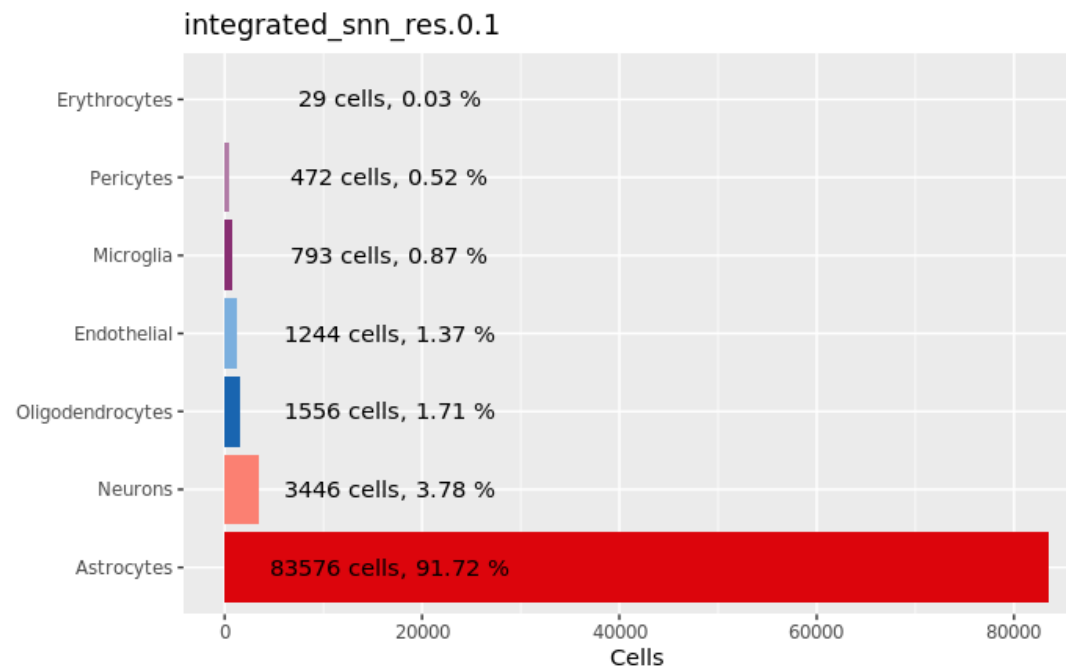
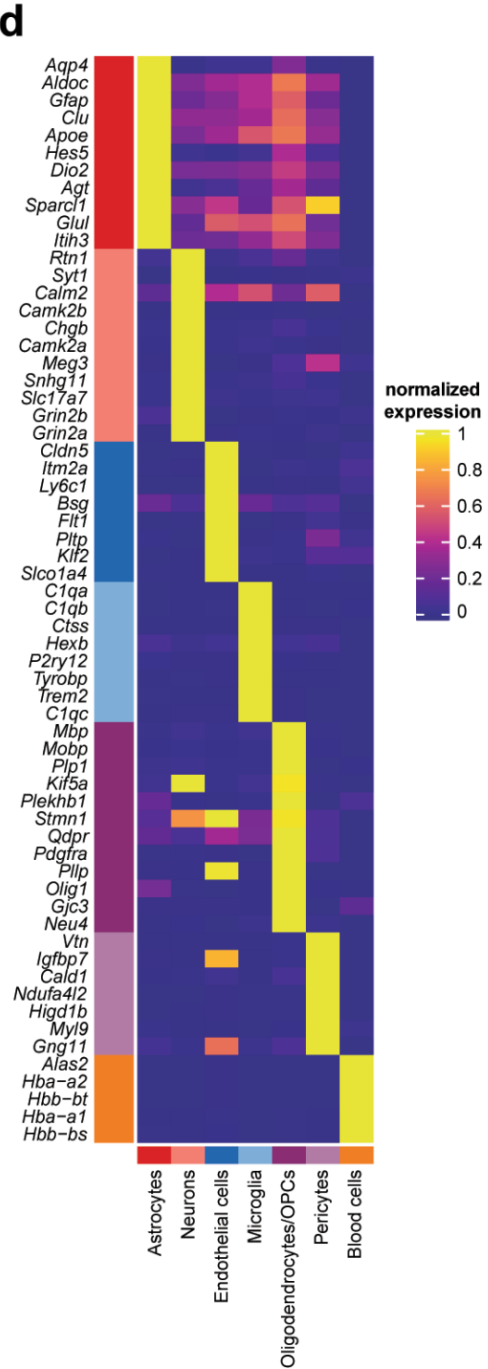


**Extended Data Fig. 1c**, Initial clustering of all 91110 cells across all 12 animals identifies presence and percentage of non-astrocytic cells as defined by cell type marker genes

