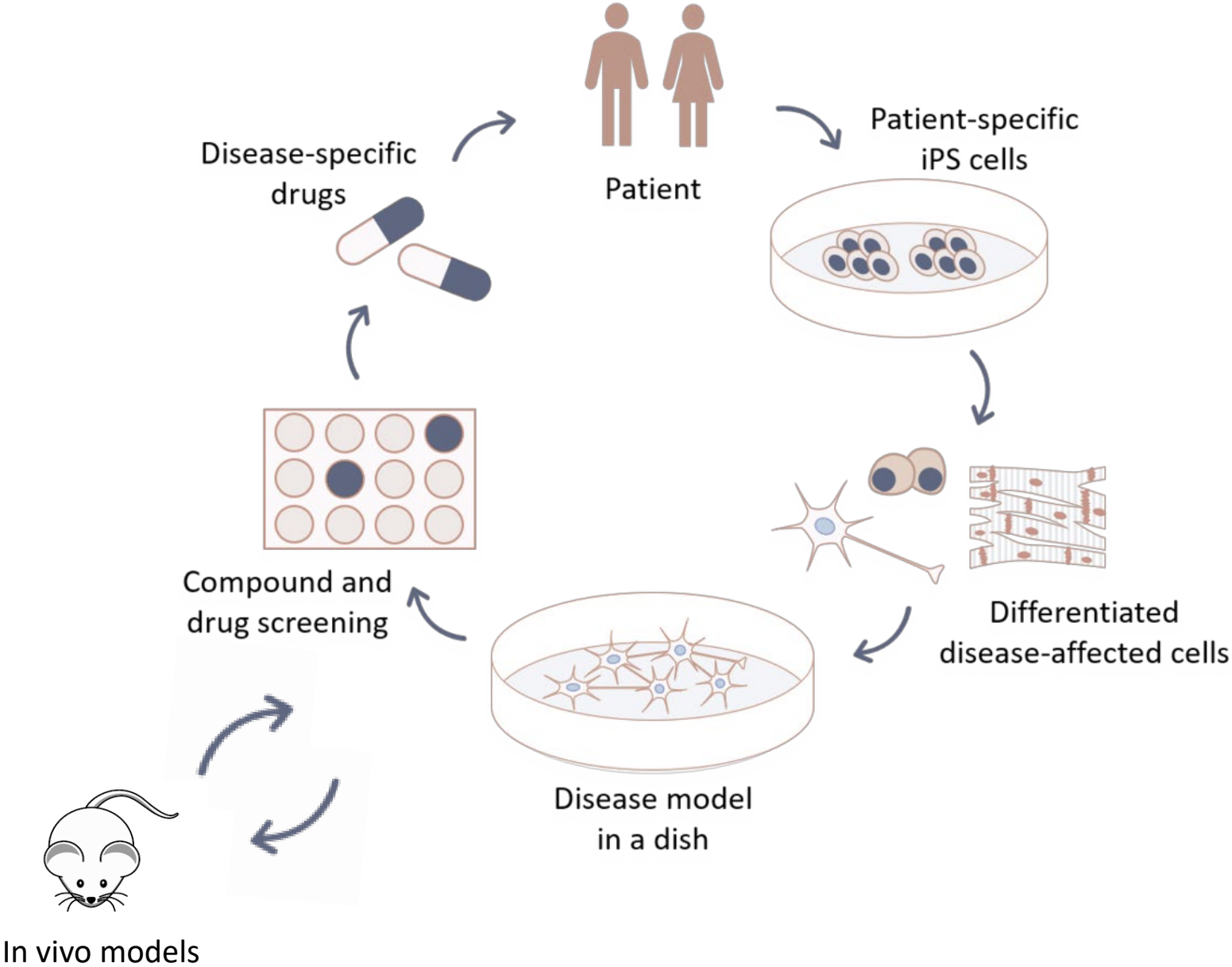
KCC2/NKCC1 expression



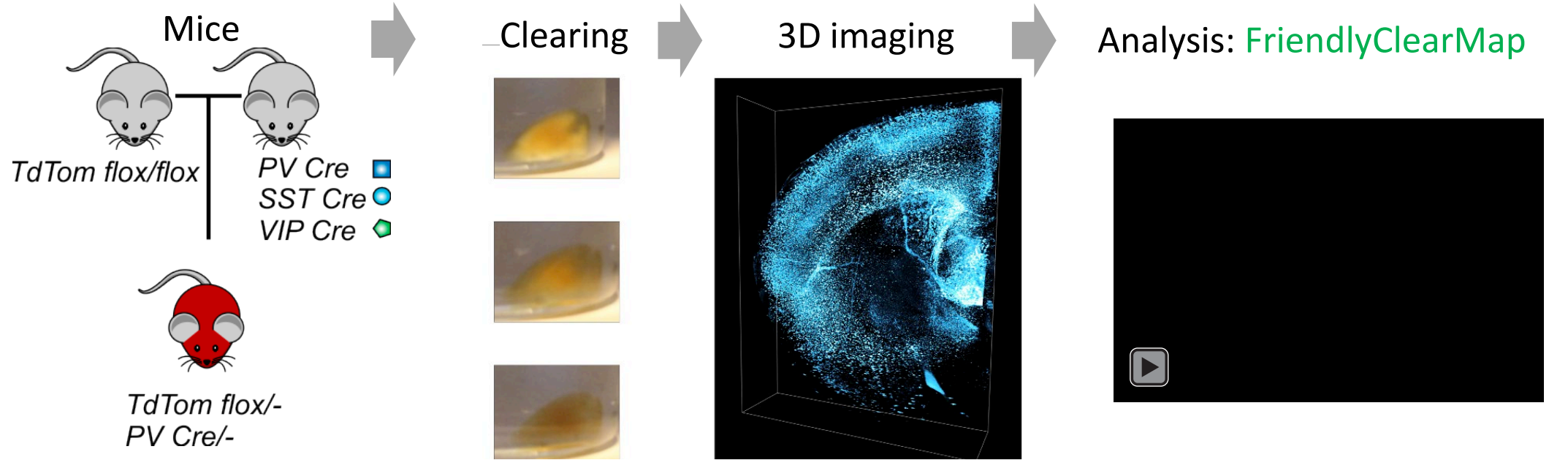
development



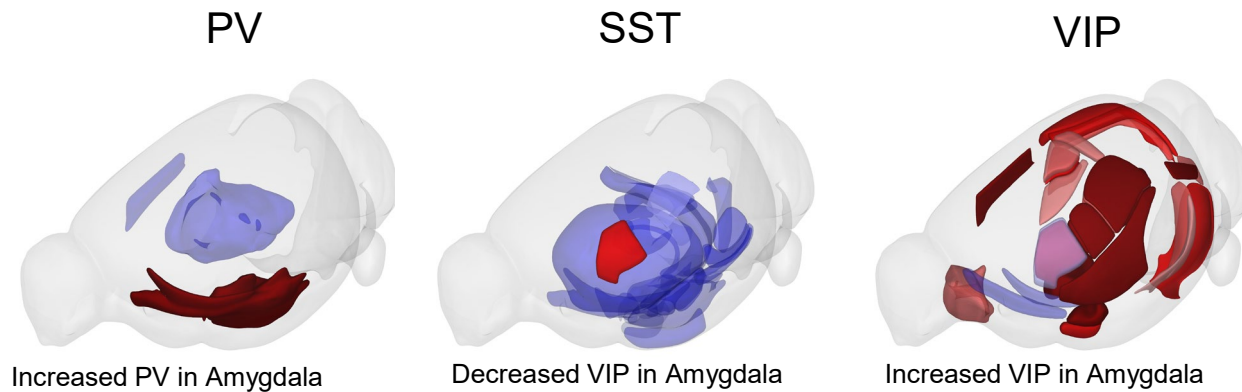
How do findings in a dish translate to *in vivo*?



Altered development of GABAergic neurons in *EHMT1*^{+/-} mice?



