

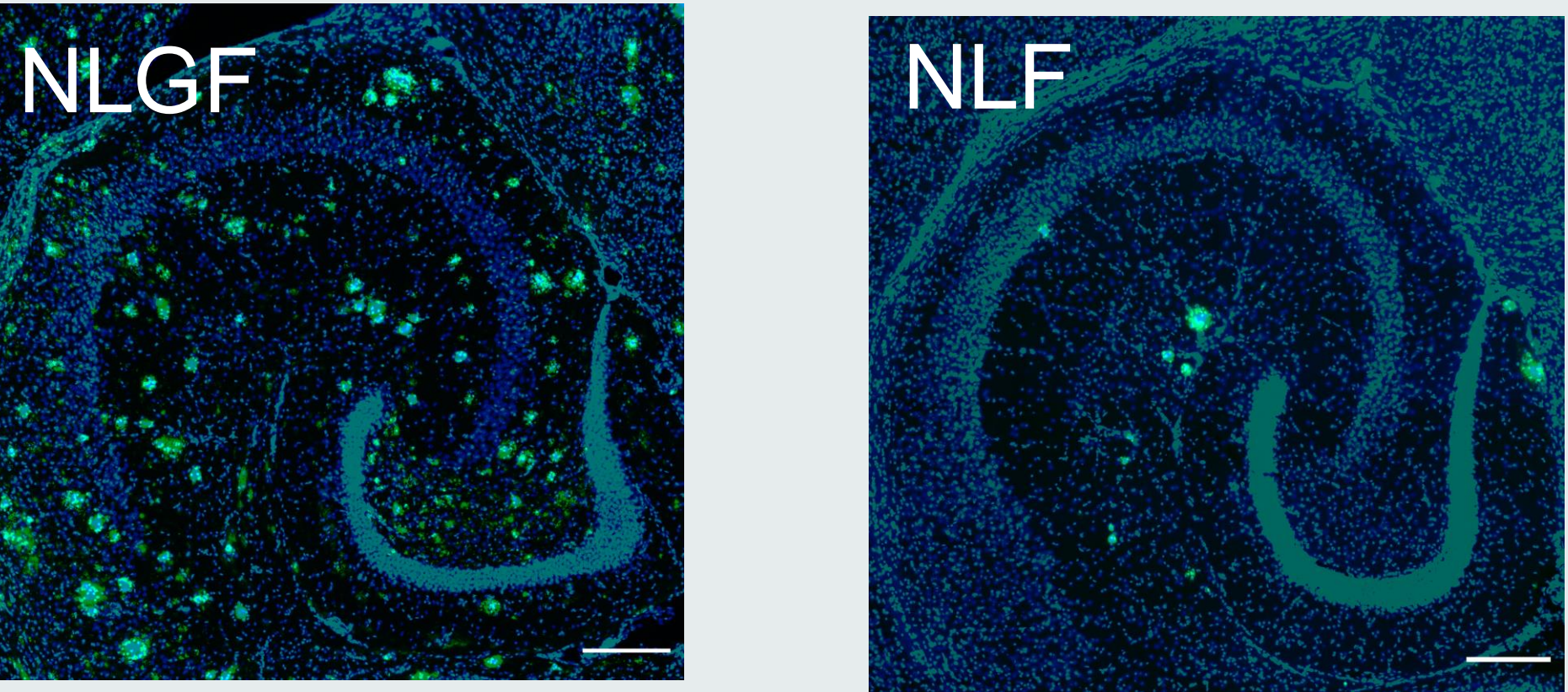
Knock-in mouse model, App^{NLF/NLF}, with slow plaque development into old age, shows plaque specific gene expression pattern more similar to that found in post-mortem tissues from people with Alzheimer's disease.