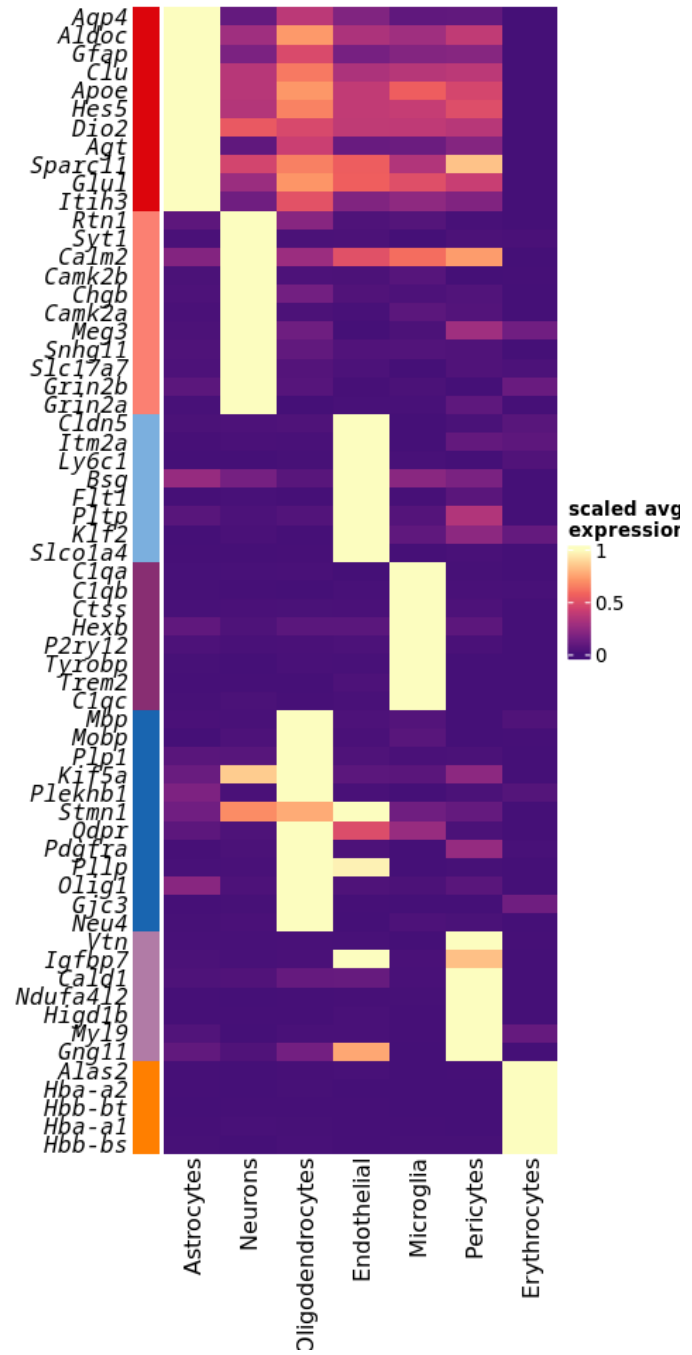


Reproduction of cell type annotation

**Extended Data Fig. 1d,**  
Heatmap of  
normalized cell type  
marker genes across  
the identified cell type  
clusters.

