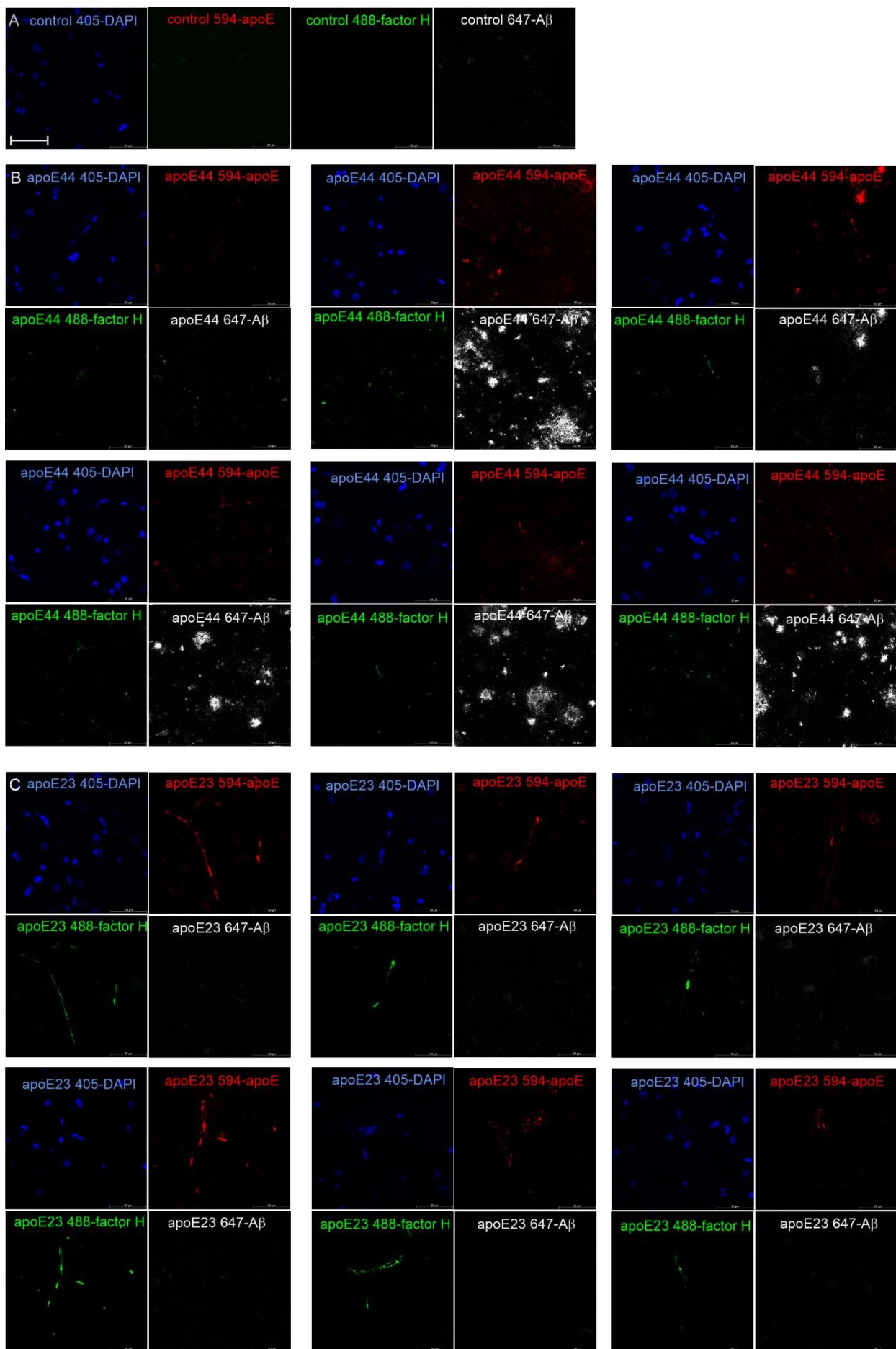


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