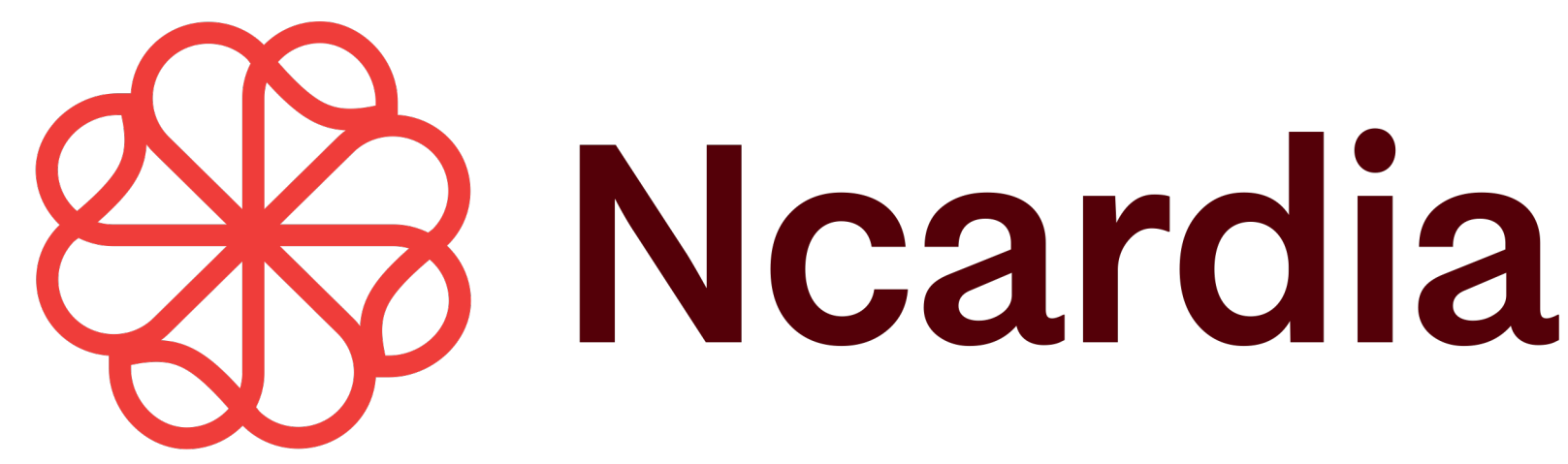


Development of iPSC-based Parkinson’s disease model for drug discovery

K. Pitsa, L. Smit, C. Van Berkel, B. Samson-Couterie, **S. Jain**

Ncardia Services BV Leiden, The Netherlands, support@ncardia.com, www.ncardia.com



