



Asaro et al., Figure S1

**Fig. S1. *APOE* genotype does not impact levels of sortilin or FABP7 in primary astrocytes**

(A) Immunodetection of FABP7 (red) and sortilin (green) in primary astrocytes from apoE3 mice either (WT) or genetically deficient for *Sort1* (KO). FABP7-stained cells are identified as astrocytes by expression of GFAP (blue). Merged images show co-expression of FABP7 and sortilin. Images represent single z-planes. Scale bar: 20  $\mu$ m. (B) Exemplary western blot analyses of sortilin, FABP5, and FABP7 in primary astrocytes from apoE3 or apoE4 targeted replacement mice, either wild-type (WT) or homozygous for the *Sort1* null allele (KO). Detection of GAPDH served as loading control. (C - D) Quantitative analysis of FABP7 (C) and sortilin (D) levels in primary astrocyte cultures using densitometric scanning of replicate blots (as exemplified in panel B). No genotype-dependent alterations in FABP7 or sortilin levels were seen comparing cells of the indicated *APOE* and *Sort1* genotypes. Data are given as mean  $\pm$  SEM with E3/WT levels set to 100% (n=9 independent cultures per genotype). (E) Levels of *Fabp7* transcript were identical in primary astrocytes from apoE3- and apoE4-expressing mice, either WT or KO for *Sort1*, as determined by quantitative RT-PCR (n=5 independent cultures per genotype). Values are given as log<sub>2</sub> fold change relative to the respective WT (set to value 0).

**Table S1.** List of selected proteins with altered cell surface localization in primary cultures of sortilin-deficient neurons or astrocytes

ACC	Protein	Gene	ratio (KO/WT)	p-value
<b>Neurons</b>				
P15209	BDNF/NT-3 growth factors receptor	<i>Trkb</i>	0.724	0.020768
Q01279	Epidermal growth factor receptor	<i>Egfr</i>	0.793	0.000107
Q04519	Sphingomyelin phosphodiesterase	<i>Asmase</i>	1.53	0.005667
Q8C0E2	Vacuolar protein sorting-associated protein 26B	<i>Vps26b</i>	1.56	0.0035004
P51880	Fatty acid-binding protein, brain (FABP7)	<i>Fabp7</i>	1.52	0.033944
P11404	Fatty acid-binding protein, heart (FABP3)	<i>Fabp3</i>	1.4	0.22835
Q05816	Fatty acid-binding protein, epidermal (FABP5)	<i>Fabp5</i>	1.25	0.11495
<b>Astrocytes</b>				
P15209	BDNF/NT-3 growth factors receptor	<i>Trkb</i>	0.774	0.03436
P28798	Progranulin	<i>Grn</i>	0.646	0.000474
Q06335	Amyloid-like protein 2	<i>Aplp2</i>	0.707	0.000877
P41233	ATP-binding cassette transporter ABCA1	<i>Abca1</i>	0.756	0.003095
P51880	Fatty acid-binding protein, brain (FABP7)	<i>Fabp7</i>	0.94	0.61984
P11404	Fatty acid-binding protein, heart (FABP3)	<i>Fabp3</i>	not detected	
Q05816	Fatty acid-binding protein, epidermal (FABP5)	<i>Fabp5</i>	not detected	

ACC, Uniprot accession number; ratio (KO/WT), ratio between protein amounts in cell surface fractions from sortilin KO and WT primary cell cultures as determined by quantitative label-free proteomics; *p*-value is given based on a resampling ANOVA-based significance test.

**Table S2.** Patient samples examined by western blot analysis. Specimens were obtained from the Netherlands Brain Bank (NL; Netherlands Institute for Neuroscience, Amsterdam) and the MRC London Brain Bank for Neurodegenerative Diseases (UK; Institute of Psychiatry, King's College London).