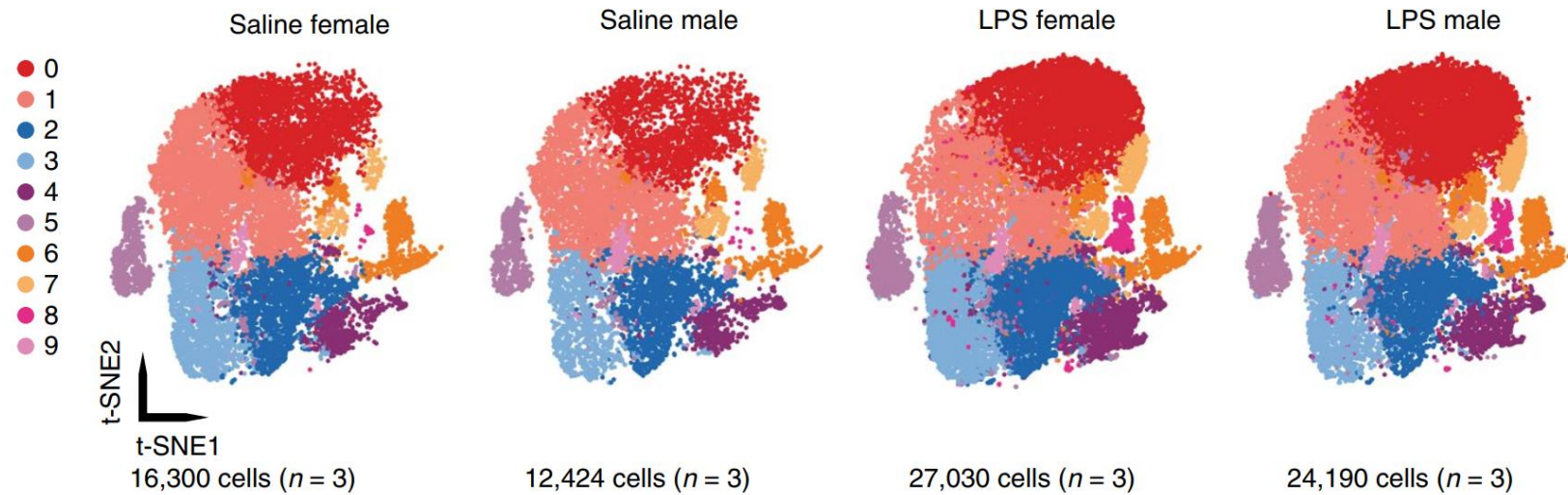


**Reproduction:**



# Fig.2b – clustered astrocytes

Paper



Reproduction

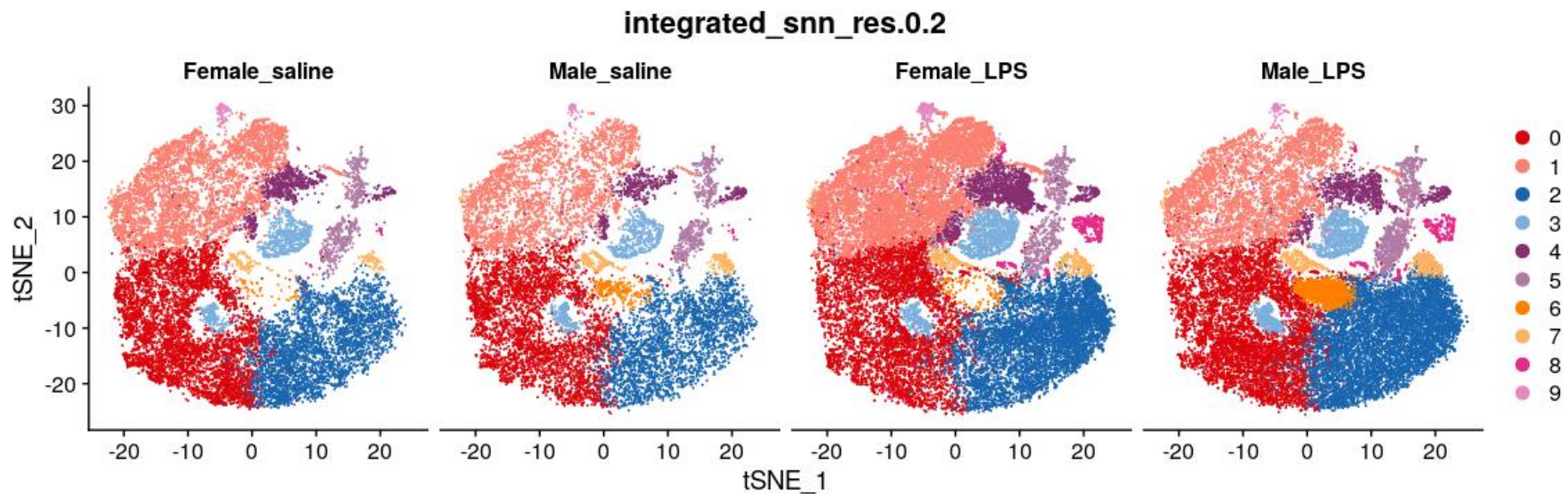


Fig.2c – gene expression heatmap

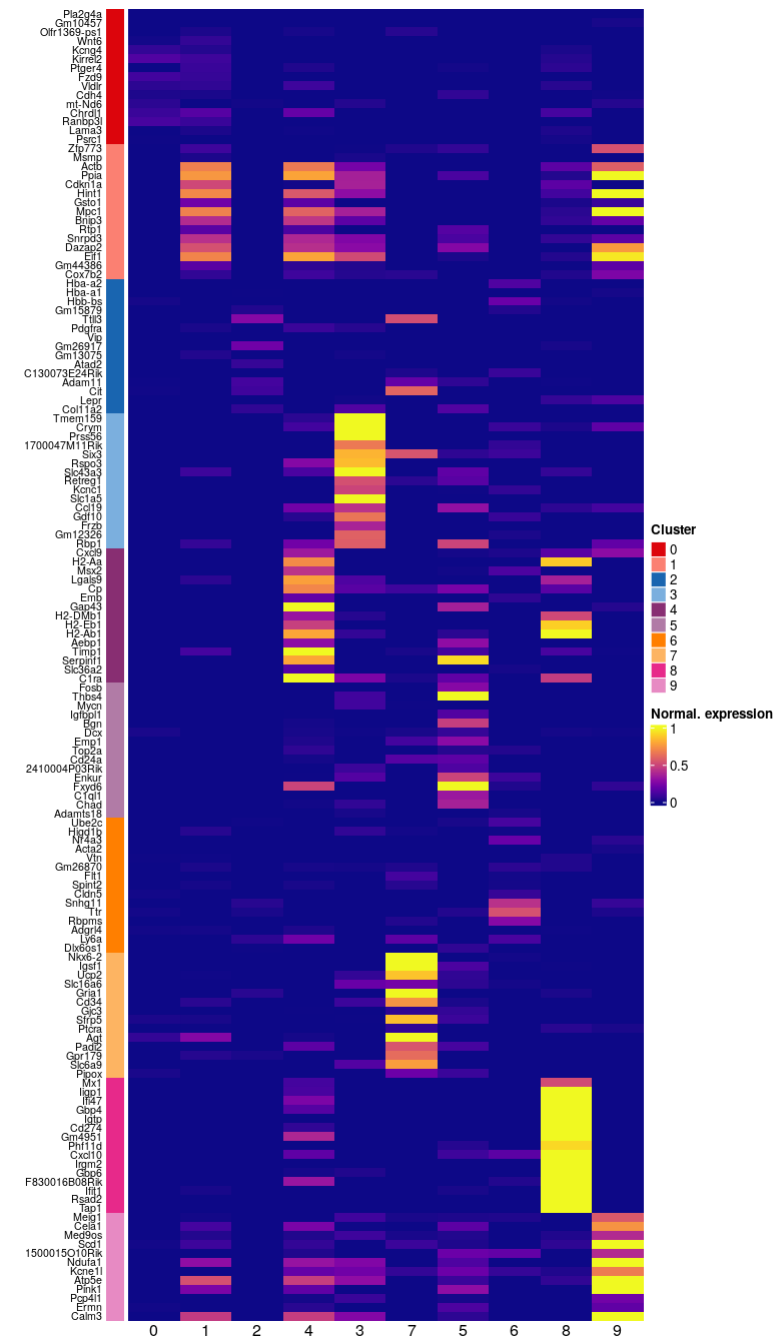