Negwer M et al., 2023 *GigaScience*



Relative cell density compared to *Ehmt1*^{+/+}

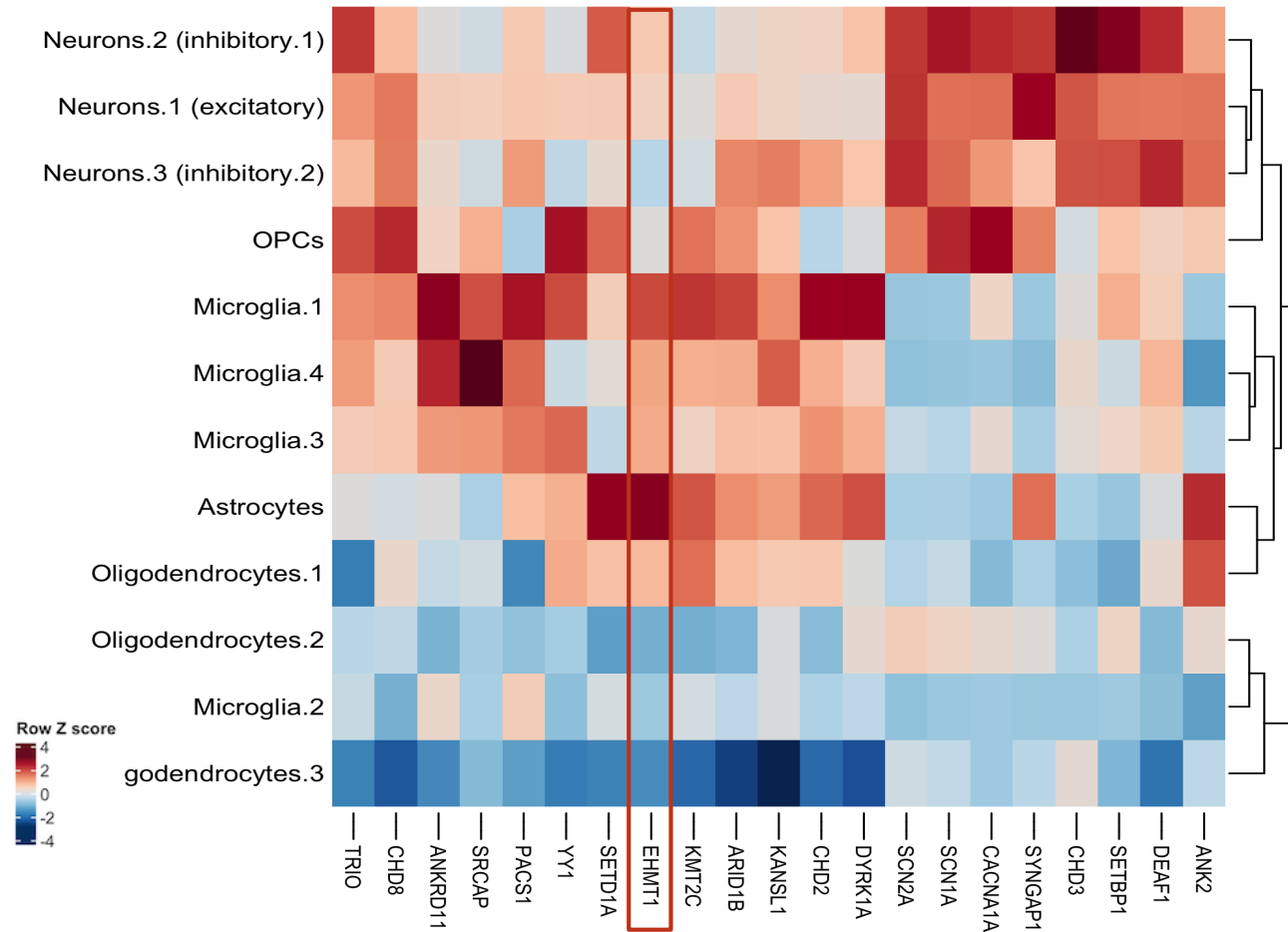


Low

High

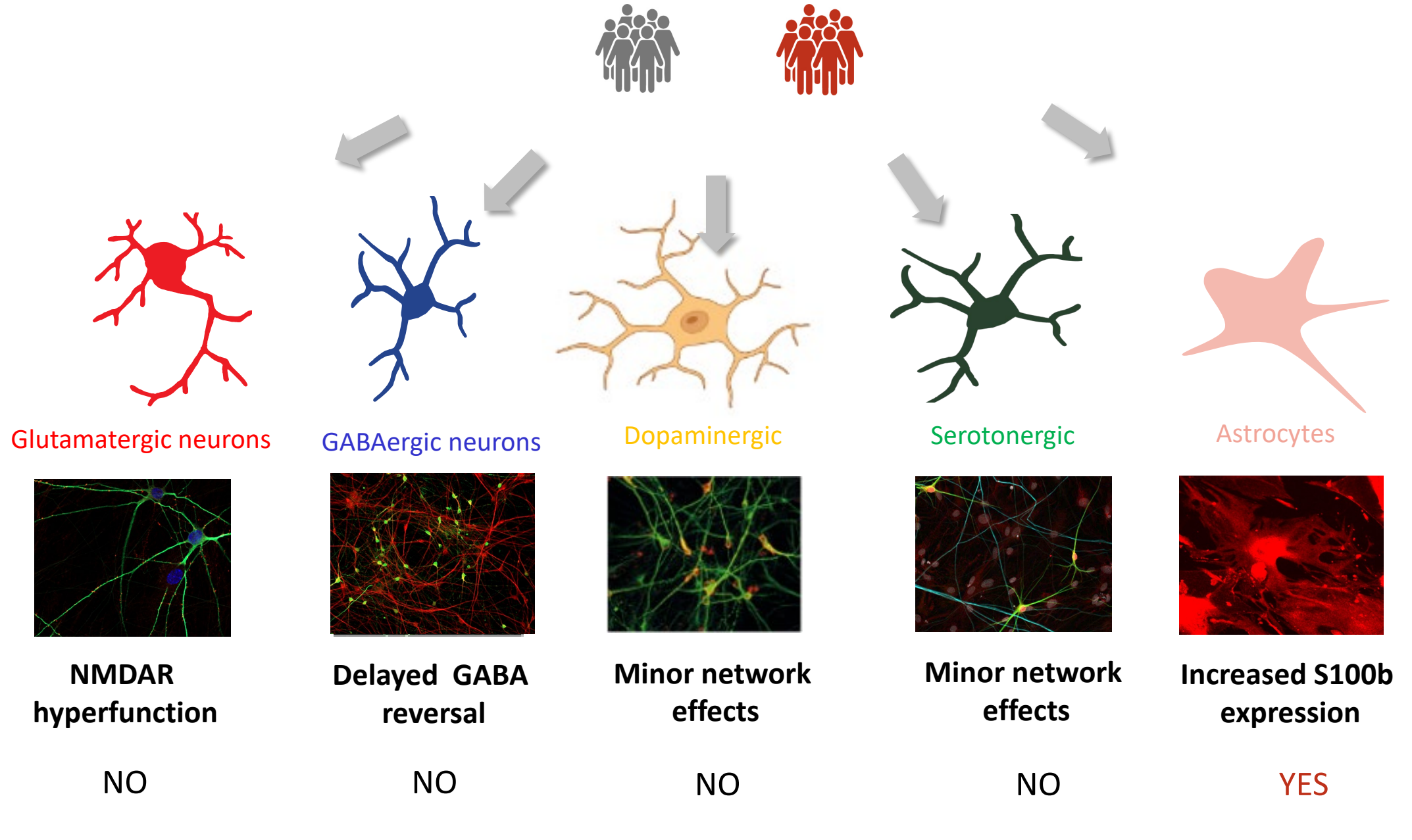
Negwer M et al, 2021, *Brain struct Funct*
Negwer M et al., *in preparation*