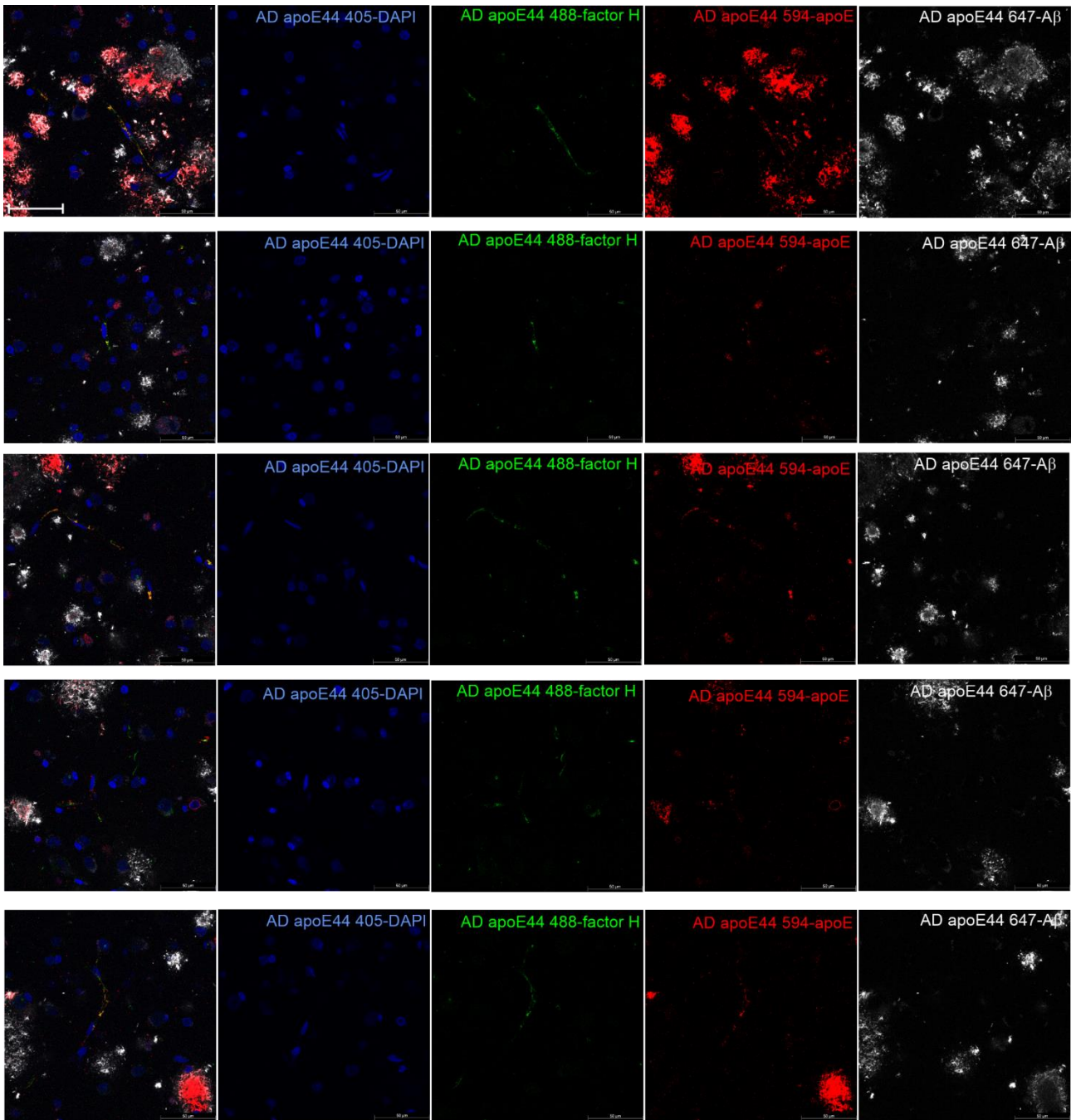
Appendix Fig. S1. Localization of apoE2/3/4 and FH in brain A β plaques. Related to Fig. 1A-B.