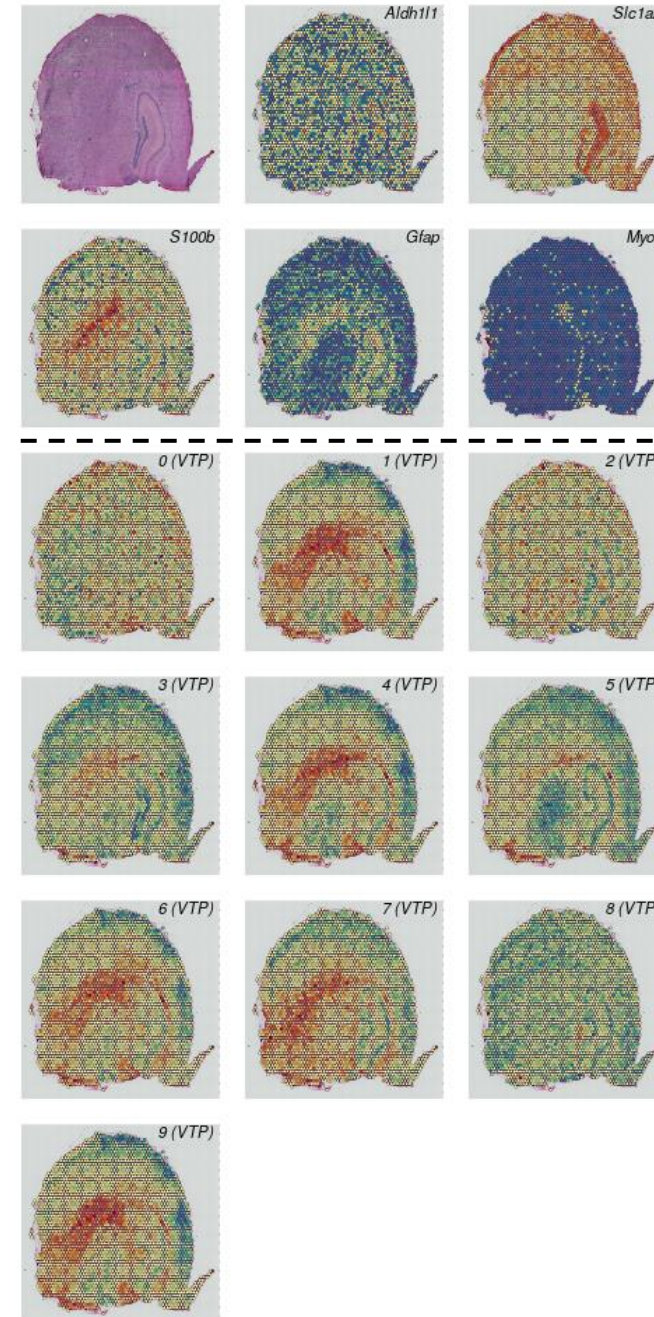




Fig.2d – Visium spatial transcriptomics plots



Astrocyte marker genes

Cluster 0-9 modules

Fig. 4 – LPS super-responder cluster that resides in the white matter (replication)