Expression of *EHMT1*



Single-cell data expression post-mortem material

Investigating the cell type-specific function of EHMT1 using iPSC models



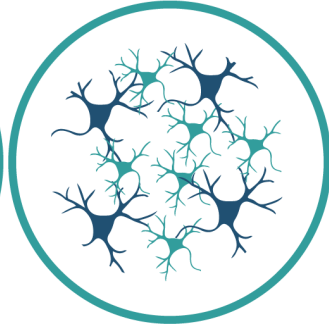


BRAINmodel

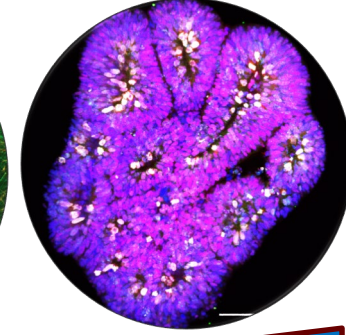
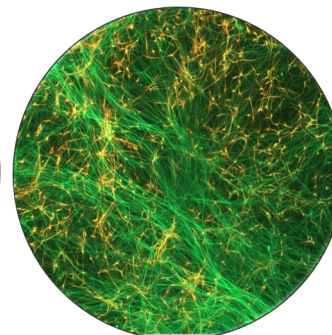
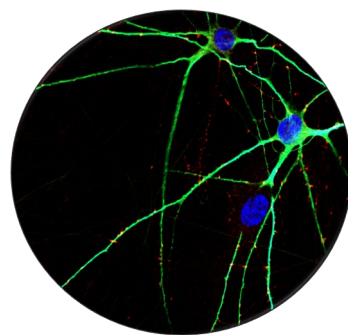
- Development of robust **stem cell-based** 2D and 3D brain models
- Increase robustness, complexity, predictability and scalability



Induced Pluripotent
Stem Cells (iPSC)



In vitro iPSC derived brain models

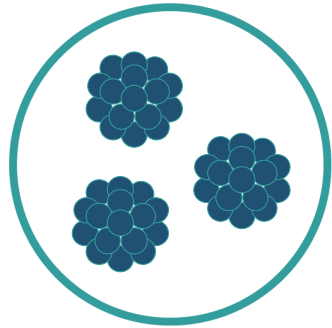


Complexity



BRAINmodel

- Development of robust **stem cell-based** 2D and 3D brain models
- Increase robustness, complexity, predictability and scalability



Induced Pluripotent Stem Cells (iPSC)