Aya Balbaa¹, Jack I. Wood¹, Aishwarya Pathak¹, Eugenia Wong¹, Ridwaan Joghee¹, Sneha Desai¹, Damian M.Cummings¹, Takshashila Tripathi¹, John Hardy¹, Frances A.Edwards¹

¹Department of Neuroscience, Physiology and Pharmacology, UCL



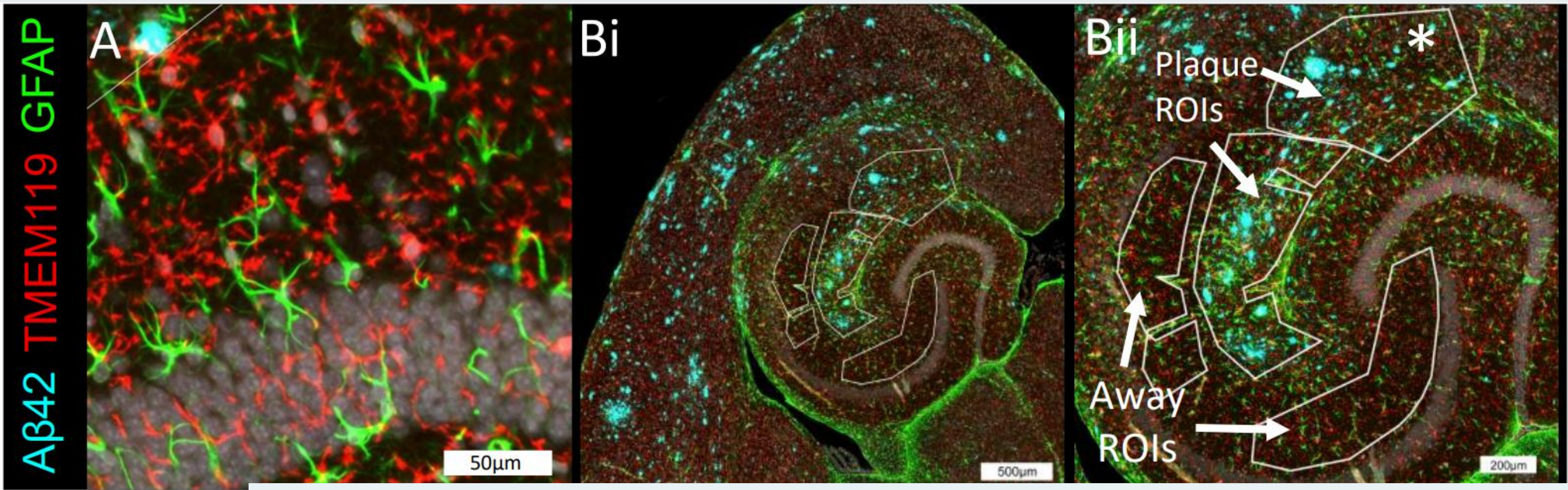
Introduction.
The APPKI mice NLF and NLGF (Saito et al., 2014) are widely used.
NLGF mouse (like many transgenic mice): an aggressive model laying down plaques rapidly from 2 months.
NLF mouse (more like progression of preclinical Alzheimer's disease): earliest plaques in middle age (around 9 months); slow progression into old age.
New technologies Nanostring GeoMx regional transcriptomics allows study of more relevant.
Age an important factor in gaining translational insights in mouse models.