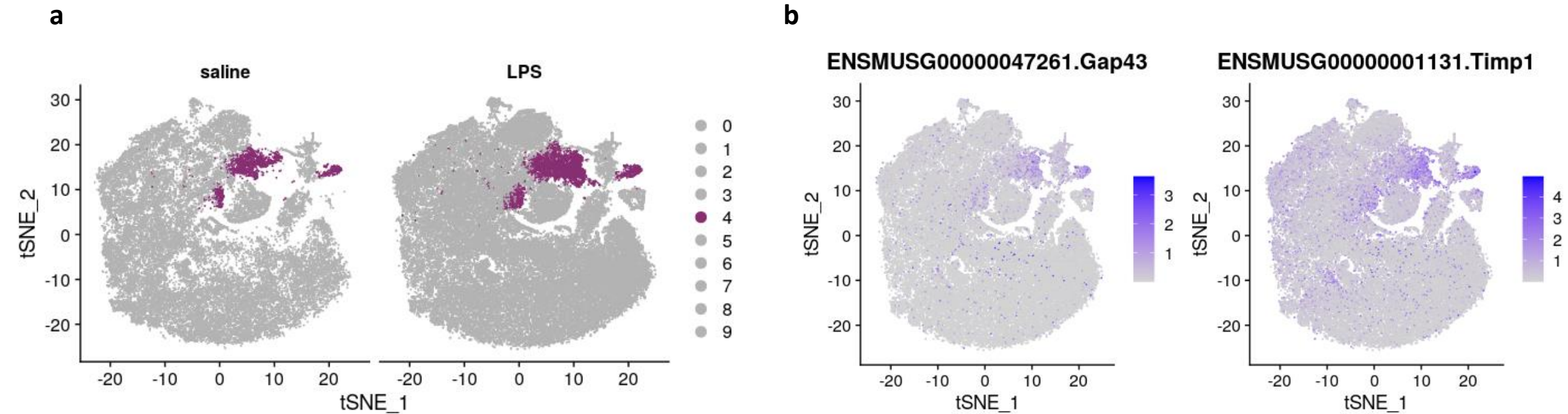


Fig. 4 – LPS super-responder cluster that resides in the white matter (replication)

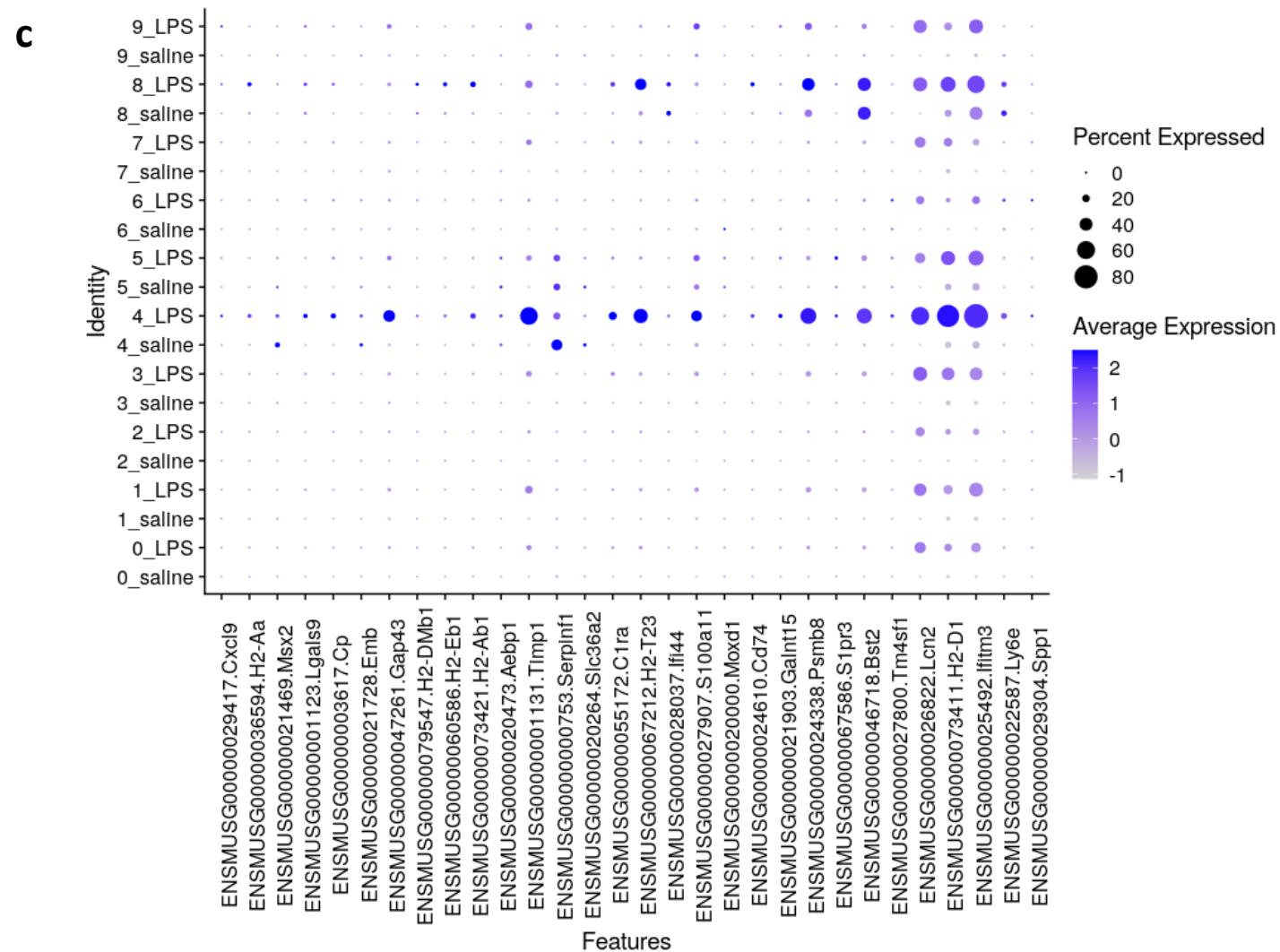
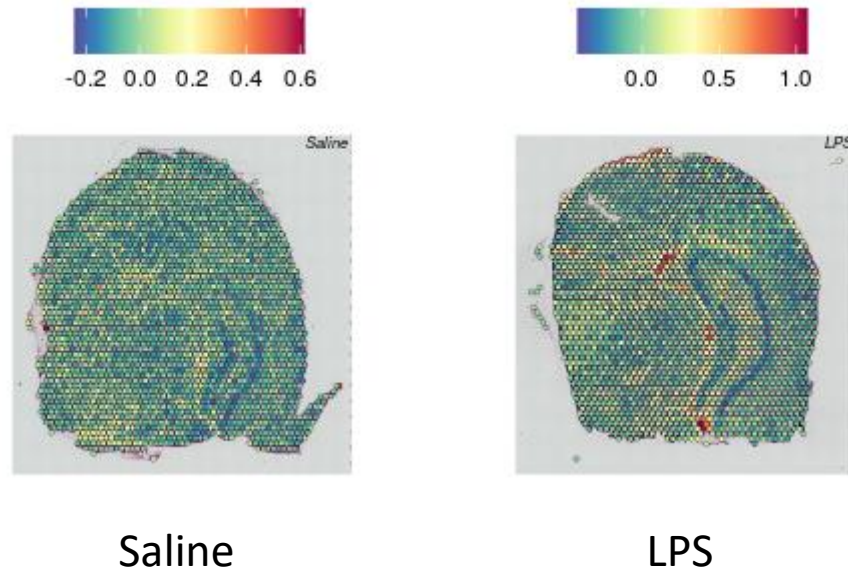