- Chromatinopathies
- Synaptopathies

H3K4Me
 KMT2A
 KMT2C
 KMT2D
 KMT2F/SETD1A
 KMT2G
 KDM1A
 KDM5C

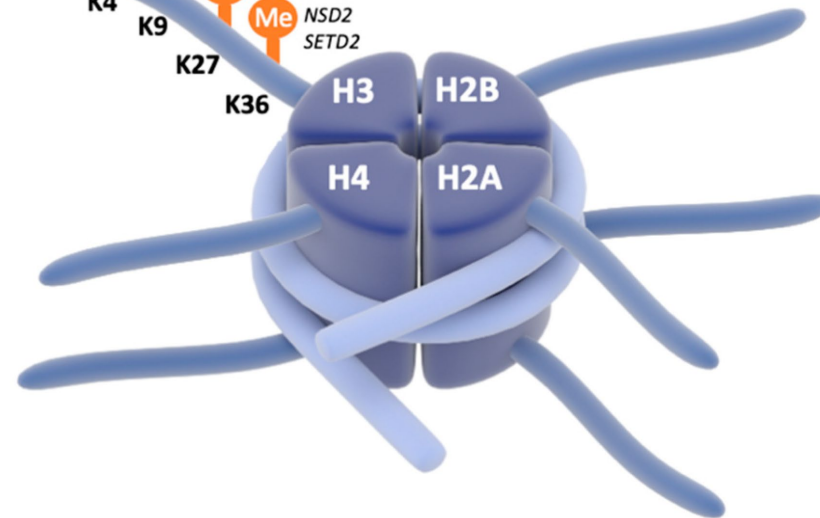
H3K9Me
 EHMT1

H3K27Me
 EZH1
 KDM6A
 KDM6B

H3K36Me
 NSD1
 NSD2
 SETD2

Me
 Me
 Me
 Me

K4
 K9
 K27
 K36



–EHMT1
 –KMT2A
 –KMT2D
 –SETD1A



BRAINmodel

- Linking **in vitro** measurements to **patient** measurements



Balance
Excitation-Inhibition



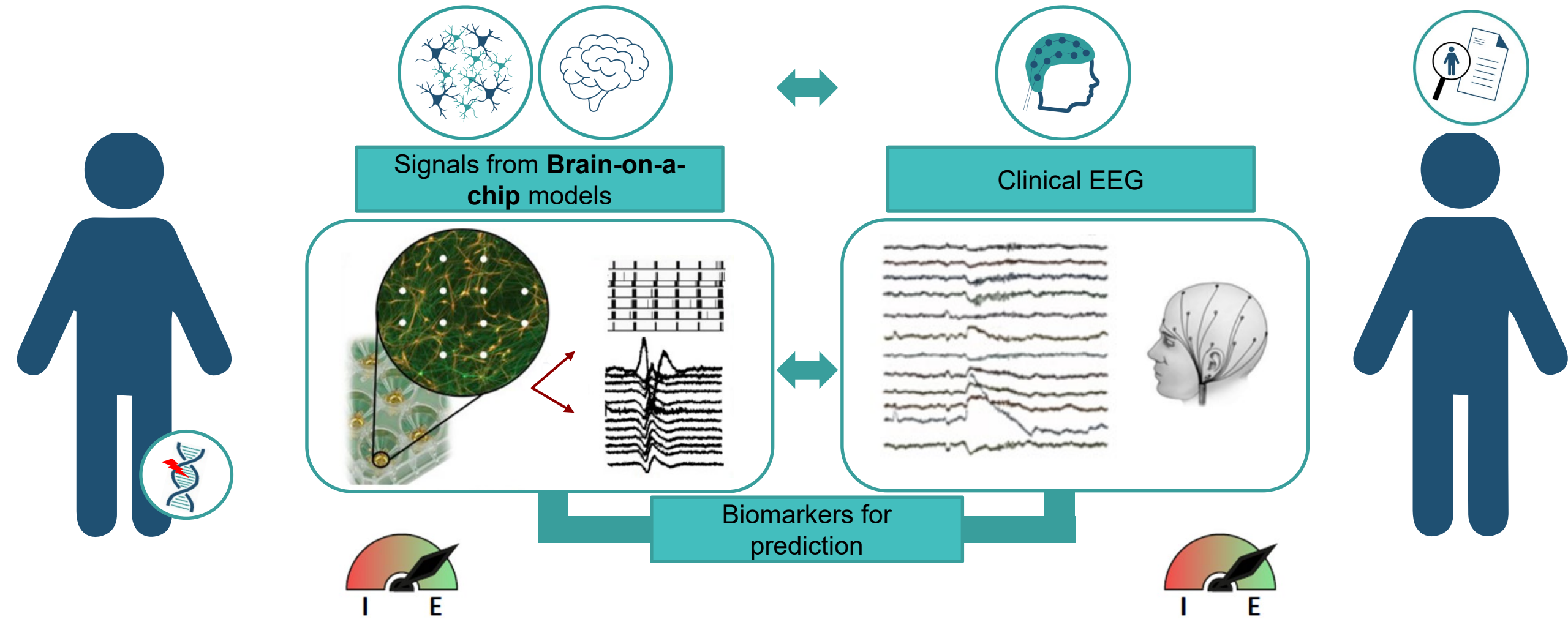