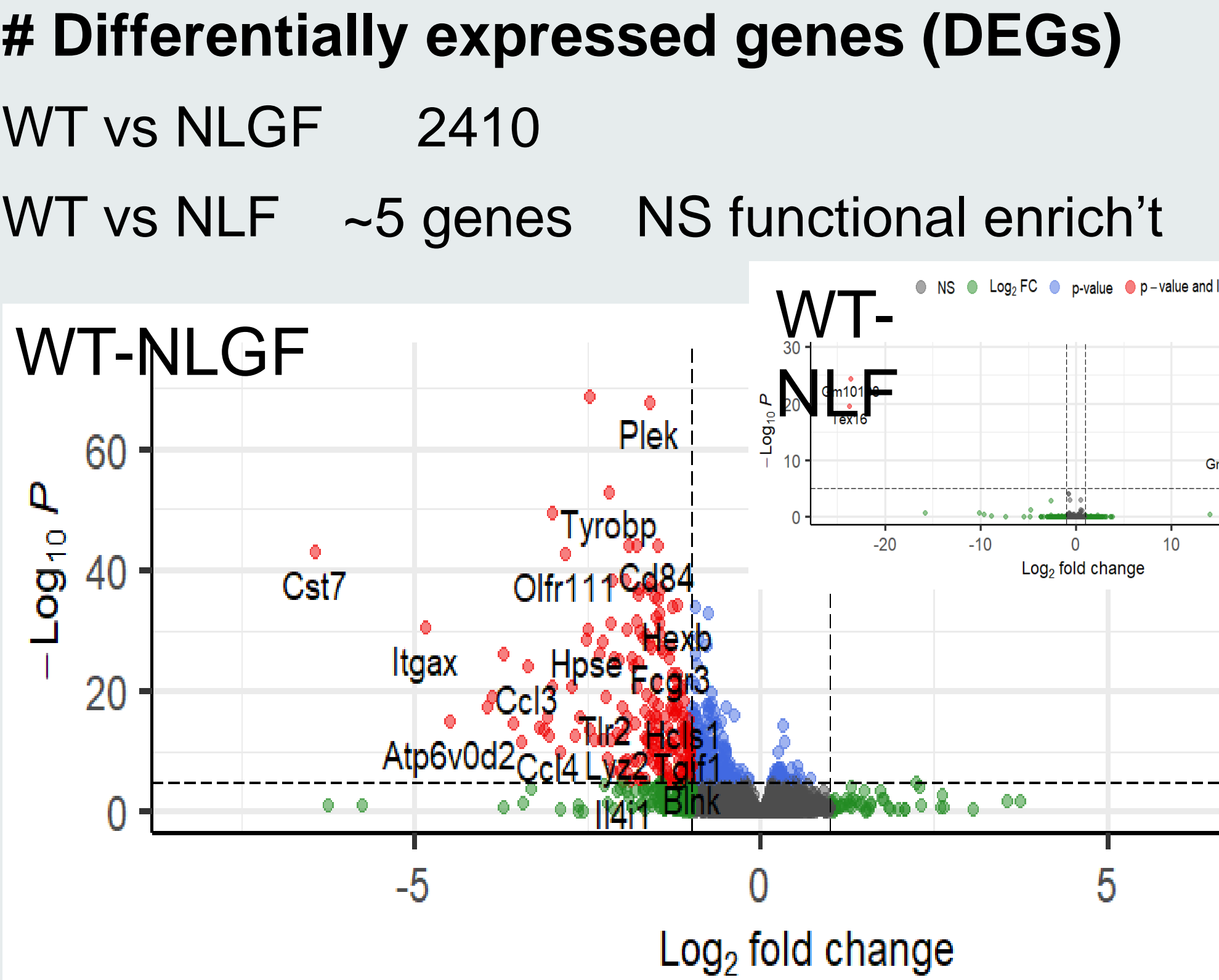
Methods: Whole hippocampal RNA was extracted from WT, NLF and NLGF mice (n=9-12/genotype; mixed sex; 18 months old).
RNAseq: Differential expression analysis compared WT to NLF or NLGF using DESeq2 R package, batch effect was accounted for as needed.

Regional cell-type enriched transcriptomics.
Nanostring GeoMx technology: AOIs microglia, astrocytes (and underlying synapses) on, near or far from plaques