Fig. 4 – LPS super-responder cluster that resides in the white matter (replication)

d



f

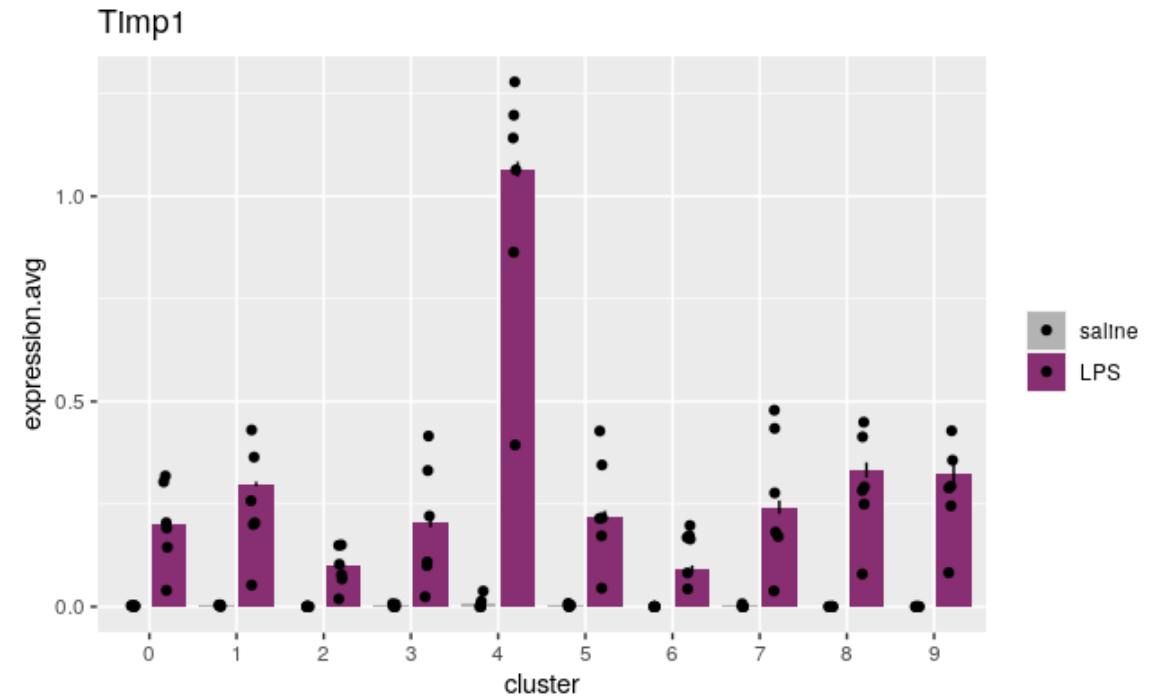


Fig. 5 – LPS super-responder cluster associated with the brain surface (replication)