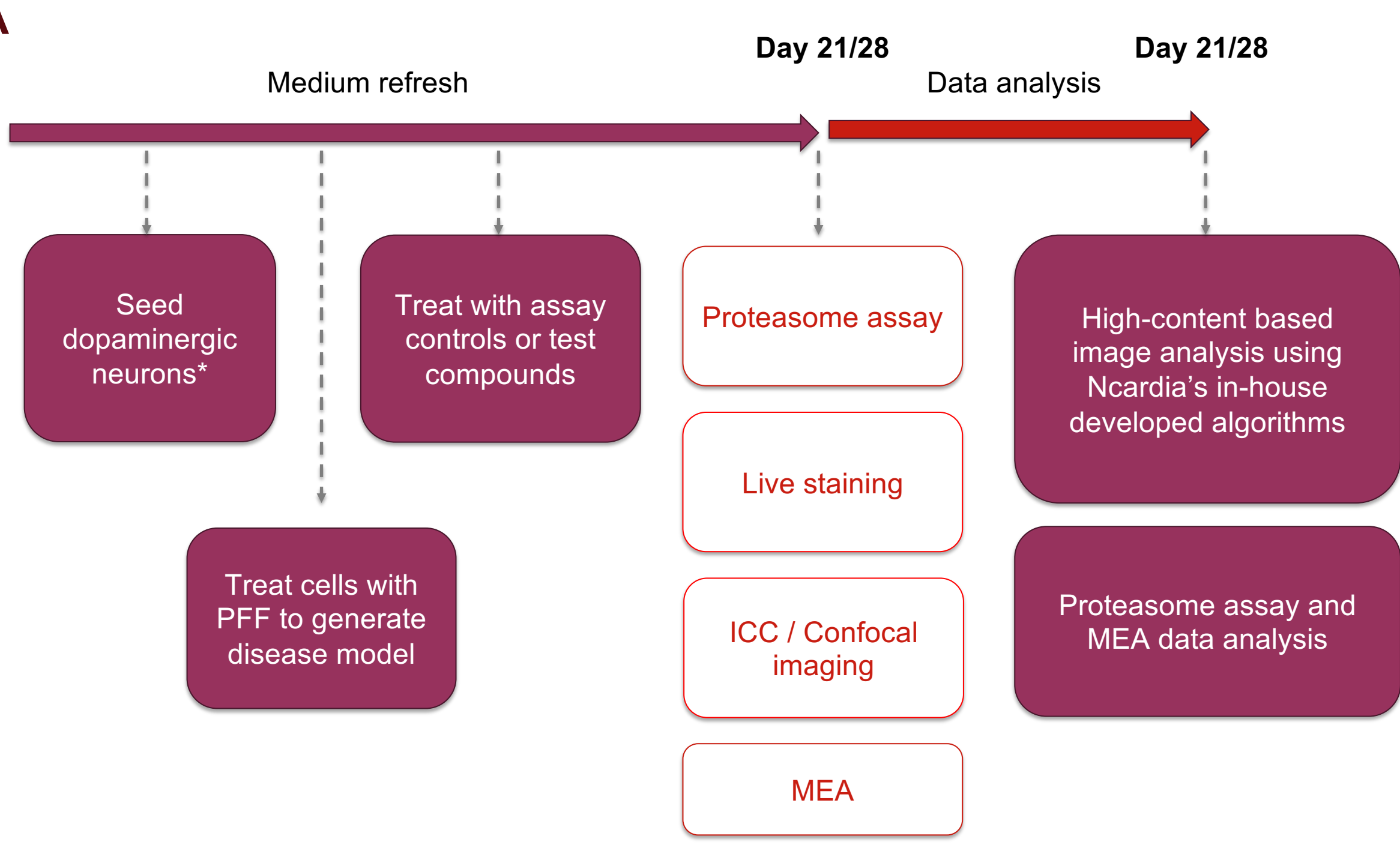
Background

Parkinson’s disease (PD) is the second most common neurodegenerative disorder, with a prevalence that has more than doubled over the past decades and is expected to double again in the coming years. Despite extensive research, the precise causes of PD remain incompletely understood, and a definitive, disease-modifying treatment is still lacking. One of the key challenges lies in the use of suboptimal preclinical models, which fail to accurately reproduce the wide range of pathological phenotypes observed in PD patients.

Induced pluripotent stem cells (iPSCs), particularly patient-specific iPSCs, represent a powerful and physiologically relevant system to overcome many of these limitations.

