

Cell line	CRISPR edit	Genotype	NYSCF ID	Status / Estimated Availability
WT parental line: 7889SA			CO0002-01-SV-003	Available
1	<i>APP</i> Knock-In (pathogenic)	WT/Swe	CO0002-01-CS-003	Available
2		Swe/Swe	CO0002-01-CS-004 / BN0013	Available
3		WT/A692G*	BN0009	Available
4		A692G/A692G*	BN0010	Available
5		WT/V717G	BN0011	Available
6		V717G/V717G	BN0012	Available
7	<i>PSEN1</i> Knock-In (pathogenic)	WT/M146V	CO0002-01-CS-002	Available
8		M146V/M146V	CO0002-01-CS-001	Available
9		WT/L166P^	BN0007	Available
10		L166P/L166P	BN0001	Available
11		WT/M233L	BN0005	Available
12		WT/A246E	BN0004	Available
13	A246E/Null	BN0006	Available	
14	Double Knock-In (pathogenic)	Swe/Swe M146V/M146V	BN0002	Available
15	<i>APP</i> Knock-In (protective)	A673T/A673T	BN0003	Available
16	<i>APP</i> Knock-Out	APP-KO/KO	BN0008	Available

* Trisomy 20

^ Partial Trisomy 17

Reference:

Kwart, D. et al. A Large Panel of Isogenic *APP* and *PSEN1* Mutant Human iPSC Neurons Reveals Shared Endosomal Abnormalities Mediated *APP* β -CTFs, Not $A\beta$. *Neuron* **104**, 256–270, e255 (2019).