



Amyloid Structural Changes Studied by Infrared Microspectroscopy in Bigenic Cellular Models of Alzheimer's Disease

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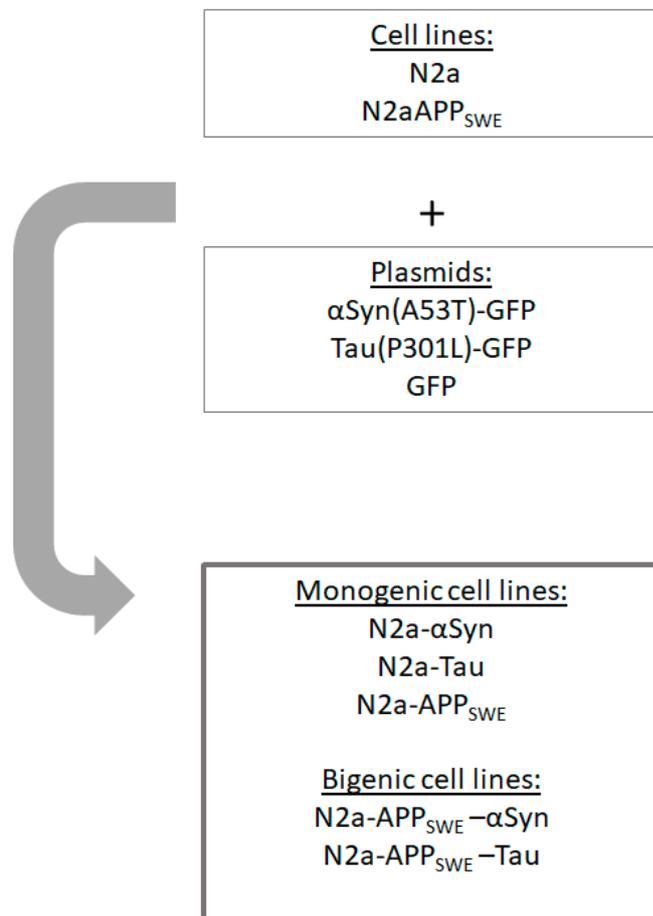


Figure S1. Cell models to study aggregation of amyloid proteins.

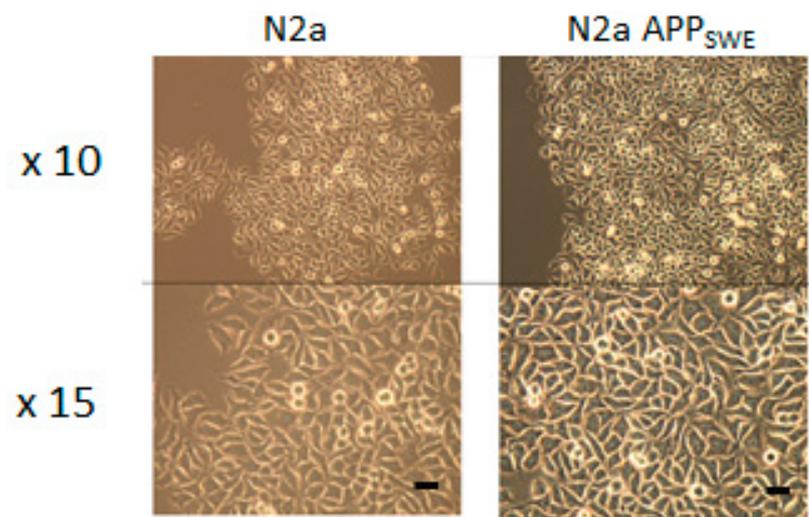


Figure S2. Bright field images of cells before transfection. Scale bar is 20 μ m.