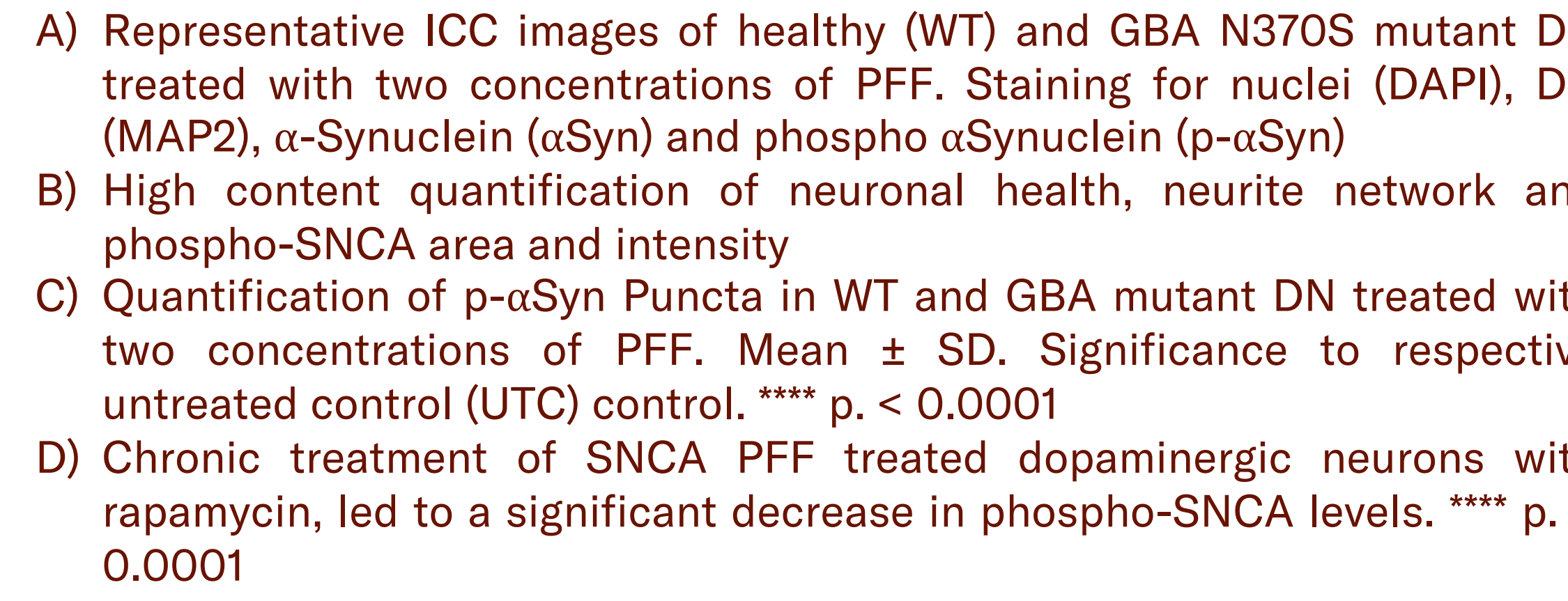
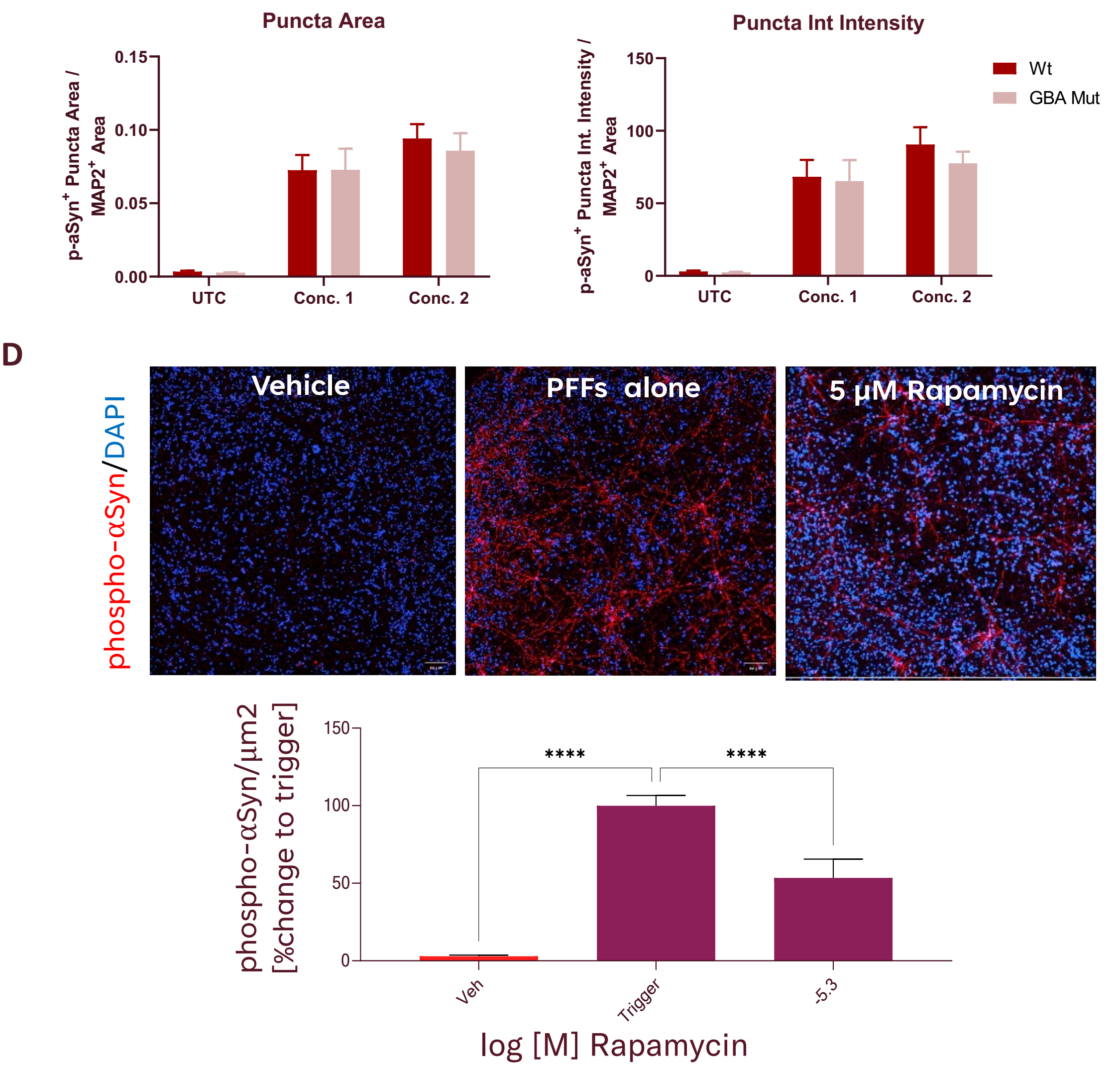
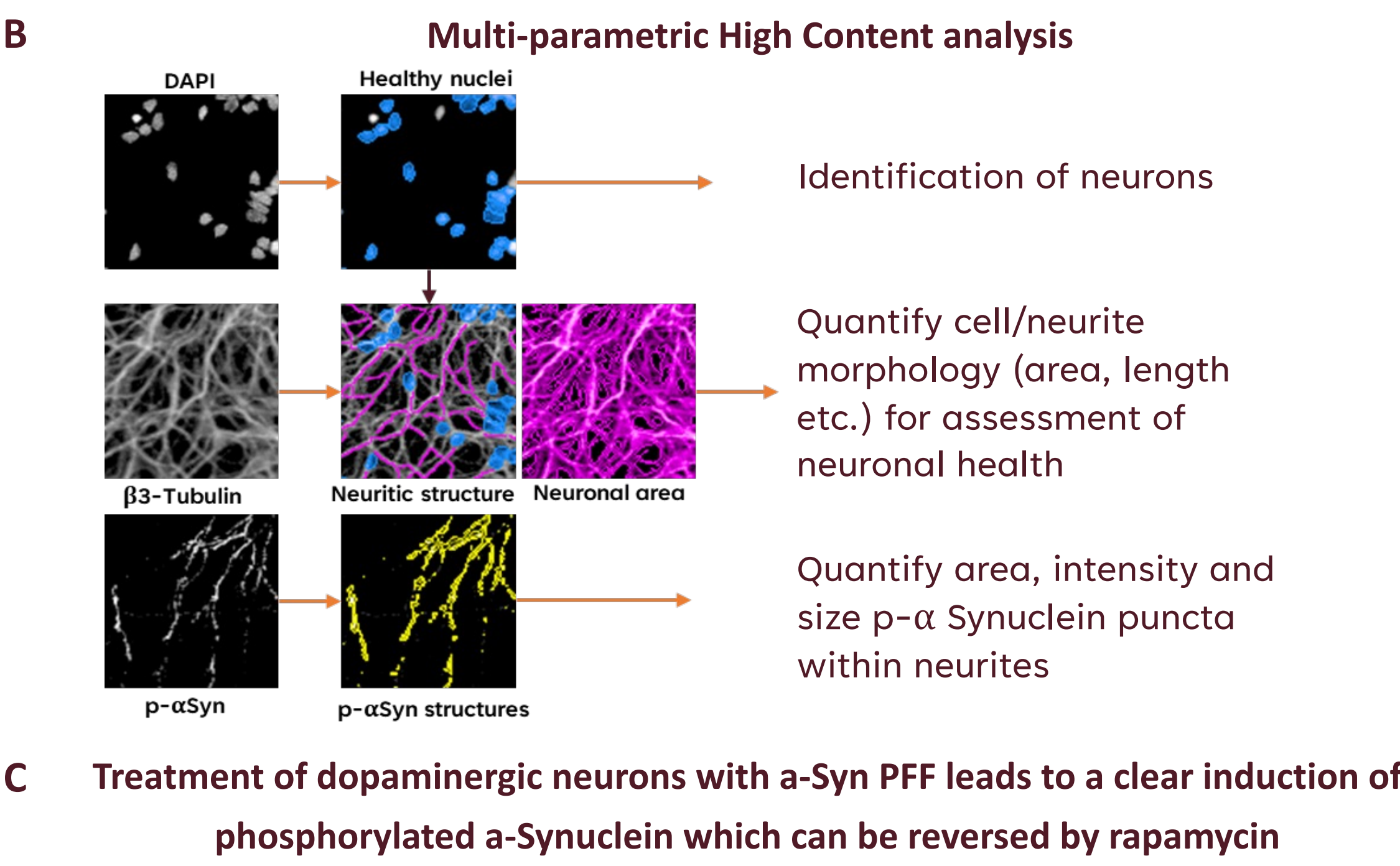
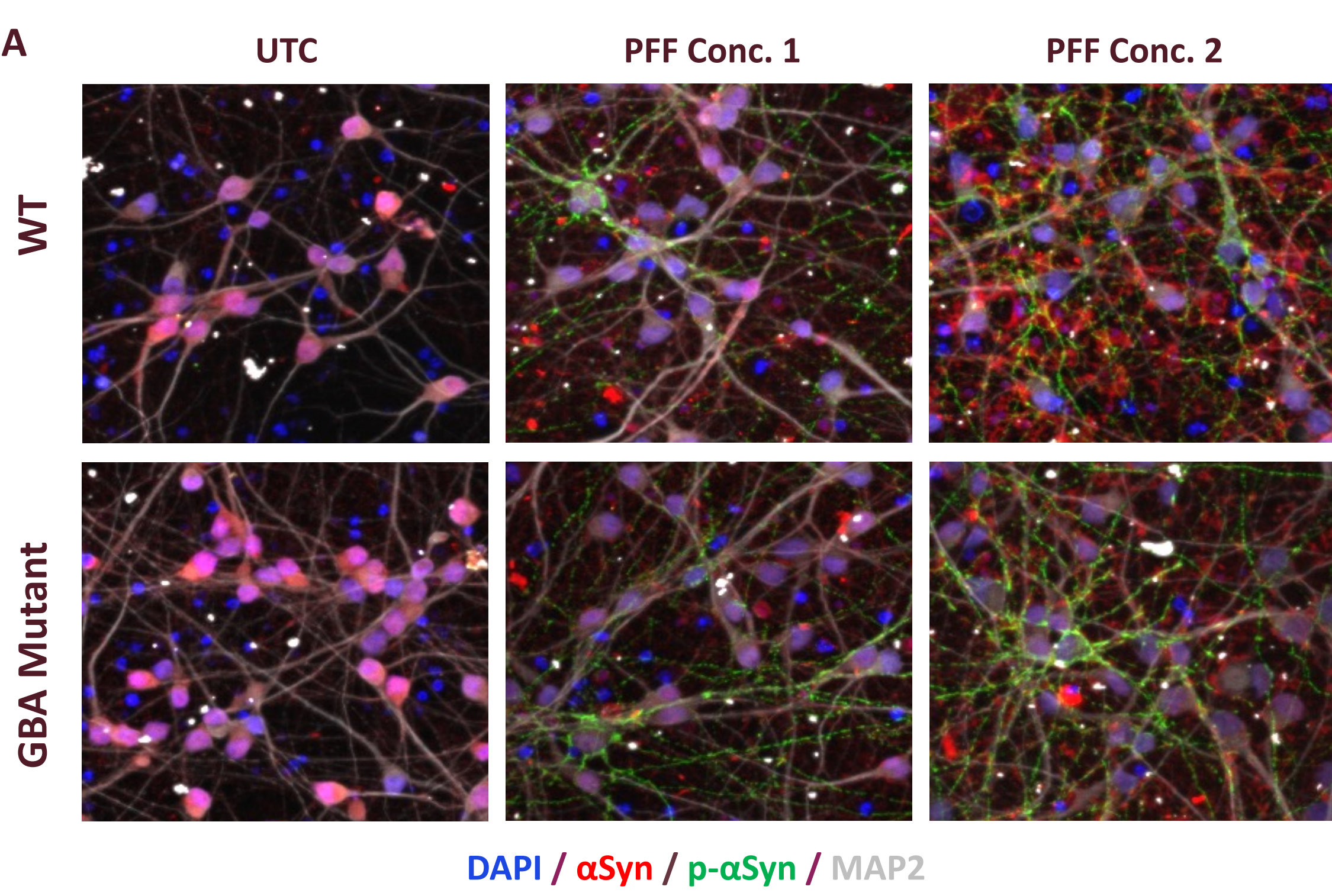
At Ncardia, iPSC-derived dopaminergic neurons (iPSC-DNs) were treated with α -synuclein (SNCA) recombinant pre-formed fibrils (PFFs) to model disease-relevant PD phenotypes. Using high-content imaging, several quantitative readouts were analyzed, including phosphorylated alpha-synuclein (pS129 α -Syn) area, number of puncta, and intensity. Furthermore, Ncardia investigated the impact of α -Syn accumulation on autophagy dynamics, mitochondrial density and functionality, as well as proteasomal activity and neuronal network activity using microelectrode array (MEA) recordings.

