

# An *in vitro* and *in silico* neuronal network model to unravel genetic encephalopathies

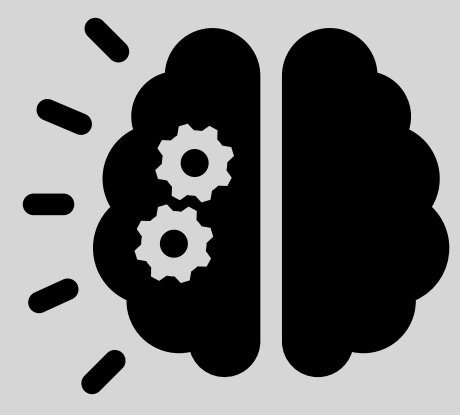
Contact



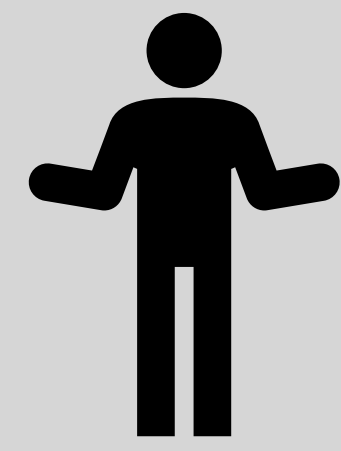
*In vitro* article



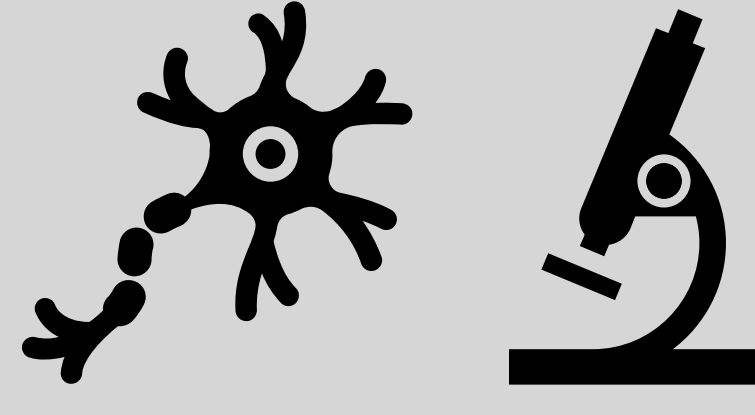
## Introduction



Neurological disorders are poorly understood



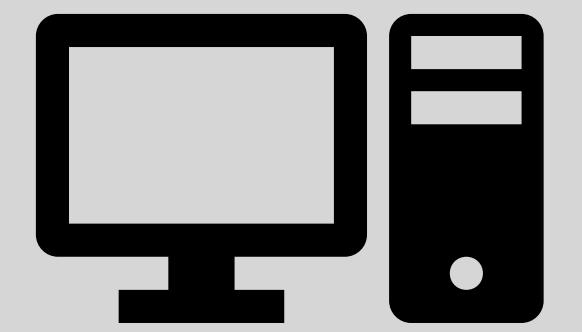
Limited experimental freedom and insight *in vivo*