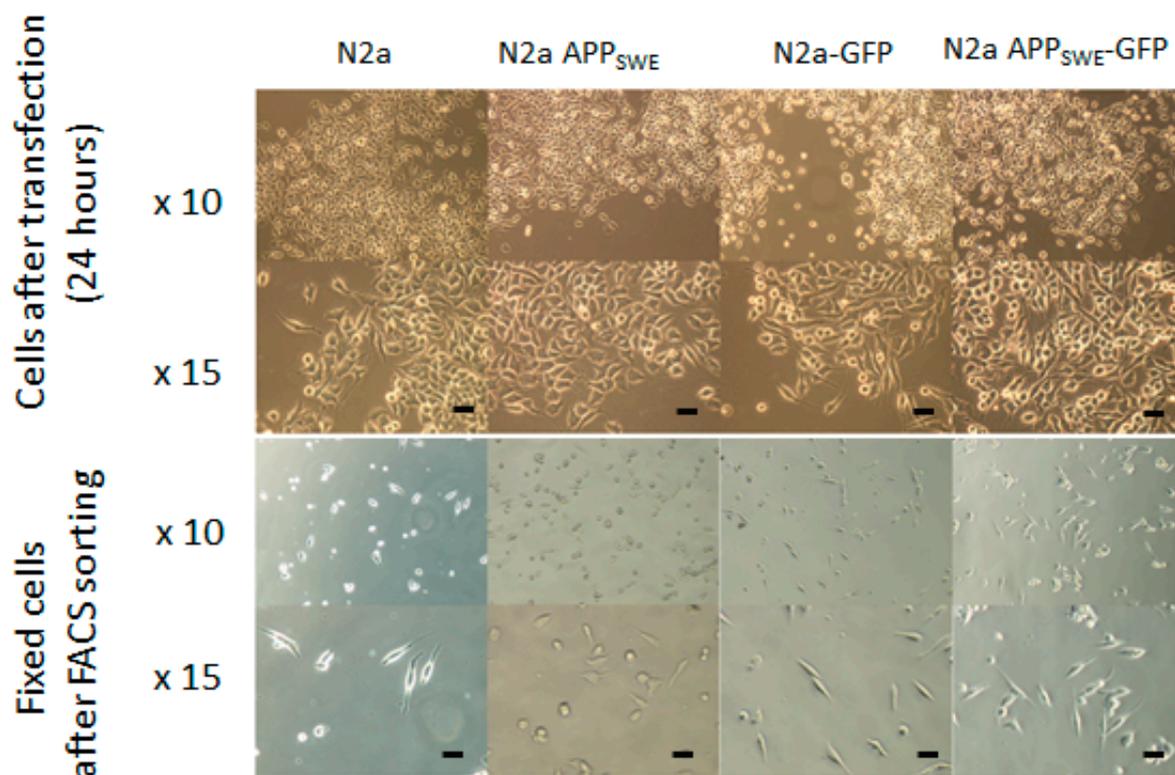


Figure 3. Bright-field images of cells after transfection and FACS sorting. Scale bar is 20 μ m.

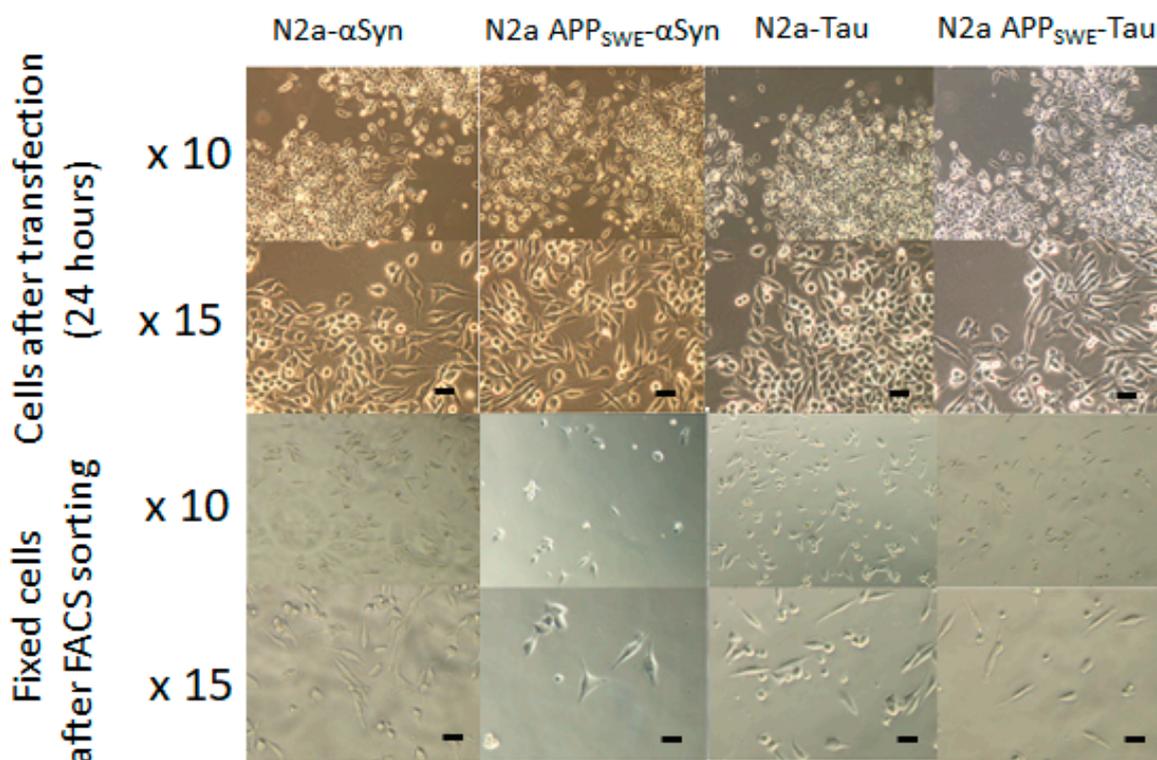


Figure S4. Bright-field images of cells after transfection and FACS sorting. Scale bar is 20 μ m.