Immunofluorescence staining of

(A) control and six microscope images (size of the scale bar = 50 μ M) from the

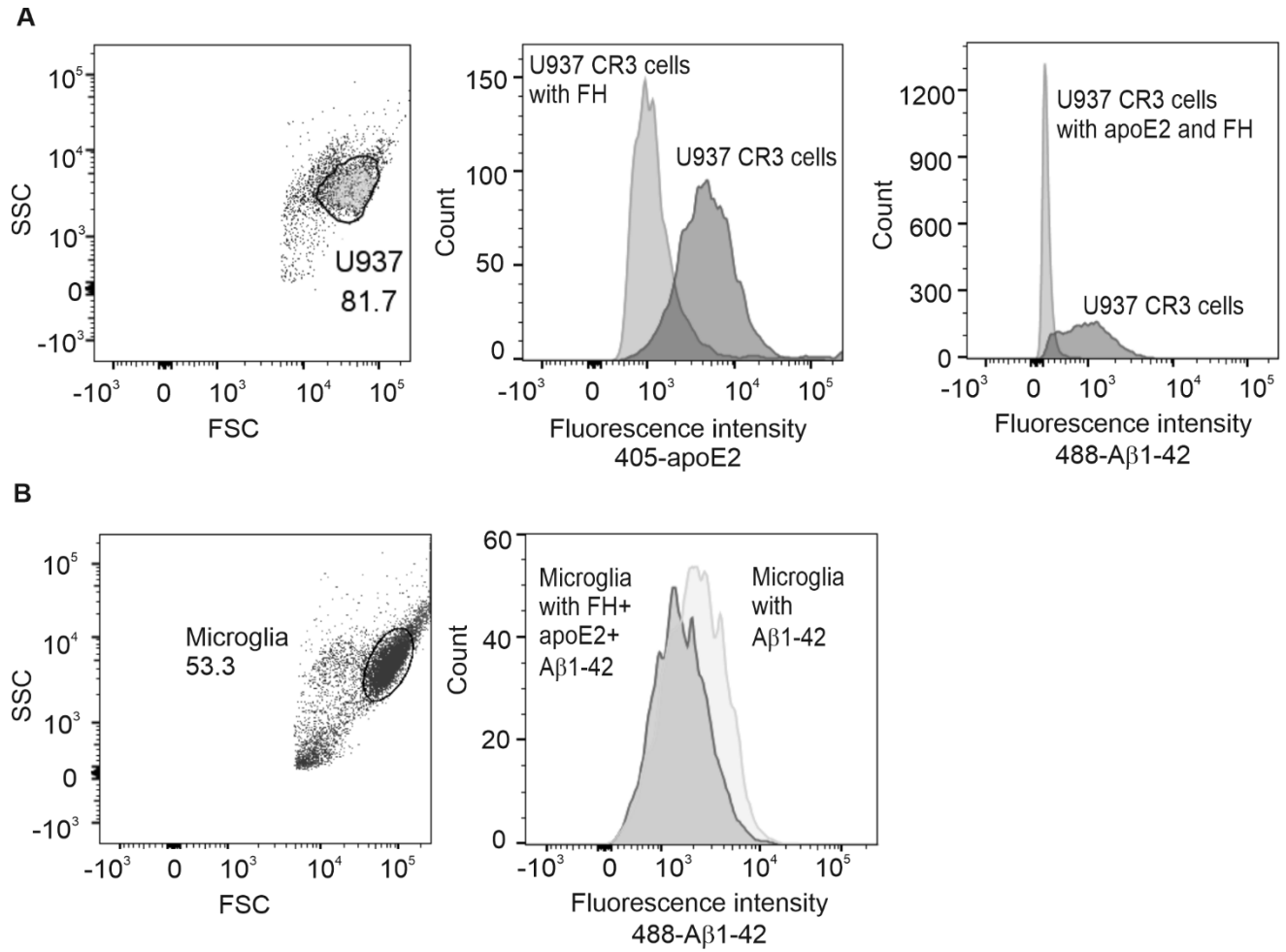
(B) apoE44 (n=2) and

(C) apoE23 (n=2) genotyped iNPH patient biopsy samples shown in separate channels. Staining of (green) FH, (red) apoE (white) A β plaques and (blue) nuclei are shown.