Approach: stem-cell derived neuronal networks *in vitro*

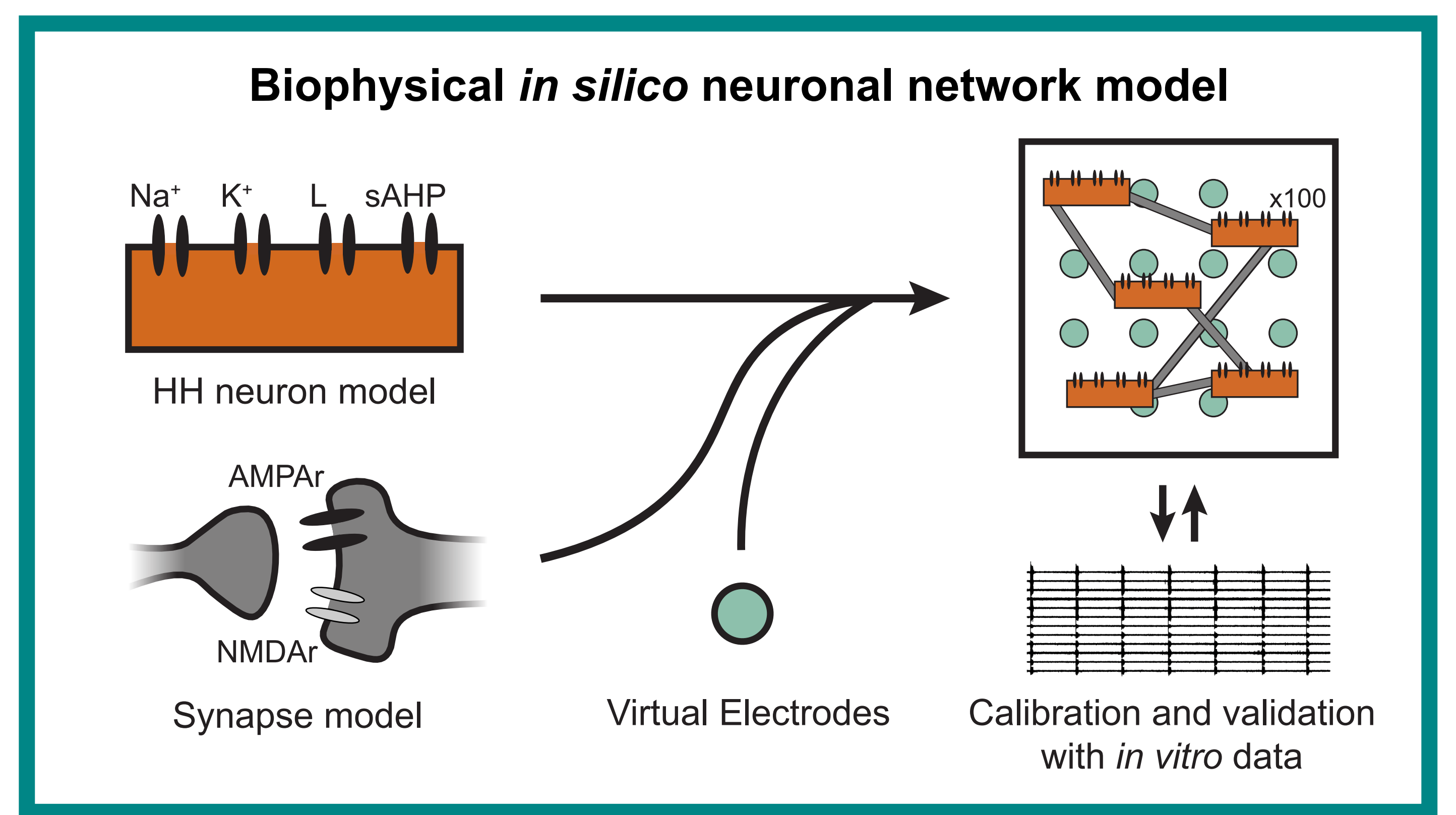
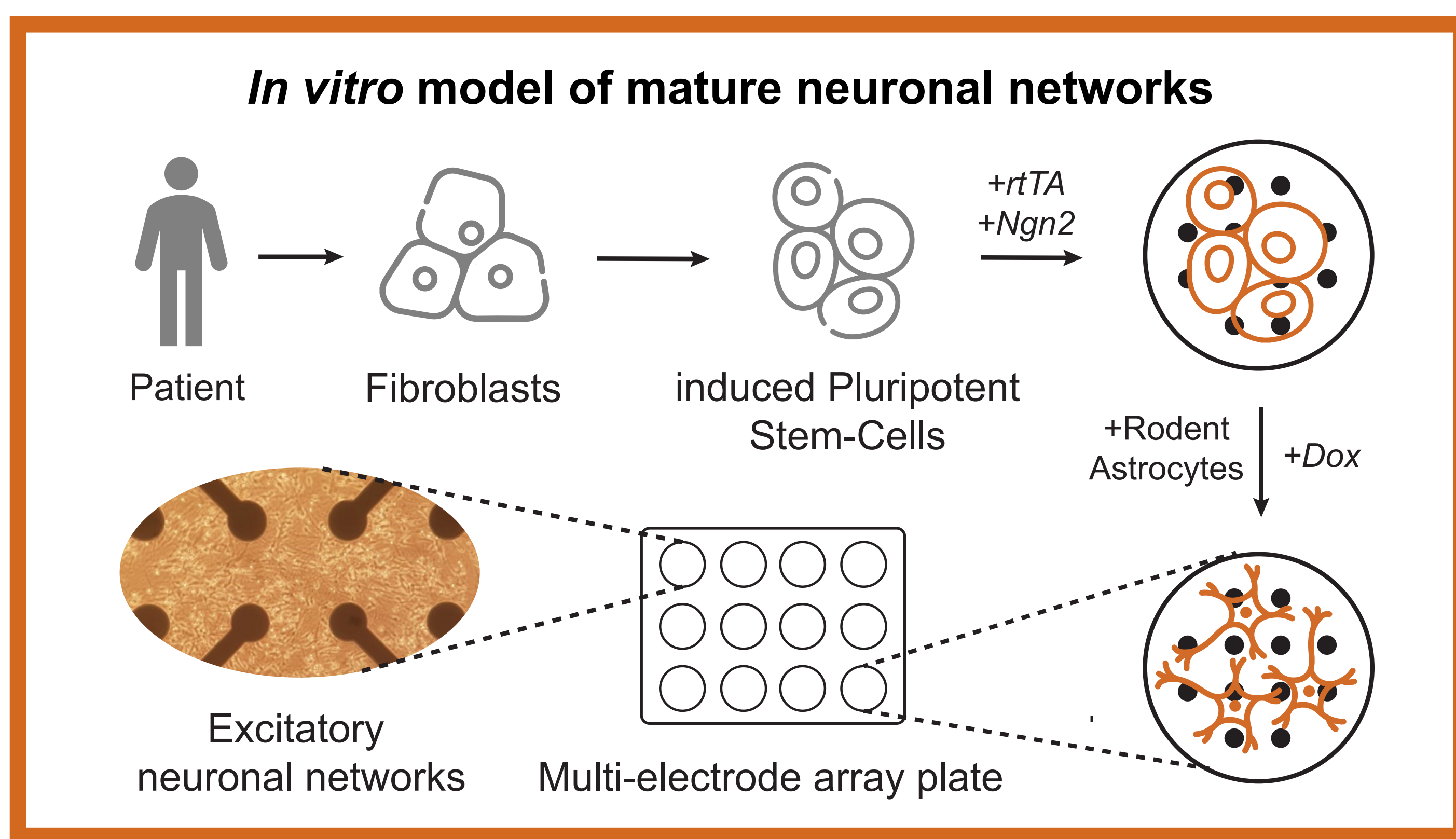