- Based on mechanisms: E/I imbalances - drug repurposing
- Epigenetic compounds to restore “balance”



BRAINmodel



<https://brainmodel.nl>





UNIVERSITY OF TWENTE.
CLINICAL NEUROPHYSIOLOGY

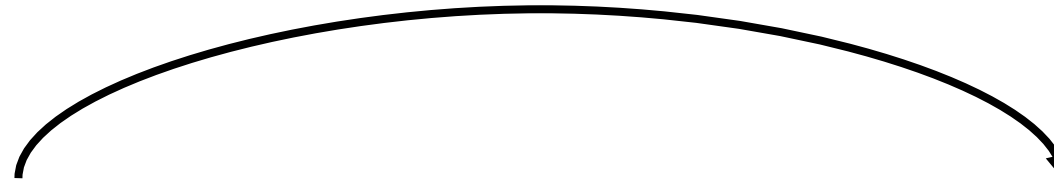
Nina Doorn

Dr. Monica Frega

Prof. dr. ir. Michel van Putten

<https://brainmodel.nl>

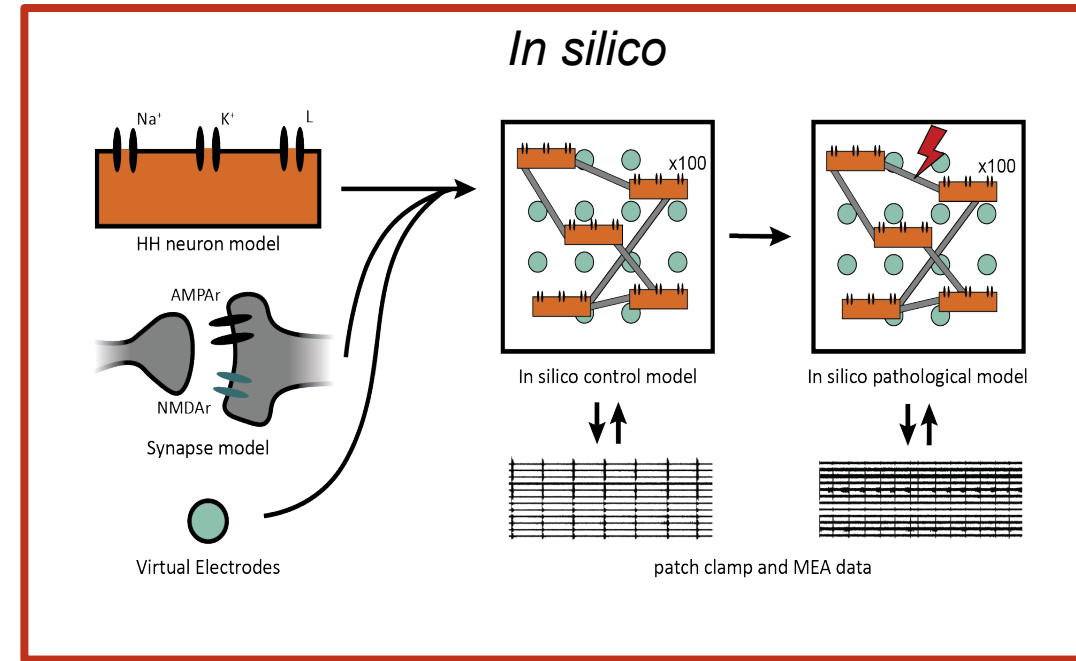
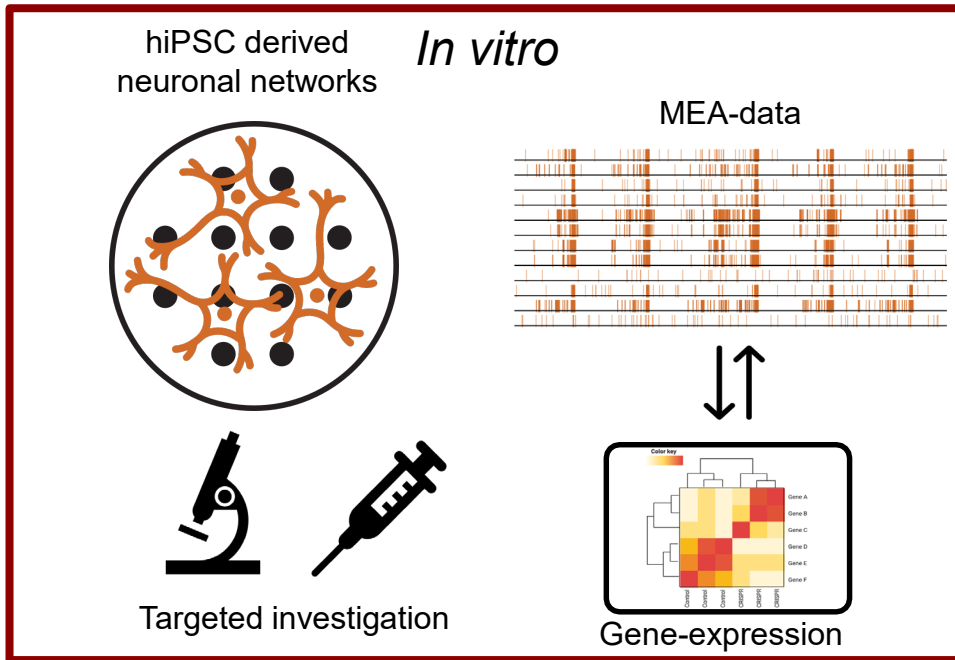
Generate model with data