Appendix Fig. S2. Localization of apoE4 and FH in brain Aβ plaques. Related to Fig. 1C.

Immunofluorescence staining of six microscope images shown in separate channels from a biopsy sample obtained from an iNPH patient diagnosed with Alzheimer's clinical syndrome (ACS) (size of the scale bar = 50 μM). Staining of (green) FH, (red) apoE (white) Aβ plaques and (blue) nuclei are shown.



Appendix Fig. S3. Gating and histograms of U937 CR3 cells and SV40 microglia.

(A) Related to Fig. 3F-I. (left) Gating of Mn^{2+} activated U937 CR3 cells (81.7% of all cells in the gate). (Center) Inhibition of apoE binding by FH. (Right) Inhibition of $A\beta$ binding by apoE2 and FH.

(B) Related to Fig. 5A (left) Gating of microglia SV40 cells (53.3% of all cells in the gate). (right) Inhibition of $A\beta$ phagocytosis by apoE2 and FH. SSC, side scatter and FSC, forward scatter.

Appendix Table S1. Patient data

NPH case number	age	sex	APOE genotype	Aβ pathology	Tau
1	82	M	34	yes*	no
2	79	M	23	no*	no
3	84	M	34	yes*	no
4	71	F	33	no*	no
5	59	M	33	yes*	no
6	72	M	33	yes*	no
7	80	M	44	yes*	no
8	66	M	33	yes*	no
9	58	F	33	no*	no
10	65	M	34	no*	no
11	87	M	33	yes*	no
12	79	M	33	no*	no
13	82	M	34	yes*	no
14	79	F	33	yes**	yes
15	77	F	33	yes*	no
16	72	F	44	yes**,***	yes
17	78	F	33	no*	no
18	71	M	23	no*	no
19	85	F	34	yes*	no
20	83	F	34	yes*	no
21	84	M	23	no*	no
22	70	F	34	no*	no
23	74	F	44	yes**	yes

Estimated Braak stage *1-4 or **5-6, *** Patient diagnosed for Alzheimer's clinical syndrome (ACS)

Appendix Table S2. Quantification of binding parameters using dissociation constants (K_d) fit model of normalized (ΔF_{norm}) values in MST (Related to Fig. 1I-K).

Ligand name	n	FH (nM)	K_d (μM)*	K_d Confidence	Reduced χ^2	Signal to Noise
native apoE2	3	2	0.002	0.0007		
native apoE4	3	2	0.3	0.2		
Ligand name	n	NT657-apoE2(1-165) (nM)	K_d (μM)*	K_d Confidence	Reduced χ^2	Signal to Noise
FH402H	3	2.8	0.496	0.139	1.8	16.3
FH402Y	3	2.8	0.253	0.96	1.7	10.2
FH1-4	3	2.8	0.999	0.598	2.1	9.6
FH19-20	3	2.8	no binding		3.1	0
Ligand name	n	NT647-FH5-7 402Y (nM)	K_d (μM)*	K_d Confidence	Reduced χ^2	Signal to Noise
apoE2	3	35	1.005	0.458	1.6	9.0
apoE3	3	35	1.384	0.553	3.1	11.0
apoE4	3	35	1.945	0.1162	0.7	8.1

*Reliable binding affinity was obtained in measurements with a signal to noise ratio > 5 and K_d confidence in the range of 68% using MO. Affinity Analysis Software. When a reliable binding curve could not be assigned, signal to noise ratio shows 0. Reduced χ^2 estimates the goodness of fit (1 = good fit).