Treatment of iPSC-DNs with PFFs resulted in a strong and reproducible increase in phosphorylated α -Syn, accompanied by disrupted autophagy, reduced mitochondrial density and function, and significantly decreased proteasomal activity. Together, these findings provide a comprehensive in vitro model that recapitulates key hallmarks of PD pathology and enables the assessment of potential therapeutic interventions.

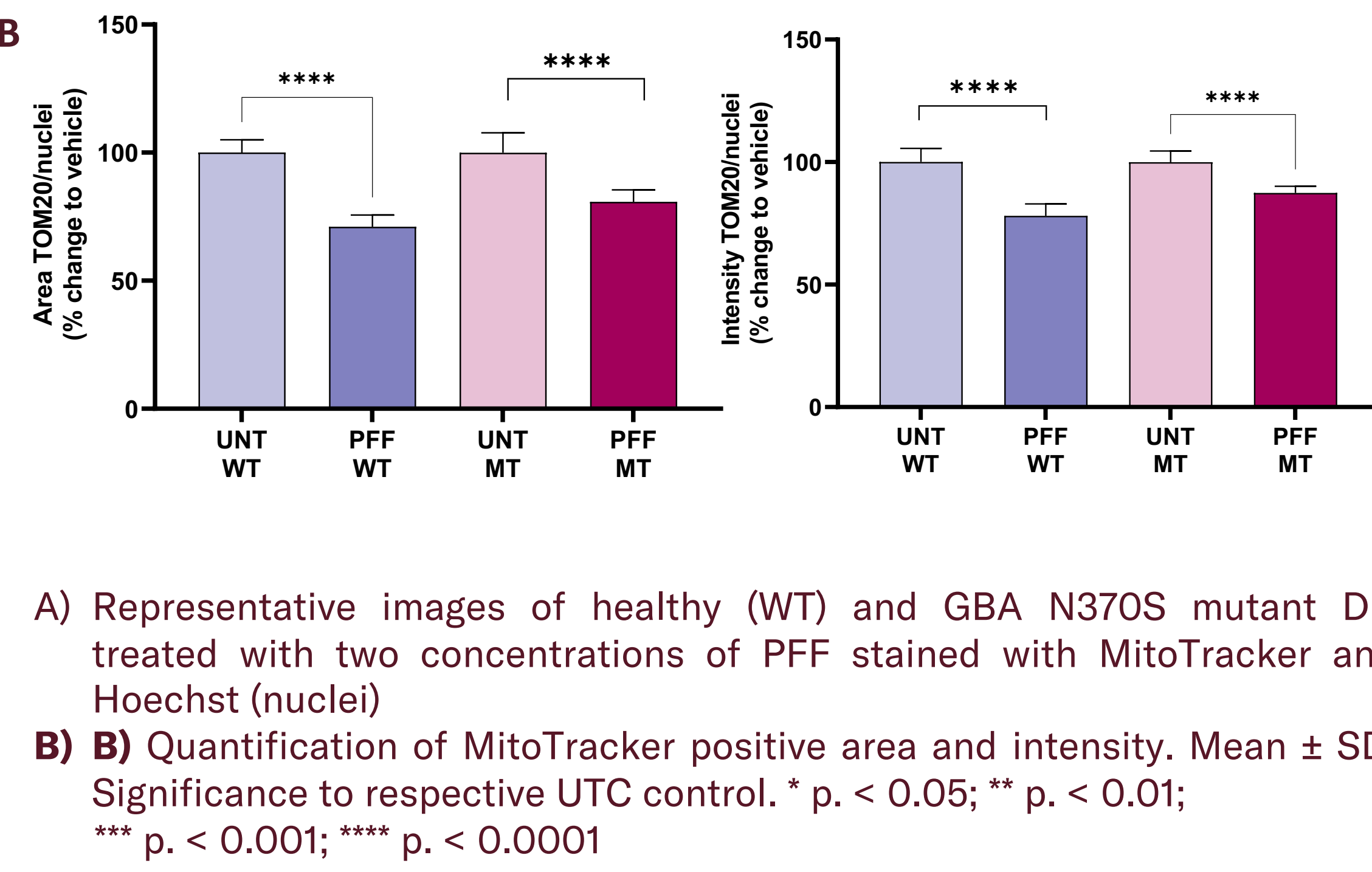
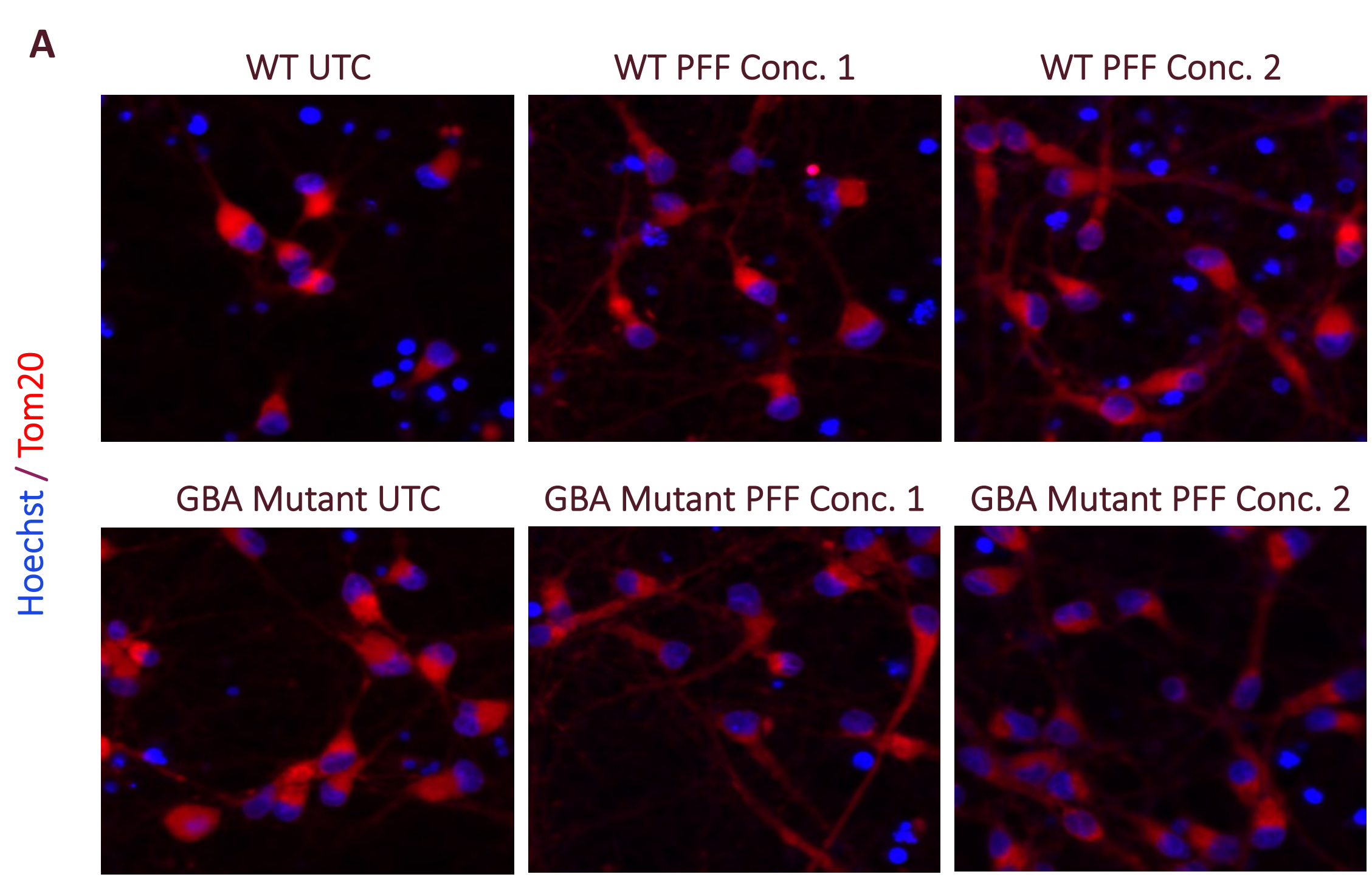