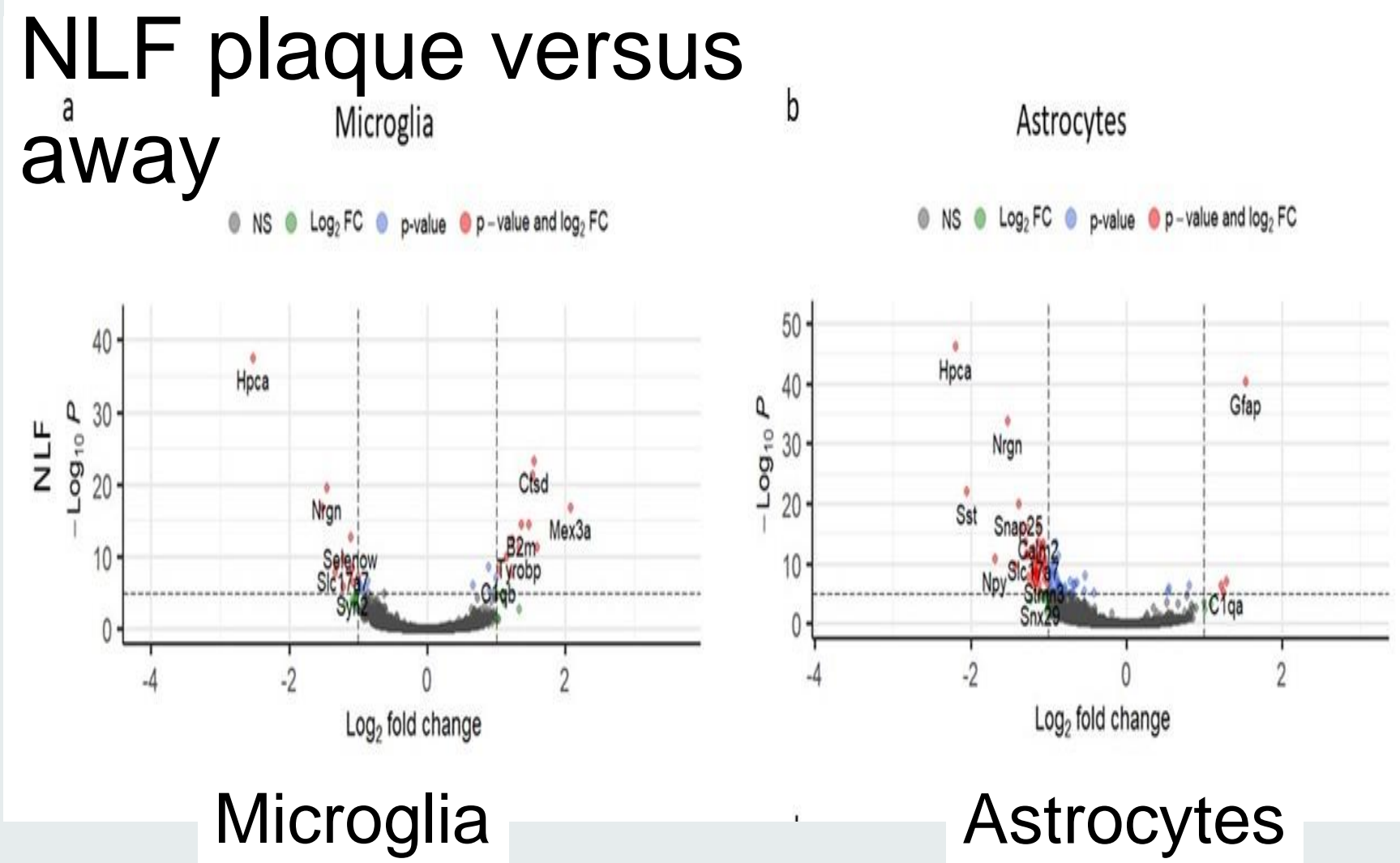
Comparing regions with dense plaques (**Plaque ROIs**) to regions away from plaques (**Away ROIs**) (n=6, 4 female, 2 male). Analysed using the limma-voom pipeline; within tissue pairing from plaque versus away.