Appendix Table S3. Band intensities in WBs and PAGE (Related to Fig. 2B)

SDS-PAGE and silver stain (Related to Fig. 2B, left)									
	description	apoE4/A β /FH (intensity)	apoE4/A β (intensity)	apoE2/A β /FH19-29 (intensity)	ApoE2/A β /FH (intensity)	apoE2/A β (intensity)	apoE2 (intensity)	A β /FH (intensity) *	A β (intensity) *
1	apoE/A β /FH or A β /FH complex	14868	0	0	18881	0	0	17819	0
2	apoE/A β complex 1	0	0	2165	0	408	0	0	0
3	apoE/A β complex 2	0	0	2795	0	908	0	0	0
4	apoE/A β complex 3	0	0	3318	0	889	0	0	0
5	apoE/A β complex 4	0	0	11519	0	4658	0	0	0
6	apoE/A β complex 5	0	0	15017	0	9765	0	0	0
7	apoE/A β complex 6	0	0	20758	2889	4746	0	0	0
8	apoE/A β complex 7/apoE	0	0	17614	5572	7229	2108	0	0
9	apoE/A β complex 8/apoE	5497	5076	112742	7516	9213	8474	0	0
10	A β	1611	3767	7351	3287	10644	0	6769	9527
WB (Related to Fig. 2B, middle)									
	description	apoE2/A β /FH19-20 (intensity)	apoE2/A β /FH (intensity)	apoE2/A β (intensity)	A β /FH (intensity)	A β (intensity)	A β monomer (intensity)	apoE4/A β /FH (intensity) *	apoE4/A β (intensity) *
1	A β aggregate	767	897	2872	16772	21369	0	0	282
2	apoE/A β /FH or A β /FH complex	0	16467	0	2089	0	0	0	0

3	apoE/A β complex 1	5347	0	425	0	0	0	0	0
4	apoE/A β complex 2	15842	0	9148	0	0	0	0	0
5	apoE/A β complex 3	18735	0	11294	0	0	0	0	0
6	apoE/A β complex 4	18932	0	12551	0	0	0	0	0
7	apoE/A β complex 5	17650	0	14099	0	0	0	0	0
8	apoE/A β complex 6	15514	3667	13637	0	0	0	0	0
9	apoE/A β complex 7	23919	5862	10544	0	0	0	0	0
10	apoE/A β complex 8	18682	3166	7645	0	0	0	18118	20733
11	A β	7381	8376	10658	3639	6536	12186	18775	15802

WB (Related to Fig. 2B right)

	description	apoE2/A β (intensity)	apoE3/A β (intensity)	apoE4/A β (intensity)	apoE2/A β /FH (intensity)	apoE3/A β /FH (intensity)	apoE4/A β /FH (intensity)
1	apoE/A β /F H complex	0	0	0	33129	3825	0
2	apoE/A β complex 1	32125	0	0	0	0	0
3	apoE/A β complex 2	36240	1612	0	0	0	0
4	apoE/A β complex 3	48769	27212	0	41821	10717	0
5	apoE/A β complex 4	31930	40914	0	46849	36685	0
6	apoE/A β complex 4	38548	71915	43404	52189	58330	57632
7	A β monomer	37307	52884	52884	52694	48099	57091

*not shown in the gel image

Appendix Table S4. Band intensities in WB (Related to Fig. 4E)

		C3 (intensity)	C3b (intensity)	iC3b (intensity)	A β /BSA (intensity)	A β (intensity)	A β /FH (intensity)	E2/A β (intensity)	E2/A β /FH (intensity)	E4/A β (intensity)	E4/A β /FH (intensity)
1	C3 α and C3b α	16126	14434	3049	13526	19537	13665	17663	12098	16637	15262
2	C3 β	21402	7478	24132	6985	13315	13265	13856	8094	10099	13884
3	iC3b α	0	0	27006	0	0	13309	1572	5116	0	10140

Appendix Table S5. Differential expression of transcripts in microglial cells upon stimulation with different combinations of apoE, A β 1-42 and FH (Related to Table 1 in the manuscript).