Schematic of the PD model using iPSCs derived neurons, astrocytes treated with SNCA PFFs.
* iCell® DopaNeurons 01279 (WT) and GBA N370S 11344 (GBA Mutant) from FUJIFILM Cellular Dynamics, Inc.

1. Induction of phospho-SNCA pathology

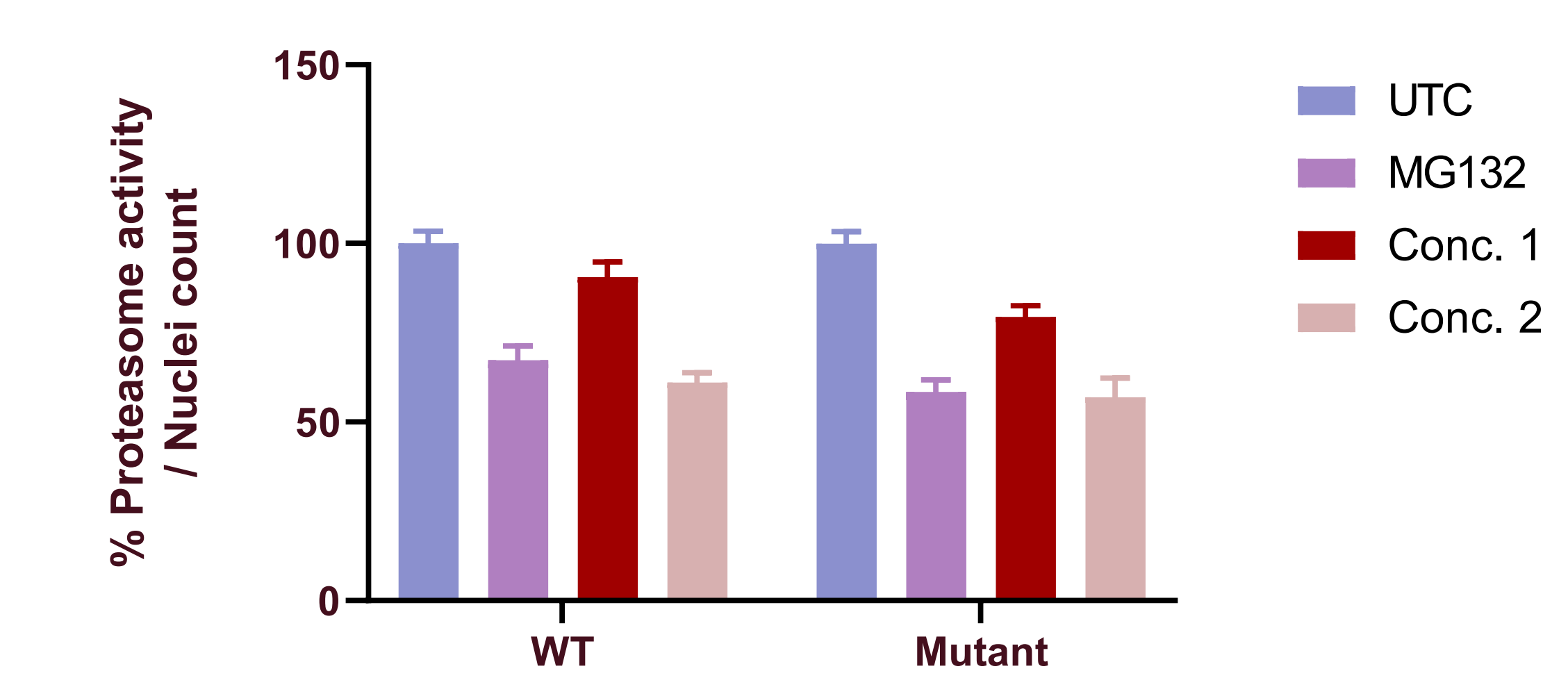


3. Mitochondria staining with phospho-SNCA pathology



Treatment of dopaminergic neurons with α -Syn PFFs leads to a reduction of mitochondria density