Difficult to deduce disease mechanisms from data



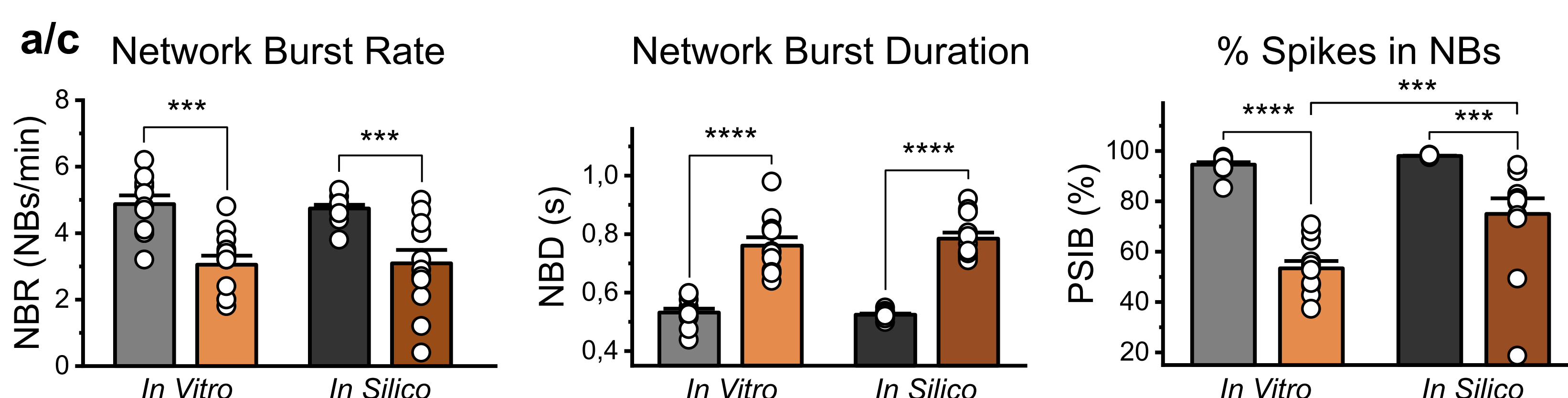
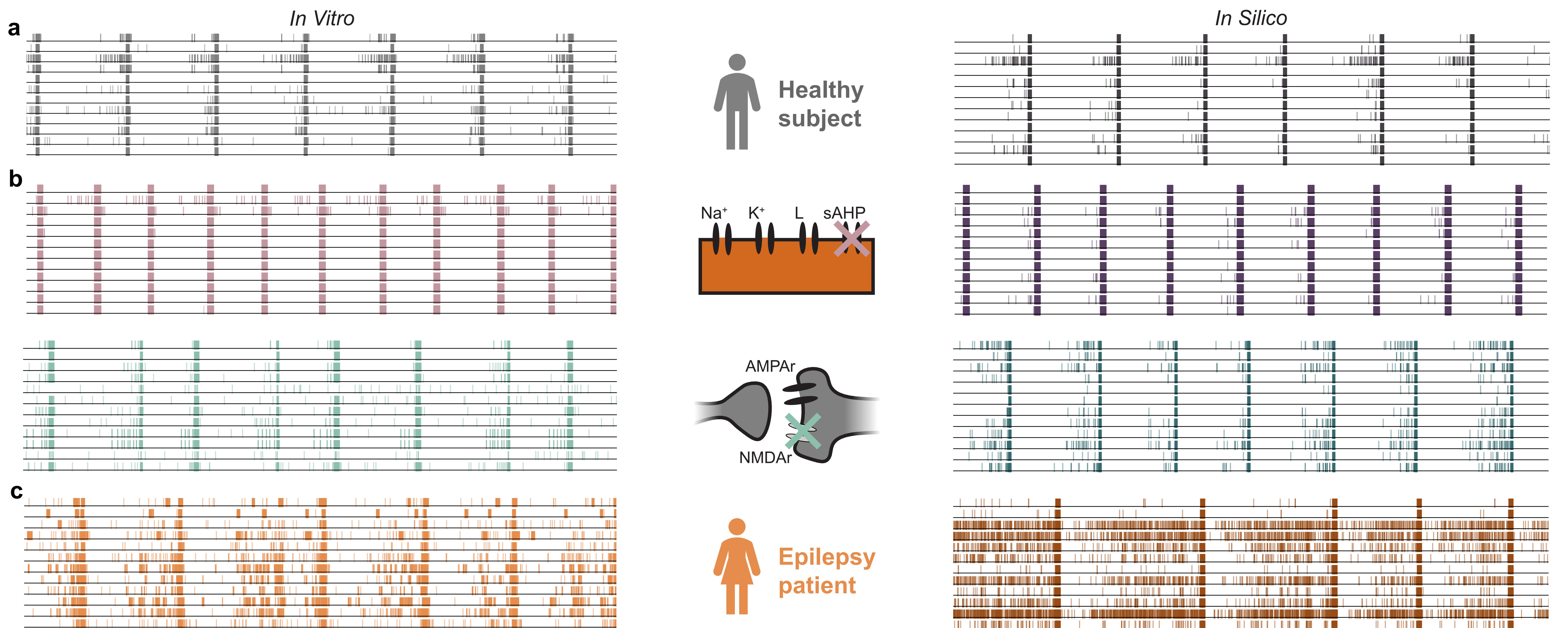
Solution: *In silico* models to leverage the data

## Methods



## Results

Exemplary rasterplots of *in vitro* measurements & *in silico* simulations in a) physiological, b) pharmacological and c) pathological conditions



In order to model the epilepsy patients' networks activity, we had to lower the **sAHP current** and **synaptic strengths**. We subsequently found evidence for these changes in the *in vitro* networks, highlighting the power of the model to **identify disease mechanisms** in patient derived networks.

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