Gene name (alternative name)	Gene name extended/encoded protein	Description	Cited in
Upregulation with FH and apoE2			
CIAPIN (Anamorsin)	Cytokine induced apoptosis inhibitor	CIAPIN has been suggested to inhibit neuronal apoptotic cell death.	(Yun <i>et al</i> , 2014)
GPR176	G protein coupled receptor 176	No known association of this particular GPR. Several studies have suggested that GPRs play a role in AD pathogenesis.	(Thathiah & De Strooper, 2011)
TRIM11	Tripartite motif containing 11	TRIM11 can suppress induced neurotoxic effects due to its interaction with, humanin, a neuroprotective peptide.	(Niikura <i>et al</i> , 2003)
MADD	MAP kinase activating death domain	MADD expression has been found to be inhibited specifically in hippocampal neurons in AD; this is associated with an increased neuronal cell death. Some mutations of MADD are linked with AD onset.	(Del Villar & Miller, 2004; Saad <i>et al</i> , 2015)
HGS (Hrs)	Hepatocyte growth factor- regulated tyrosine kinase substrate	HGS has been found to be significantly decreased in AD brains.	(Gireud-Goss <i>et al</i> , 2020)
NISCH	Nischarin	Correlates negatively with the presence of apoE4 disease allele in this study	
CES2	Carboxylesterase 2	CES2 may play a role in protecting the central nervous system from toxic esters. Perhaps a component of the BBB system.	(Zhang <i>et al</i> , 2002)
ITSN2 and ITSN2	Intersectin 1 and Intersectin 2	ITSN1 and ITSN2 are found to be one of the most expressed genes in AD.	(Herrero-Garcia & O'Bryan, 2017)
FHL2	Four and a half LIM domains 2	Interacts with presenilin 2 that has a role in APP processing.	(Tanahashi & Tabira, 2000)
FBN1	Fibrillin-1	FBN1 has a role in supporting the BBB.	(Van der Donckt <i>et al</i> , 2015)
PLXNA1	Plexin A1	Correlates negatively with A β pathology.	(Huang <i>et al</i> , 2021)
PTPRF	Protein Tyrosine Phosphatase Receptor Type F	Is involved in cell adhesion pathway and correlates negatively with A β pathology and with the presence of apoE4 allele in this study.	(Huang <i>et al</i> , 2021; Jiang <i>et al</i> , 2021)
RBBP7	Histone-binding protein RBBP7	Correlates negatively with A β pathology.	(Huang <i>et al</i> , 2021)

Upregulation with FH and apoE4			
ADNP	Activity dependent neuroprotector homeobox	ADNP is found to be significantly decreased in the serum of AD patients.	(Gozes, 2015)
RAP2B	Member of RAS oncogene family	A significant AD associated SNP is located near RAP2B on chromosome 3.	(Yan <i>et al</i> , 2021)
PAFAH1B2	Platelet activating factor acetylhydrolase α	PAFAH1B2, if less expressed, decreases A β formation helping its degradation at its previous stages.	(Page <i>et al</i> , 2012)
ERO1A	Endoplasmic reticulum oxidoreductase 1 alpha	No directly known association. The role of ERO1A is implied in promoting calcium induced apoptosis.	(Seervi <i>et al</i> , 2013)
LARP1	La-related protein 1	LARP1 associates mTOR complex and can be functionally linked to AD.	(Wang <i>et al</i> , 2015)
WNT5A	Wnt family member 5A	Abnormal expression of WNT5A is linked with inflammation and cytotoxicity promoted by A β in neurons.	(Li <i>et al</i> , 2011)
FOXP1	Forkhead box protein P1	No directly known association.	
PDK1	Pyruvate dehydrogenase kinase 1	PDK1 is linked with toxicity induced by A β since it decreases α -secretase activity and promotes AD disease progression.	(Checler, 2013; Pietri <i>et al</i> , 2013)
FOXO3	Forkhead box O3	FOXO3 is involved in A β induced neuronal death, and in production of dysfunctional A β .	(Shi <i>et al</i> , 2016)
DAAM1	Dishevelled Associated Activator Of Morphogenesis 1	Correlates positively with A β pathology.	(Huang <i>et al</i> ., 2021)
PGBD1	PiggyBac Transposable Element Derived 1	Correlates positively with A β pathology.	(Huang <i>et al</i> ., 2021)
TRA2A	Transformer 2 Alpha Homolog	Correlates positively with the presence of apoE4 disease allele in this study	

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