Case No.	Age	Sex	Clinical diagnosis	Braak stage	ApoE Genotype	Brain Bank
94-110	82	F	AD dementia	VI	4/4	NL
94-037	63	M	AD dementia	VI	4/4	NL
94-082	64	M	AD dementia	VI	4/4	NL
98-060	66	M	AD dementia	VI	4/4	NL
94-091	83	F	AD dementia	VI	3/3	NL
94-025	85	F	AD dementia	VI	3/3	NL
94-016	86	F	AD dementia	VI	3/3	NL
93-140	87	F	AD dementia	IV	3/3	NL
97-009	89	F	AD dementia	VI	3/3	NL
92-022	91	F	AD dementia	IV	3/3	NL
93-011	92	F	AD dementia	VI	3/3	NL
99-114	92	F	AD dementia	IV	3/3	NL
99-123	93	F	AD dementia	IV	3/3	NL
96-020	58	M	AD dementia	VI	3/3	NL
94-028	70	M	AD dementia	VI	3/3	NL
95-077	72	M	AD dementia	VI	3/3	NL
90-066	89	M	AD dementia	IV	3/3	NL
92-024	89	M	AD dementia	IV	3/3	NL
99-064	89	M	AD dementia	VI	3/3	NL
A197/88	81	M	AD dementia	V-VI	3/3	UK
A005/96	89	M	AD dementia	-	3/3	UK
A012/96	89	M	AD dementia	V-VI	3/3	UK
A013/96	85	M	AD dementia	V-VI	3/3	UK
A342/96	67	M	AD dementia	V-VI	3/3	UK
A277/97	89	M	AD dementia	V-VI	3/3	UK
A065/02	82	M	AD dementia	V-VI	3/3	UK
A093/97	91	M	AD dementia	V-VI	4/4	UK
A200/97	74	M	AD dementia	-	4/4	UK
A213/94	81	F	AD dementia	III-IV	3/3	UK
A335/94	72	F	AD dementia	V-VI	3/3	UK
A125/95	88	F	AD dementia	V-VI	3/3	UK
A044/96	84	F	AD dementia	V-VI	3/3	UK
A097/96	90	F	AD dementia	V-VI	3/3	UK
A291/96	60	F	AD dementia	V-VI	3/3	UK
A407/96	75	F	AD dementia	V-VI	3/3	UK
A025/97	80	F	AD dementia	V-VI	3/3	UK

A026/98	80	F	AD dementia	V-VI	3/3	UK
A028/98	70	F	AD dementia	V-VI	3/3	UK
A207/98	91	F	AD dementia	V-VI	3/3	UK
A133/99	92	F	AD dementia	III-IV	3/3	UK
A053/95	76	F	AD dementia	V-VI	4/4	UK
A240/95	75	F	AD dementia	V-VI	4/4	UK
A366/95	79	F	AD dementia	V-VI	4/4	UK
A073/96	89	F	AD dementia	V-VI	4/4	UK
A133/97	92	F	AD dementia	-	4/4	UK
A053/98	89	F	AD dementia	V-VI	4/4	UK

**Table S3.** Patients examined by immunohistology

Case No.	Age	Sex	Post-mortem interval (hrs)	Clinical diagnosis	Thal Phase	Braak stage	CERAD neuritic plaque score	AD neuropathological change
1	71	M	12	AD dementia	4	V-VI	3	A3, B3, C3
2	75	M	6	AD dementia	4	V-VI	2	A3, B3, C2
3	82	F	24	AD dementia	4	V-VI	3	A3, B3, C3

**Neuropathological assessments:** All cases were extensively characterized according to the routine protocol for neurodegenerative diseases. Neuropathological diagnosis of AD was made following the NIA-AA criteria including Thal phasing for A $\beta$  load, Braak-and- Braak-staging for NFTs, and CERAD neuritic plaque score to assess the density of neuritic plaques. (Clin Pathol. 2019 Nov;72(11):725-735; Neuropathology of Neurodegenerative Diseases. (2014). In G. Kovacs (Ed.), Neuropathology of Neurodegenerative Diseases: A Practical Guide (pp. I-II). Cambridge: Cambridge University Press).

**Table S4. Neurons**

The complete data set of proteins identified in the surface proteome of wild-type and sortilin-deficient primary neurons is provided as an Excel file.

[Click here to download Table S4](#)

**Table S5. Astrocytes**

The complete data set of proteins identified in the surface proteome of wild-type and sortilin-deficient primary astrocytes is provided as an Excel file.

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