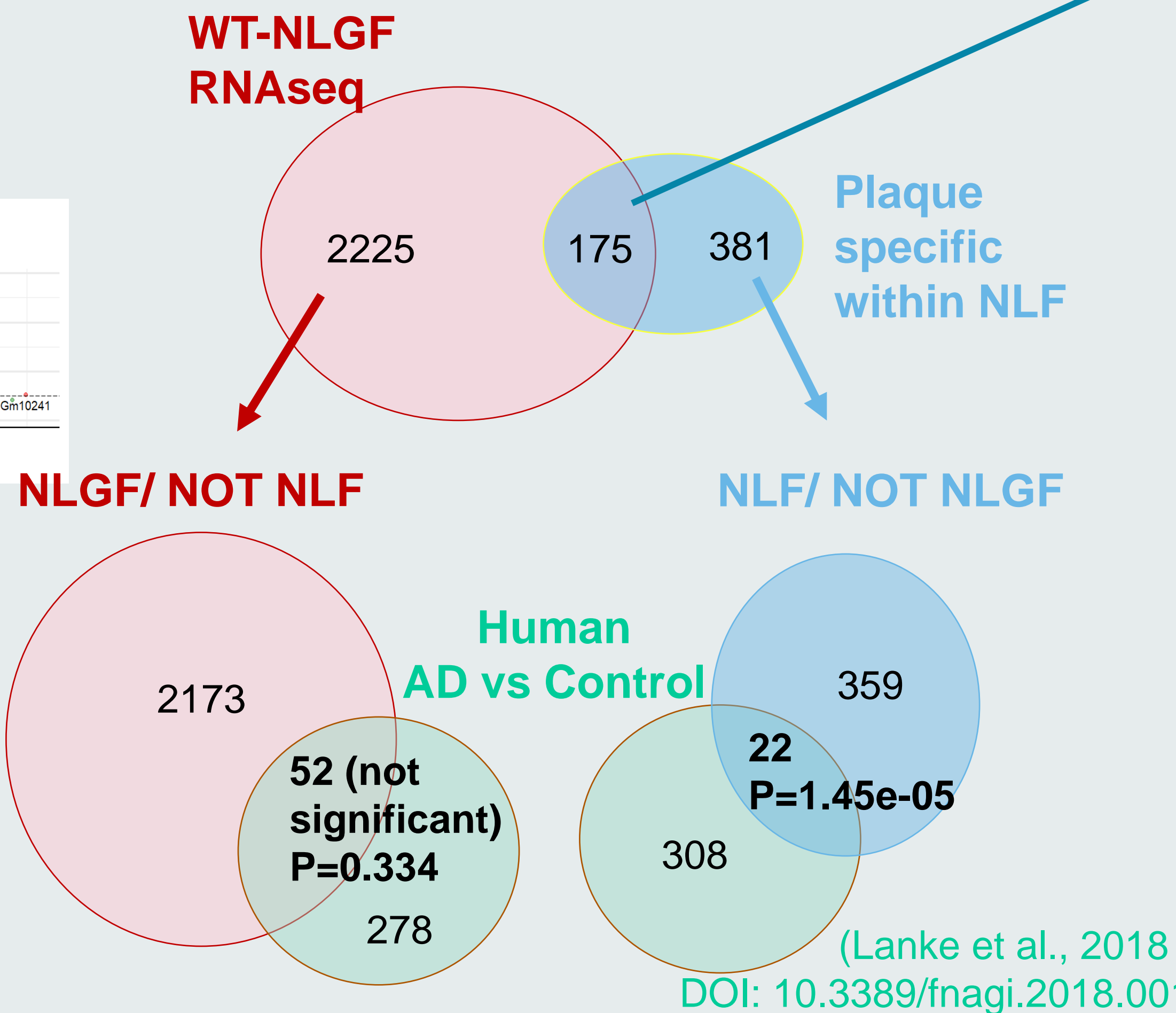
RNA-Seq from whole hippocampus only useful for NLGF:



NLF: Regional cell-enriched transcriptomics: comparing plaque and away within tissue brings up 556 DEGs



NLGF plaque-induced genes (versus WT) compared to NLF plaque-induced genes (vs far from plaque)