4. Proteasome activity in dopaminergic with phospho-SNCA pathology

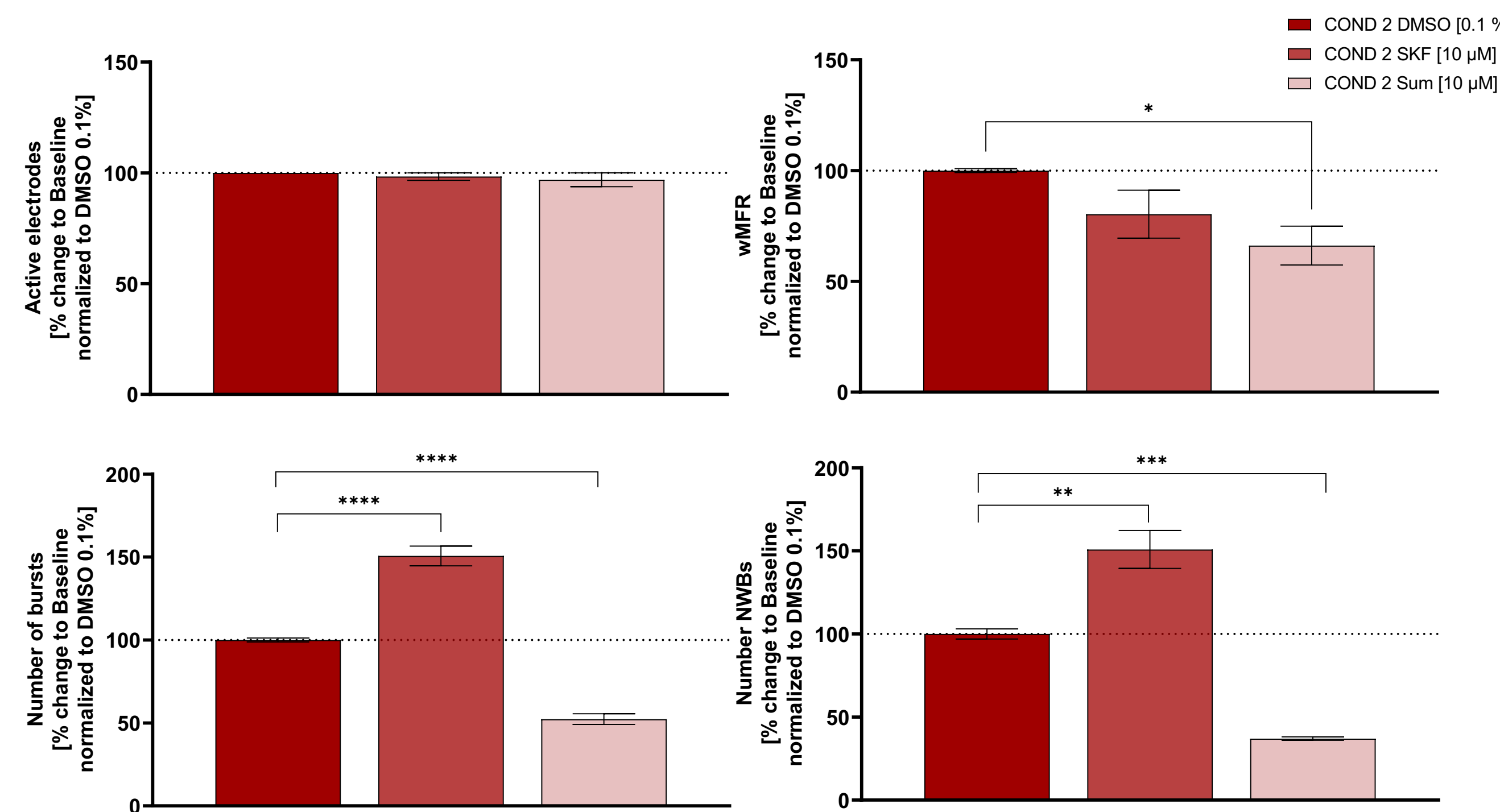
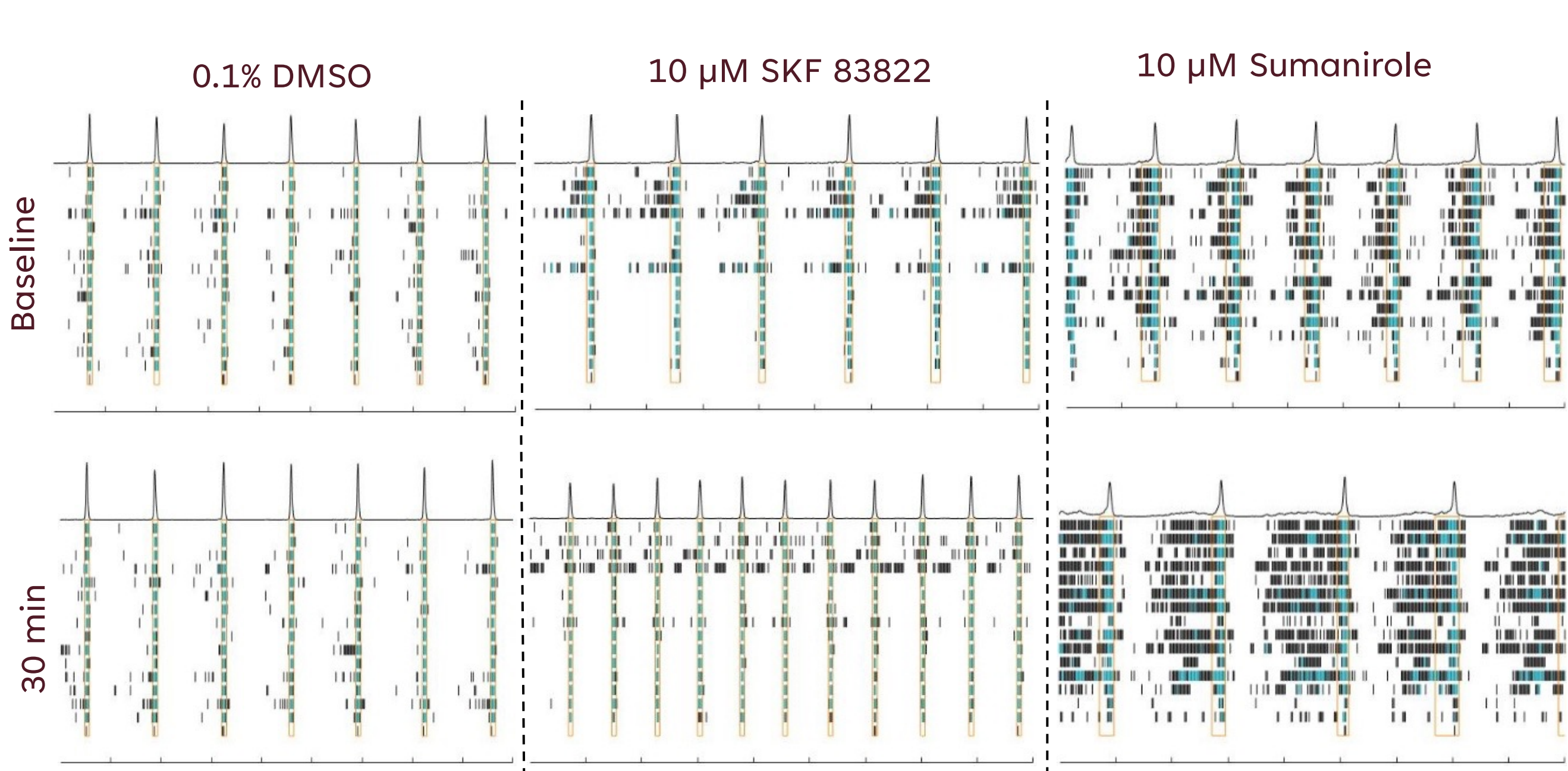