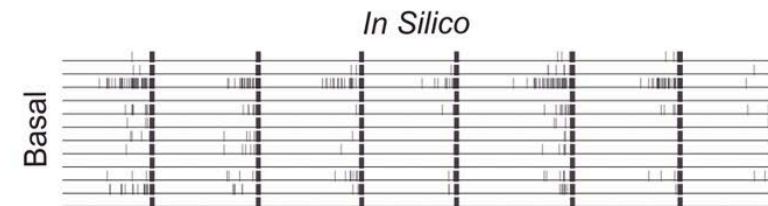
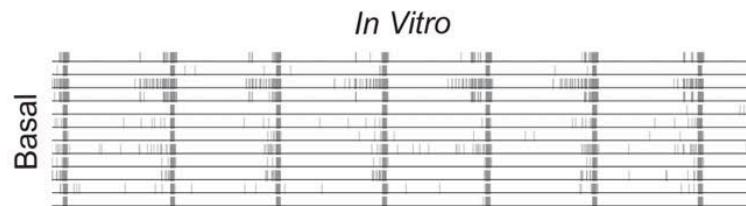
NDD patient



?



Discover potential disease mechanisms



Generate predictions about underlying cellular mechanisms

D. Schubert
H. van Bokhoven
T. Kleefstra

Shan Wang
Eline van Hugte
Katrin Linda
Chantal Schoenmakers
Astrid Oudakker
Ka Man Wu
Britt Mossink
Anouk Verboven
Elly Lewerissa
Umami Ciptasari
Rachel Mijdam
Naoki Kogo
Brooke Latour
Xiuming Yuan
Emma Dyke
Imke Schuurmans
Nicky Scheefhals
Sofia Puvogel
Kim Wijnant
Annemiek van de Steeg



Radboudumc

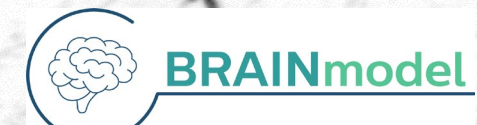
A. Bouman
B. De Vries
D. Koolen
B. Franke