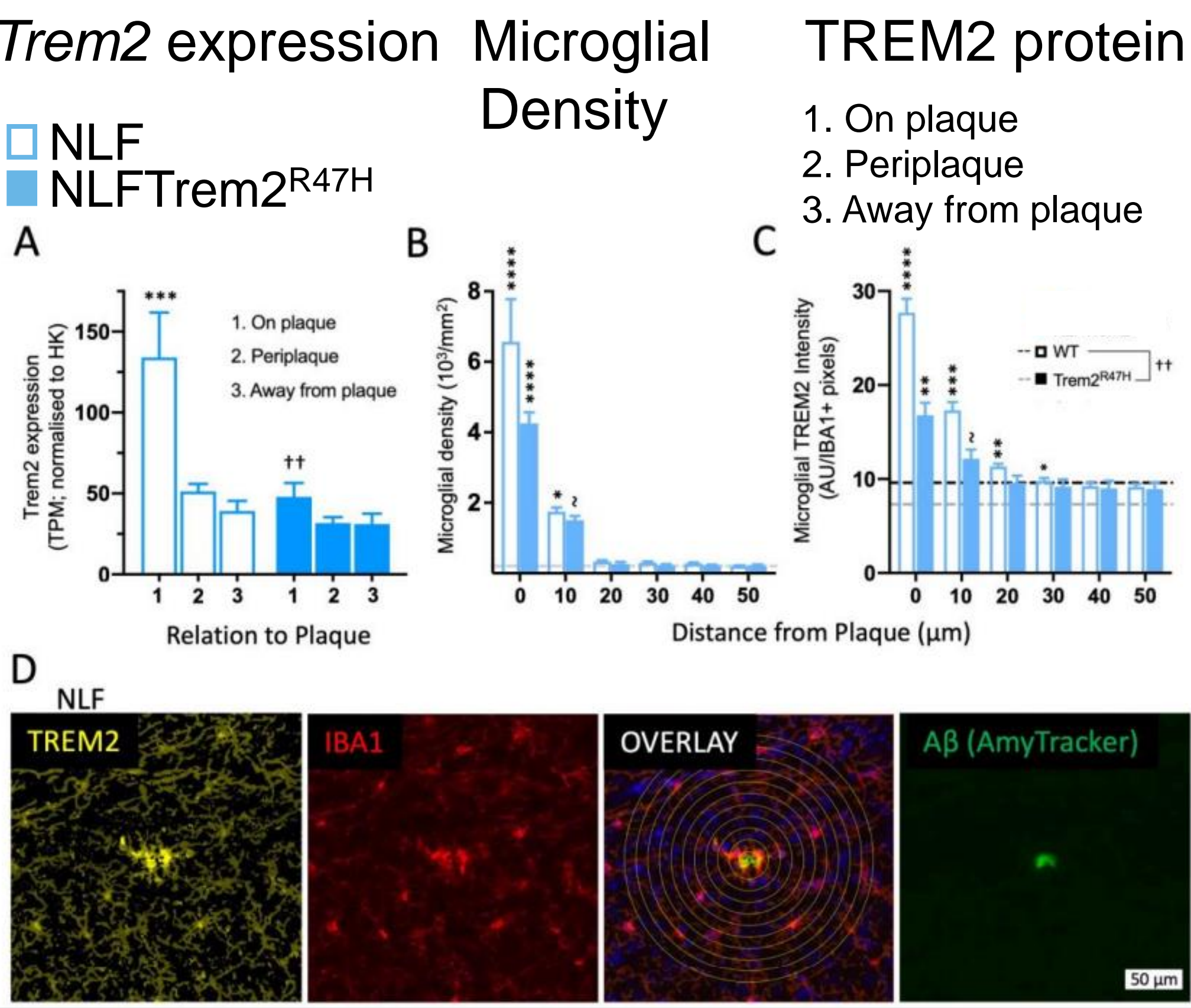
	1	2	3	4	5	6	7
AMPH							
ANK2,3							
BSN							
CACNG8							
CAMK2B							
CNIH2							
COX411							
COX5B, COX6A1, COX6B1, COX6C, COX8A							
CPLX1							
CPNE6							
DLG2							
ERC2							
FKBP1A							
GABRA5							
GRIA1, GRIA2							
GRIN2A							
HPCA							
KCNIP2							
KCNQ2							
KIF1B							
LIN7B							
MAPK8IP2							
MINK1							
NDUFA4							
NDUFAB1							
NDUFB9							
NDUF55							
NEFL							
NETO1							
NPTX1							
NPTXR							
NPY							
PLP1							
PRRT1							
PTK2B							
RIMBP2							
RIMS1							
SHISA6							
SLC12A4, SLC17A7, SLC1A2							
SNCA, B							
STX1A							
TSPOAP1							