Proteasome activity assay

Quantification of Proteasome activity in WT and GBA mutant dopaminergic neurons treated with two concentrations of PFF or MG132 (Assay Ctrl), normalized on nuclei count. Mean $\pm$ SD. Significance to respective UTC. * p. < 0.05; **** p. < 0.0001

Treatment of dopaminergic neurons with α -Syn PFF leads to a 50-40% reduction in proteasome activity

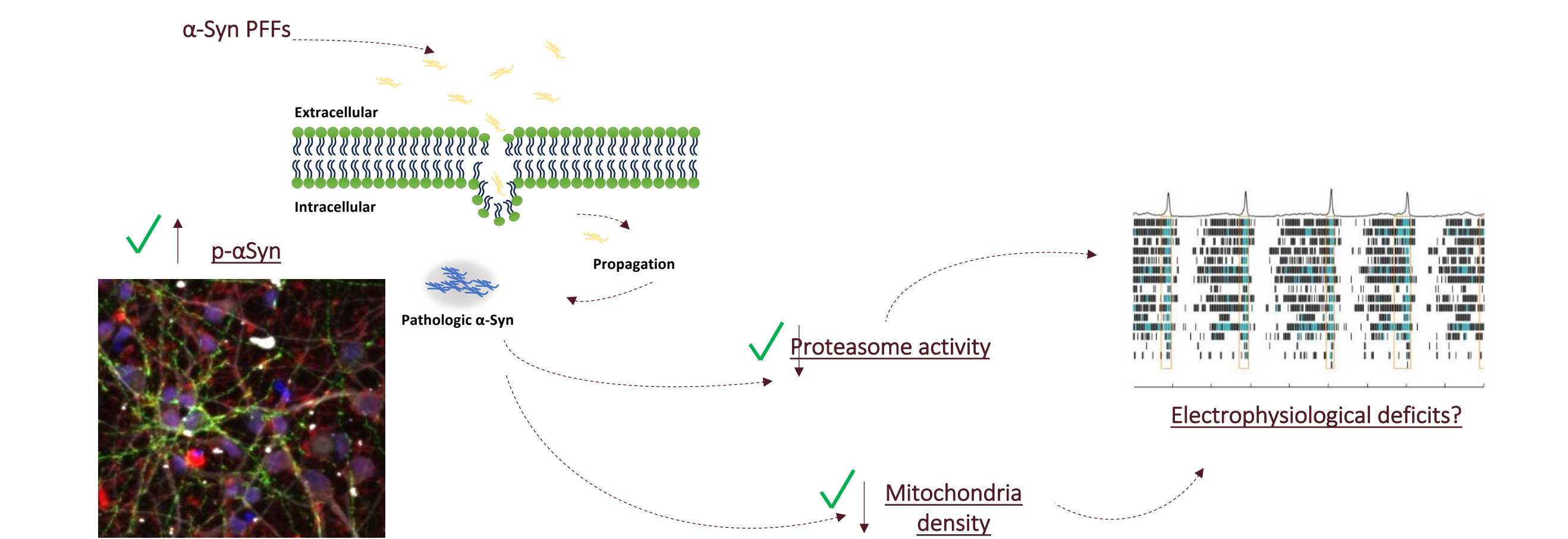
5. Electrophysiology assessment of dopaminergic neurons



Electrophysiology of WT dopaminergic neurons on MEA
Quantification of main MEA metrics in healthy (WT) and in response to D1 and D2 receptor agonists. D1 receptor agonist (SKF 83822): increased firing, bursting and number of network bursts. D2 receptor agonist (Sumanitrole): loss of higher level of firing activity

MEA assay shows a robust activity suitable compound screening and neurotoxicity assessment

Conclusions



Ncardia developed a comprehensive suite of assays designed to recapitulate the key pathological hallmarks of PD using human iPSC-derived neuronal cell models. These advanced in vitro systems provide clinically relevant and translational readouts that enable drug innovators to evaluate compound efficacy and safety at multiple stages of the drug discovery process. By leveraging human-based cellular models, Ncardia helps increase confidence in preclinical predictions, bridge the gap between in vitro and in vivo outcomes, and potentially reduce reliance on animal testing.

Advancing Parkinson’s research with human-relevant iPSC models that enhance predictability and reduce animal testing.

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