U Twente

M. Frega
M. Van Putten

VU Amsterdam

M. Verhage
N. Cornelissen



Summary

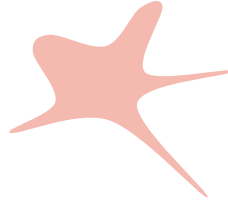
Neuronal Network



Hyperactivity

Reduced network organization

Cellular



Decreased glutamate uptake
Altered Ca^{2+} signalling
Neuroinflammation

Molecular

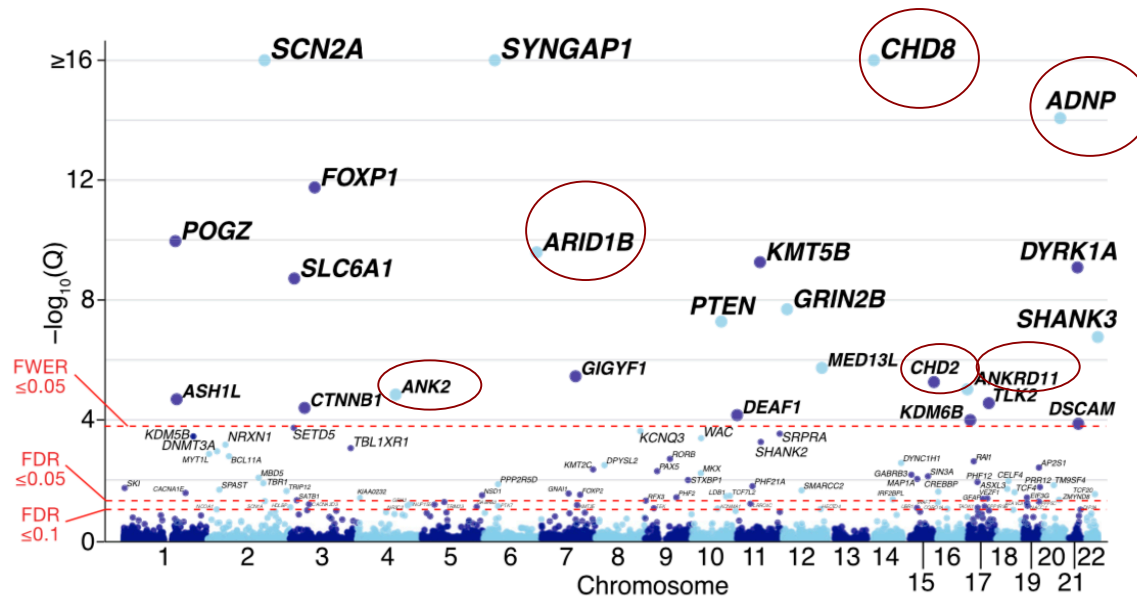


Increased *S100B* expression and decreased H3K9me2 at *S100B* loci

Rescue

Rescue with *S100B* antagonist **olanzapine**

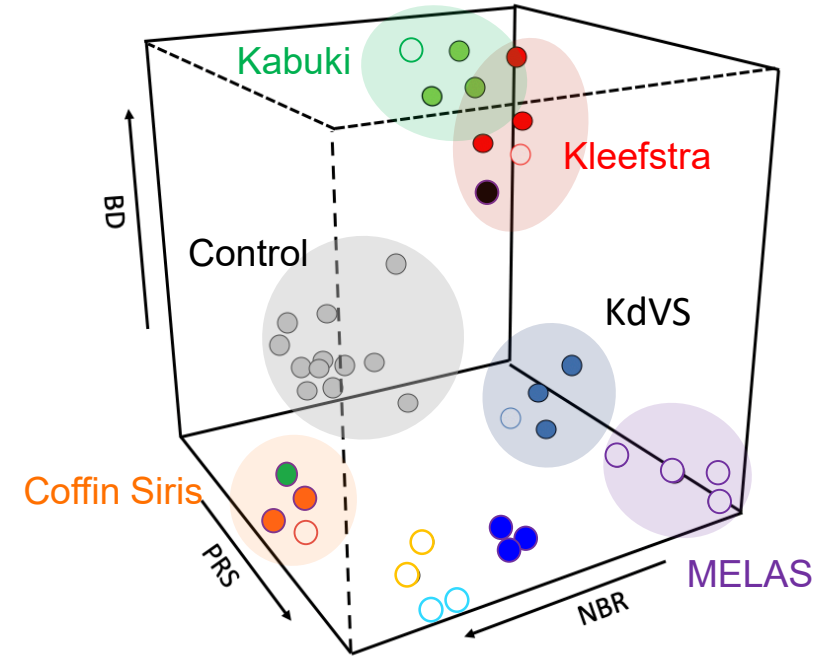
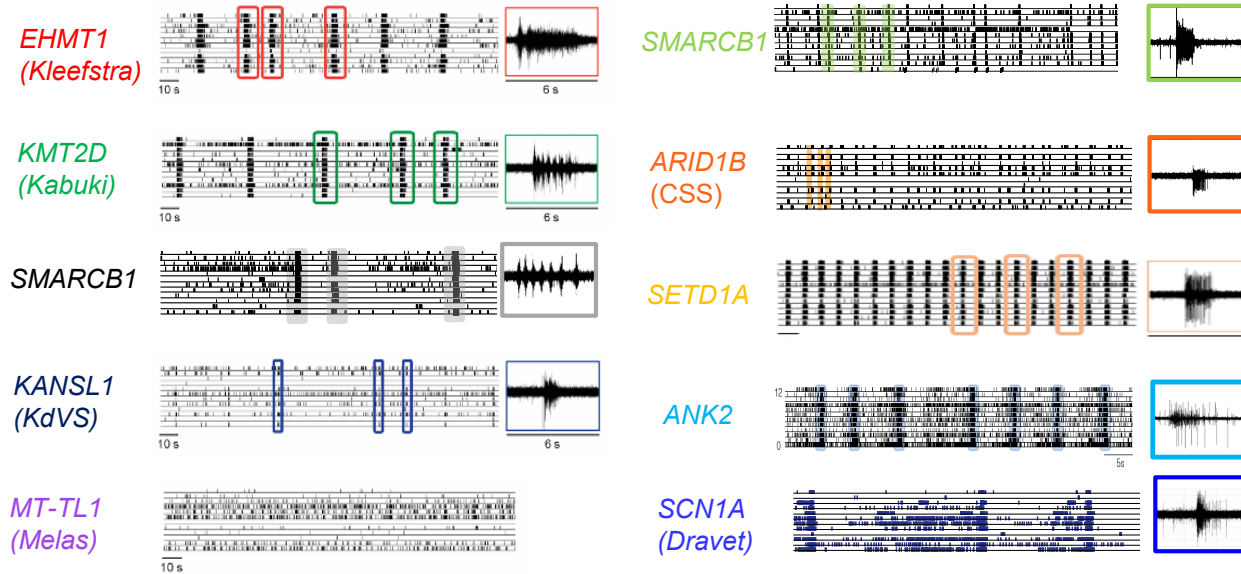
Can we identify disease/patient-specific neuronal network phenotypes?