NLF-plaque exclusive genes enriched for synaptic and mitochondrial functions

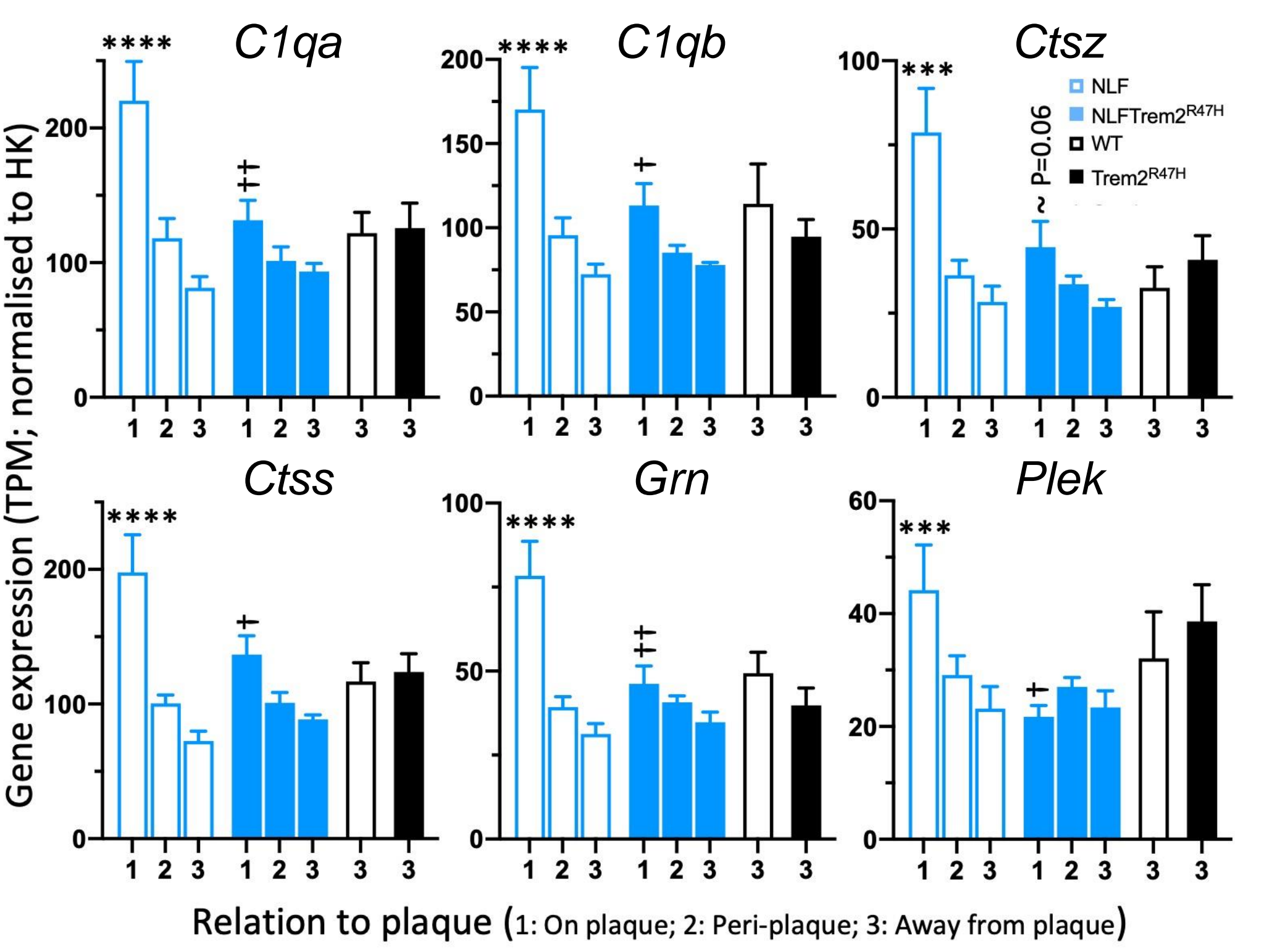
- BIOLOGICAL FUNCTIONS**
1. regulation of neurotransmitter receptor activity (GO:0099601)
 2. regulation of cation channel activity (GO:2001257)
 3. chemical synaptic transmission (GO:0007268)
 4. mitochondrial electron transport, cytochrome c to oxygen (GO:0006123)
 5. regulation of NMDA receptor activity (GO:2000310)
 6. anterograde trans-synaptic signaling (GO:0098916)
 7. aerobic electron transport chain (GO:0019646)

Plaque-dependent Microglial gene modules coexpressing in NLF and NLGF mice similar to previously studied genotypes

Some microglial genes, including *Trem2*, only increase in expression on plaque contact.



A small subset are Trem2-genotype-dependent



Conclusion
The NLF mouse shows similarities to the human condition of Alzheimer's disease that makes it likely a more translatable model. The TREM2 dependent roles of microglia depend on the microglia touching the plaque and so must depend on a lipophilic ligand