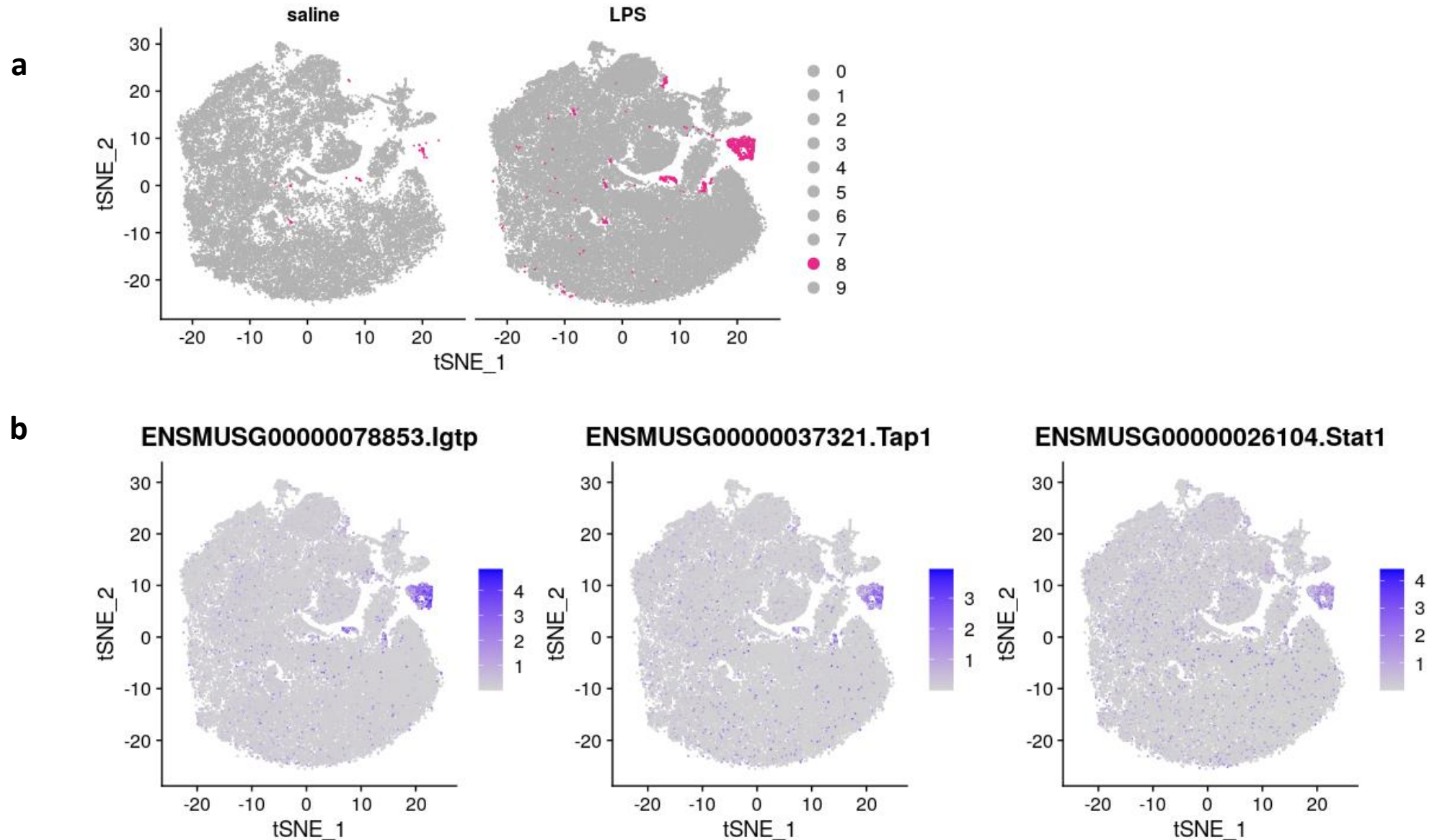


Fig. 4 – LPS super-responder cluster that resides in the white matter (replication)