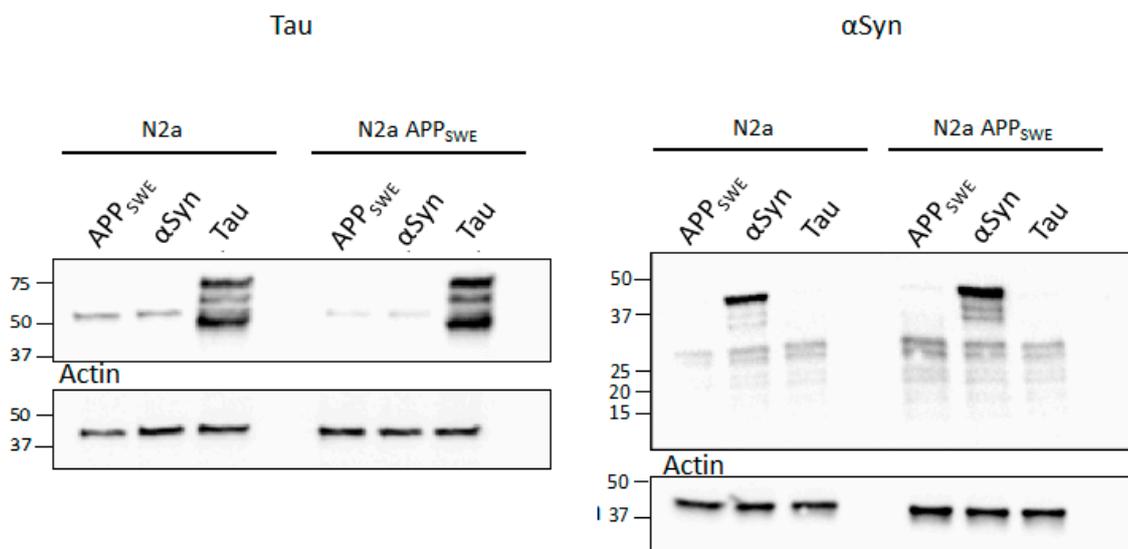


Figure S5. SDS PAGE followed by western blot analysis with specific antibodies against Tau and α Syn expression after transfection.

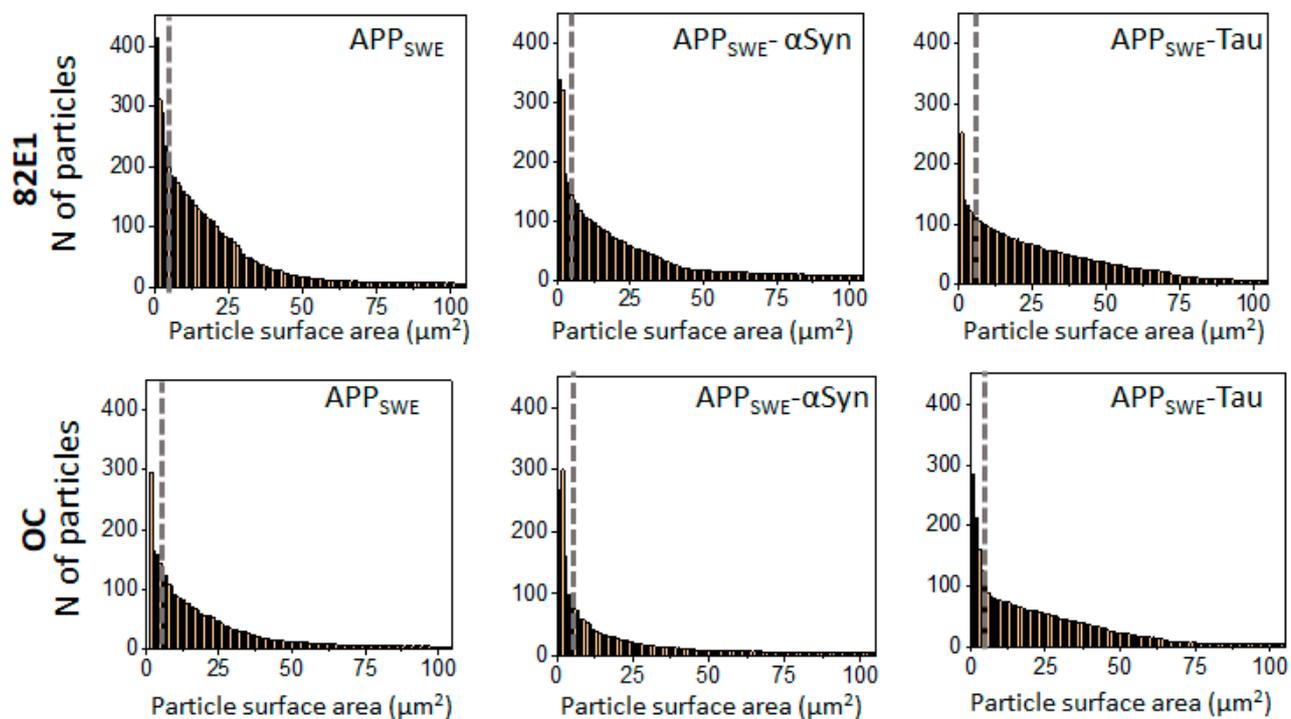


Figure S6. Particle size was quantified by Imaris (Bitplane Scientific Software, Zurich, Switzerland) after surface rendering in confocal images of the cells immunolabeled with antibodies OC and 82E1. Dashed lines indicate a threshold set for the analysis of rendered surfaces in confocal z-stacks that was used for the fluorescent signal quantification.