Satterstrom et al., 2020 Cell

- | | |
|----------------|----------------|
| EHMT1 | ANK2 |
| KMT2C | SCN1A |
| SMARCB1 | CACNA1A |
| MBD5 | MELAS |
| ARID1A | NANS |
| ARID1B | PMM2 |
| SETD1A | CDH13 |
| KANSL1 | PCDH19 |
| CHD2 | ADNP |
| CHD3 | ANKRD11 |
| CHD8 | YY1 |

Syndrome-specific neuronal network phenotypes

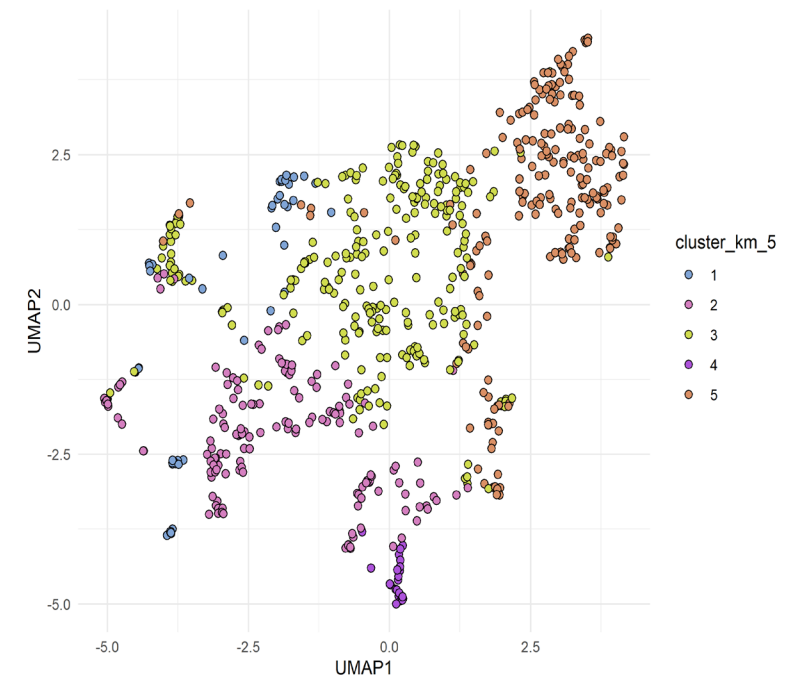
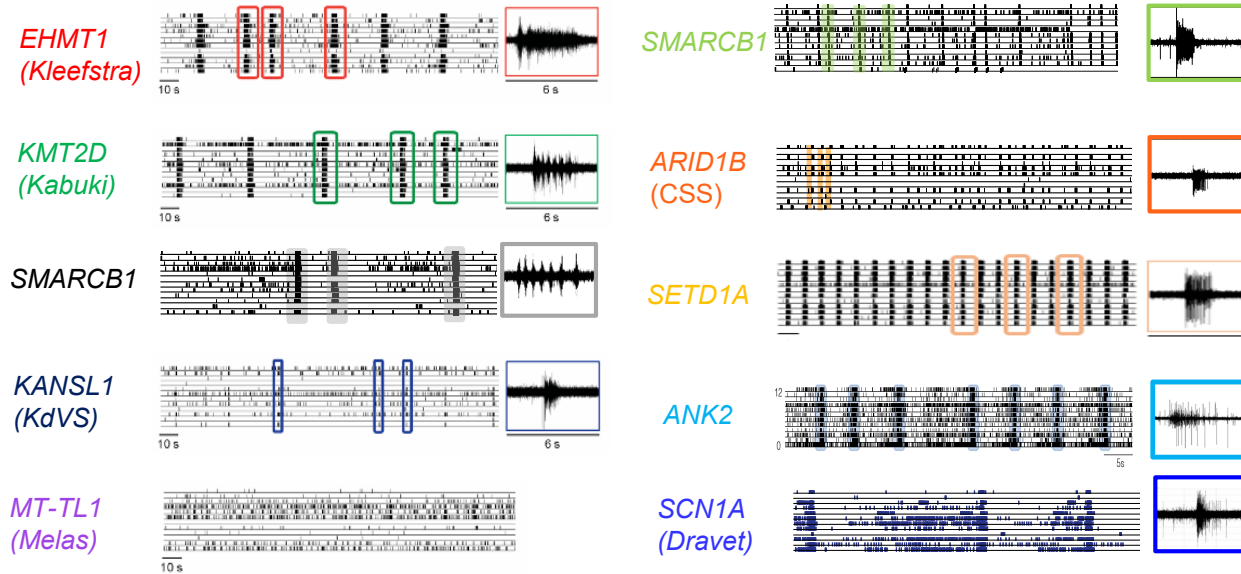


● Closed circles show independent iPSC lines
○ Open circles show CRISPR/Cas9 isogenic lines

Frega et al., Nat Comms 2019
Klein Gunnewiek et al., Cell Rep 2020
Frega et al., Cell Rep 2020
Frega et al., Cell Rep 2020
Mossink et al., Stem Cell Rep 2021
Mossink et al., Mol Psych 2021
Linda et al., Autophagy 2021
Shan et al., Cell Rep 2022
Gerosa et al., Cell Rep 2022

- 15 syndromes
- 1-5 cell lines per disease
- 39 cell lines
- >1000 recordings

Syndrome-specific neuronal network phenotypes



Frega et al., Nat Comms 2019
Klein Gunnewiek et al., Cell Rep 2020
Frega et al., Cell Rep 2020
Frega et al., Cell Rep 2020
Mossink et al., Stem Cell Rep 2021
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