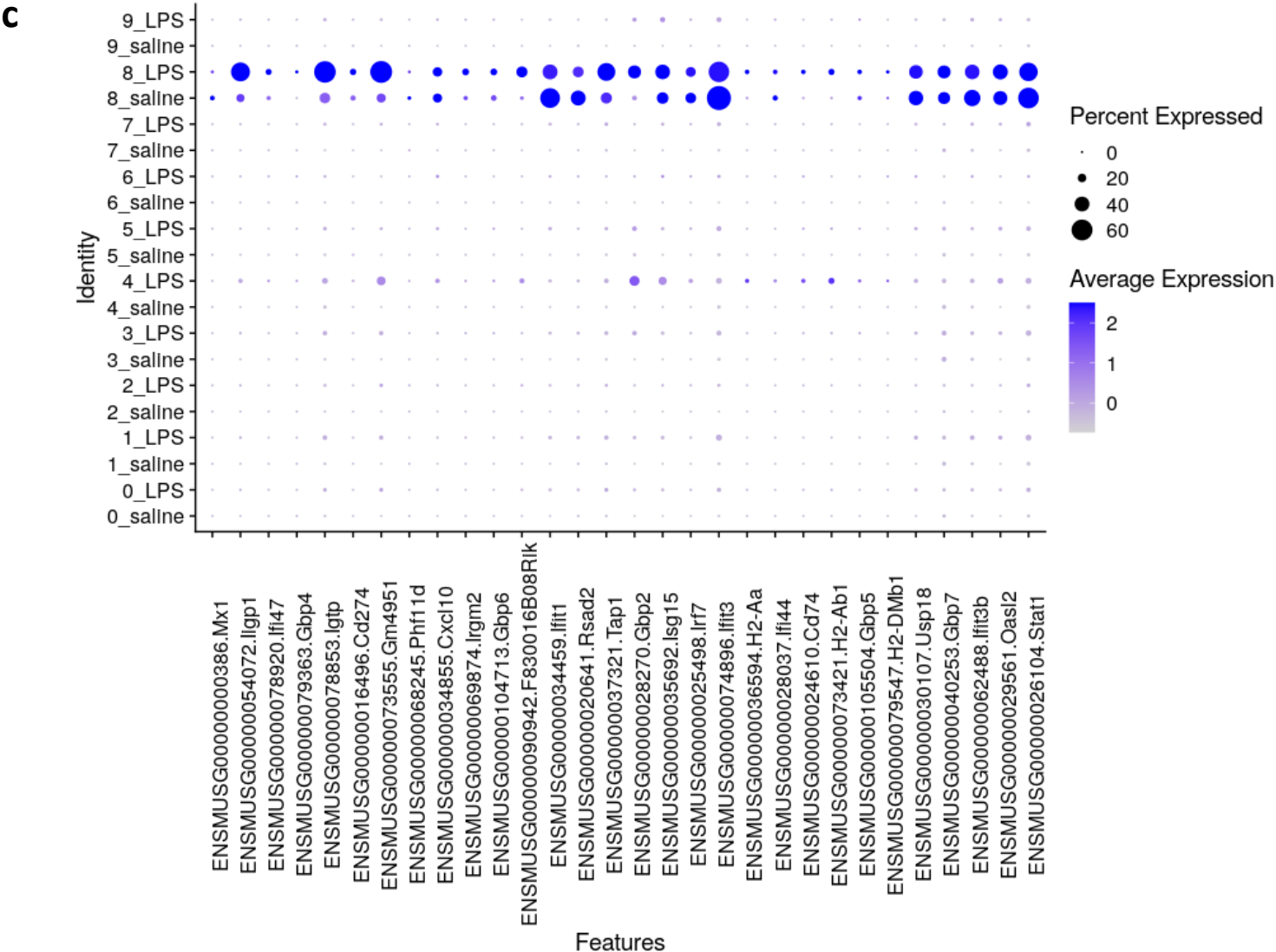
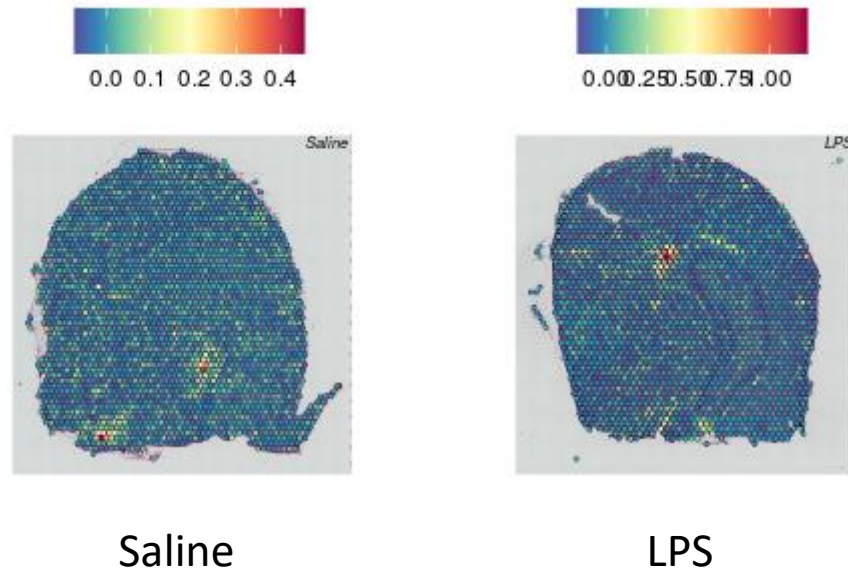


Fig. 4 – LPS super-responder cluster that resides in the white matter (replication)

d



f

