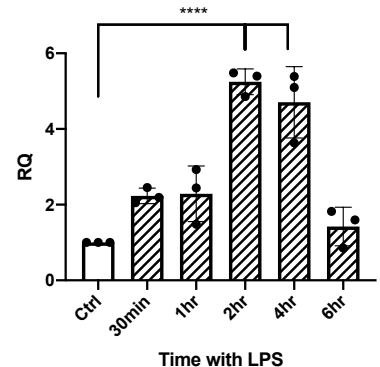
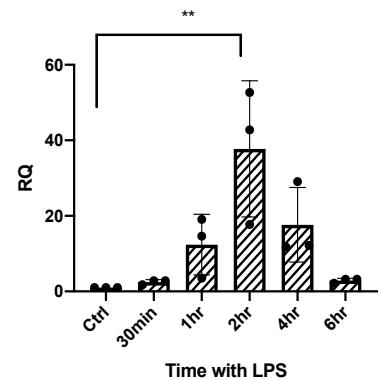


Figure. S1

NR4A1



NR4A2



NR4A3

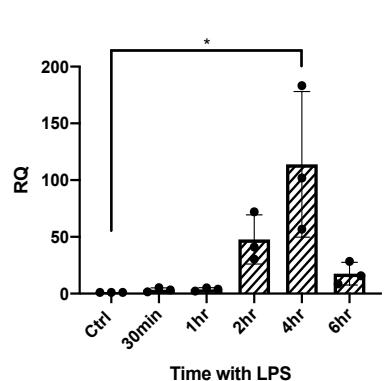
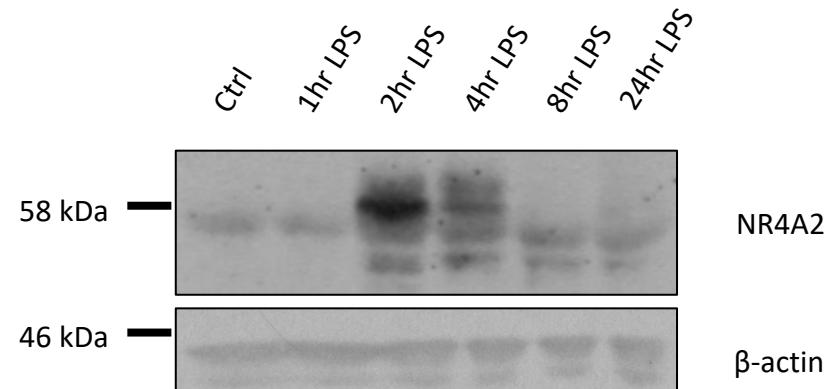
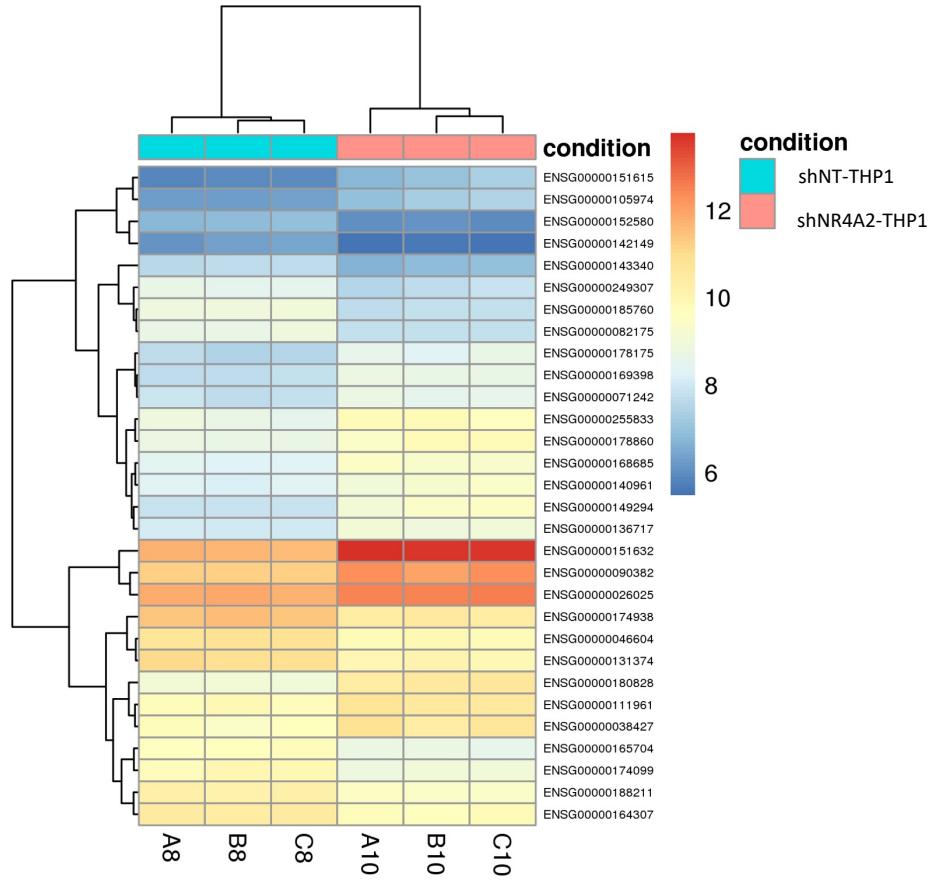
**D.**

Figure. S2

A.

shNR4A2



B.

shNR4A3

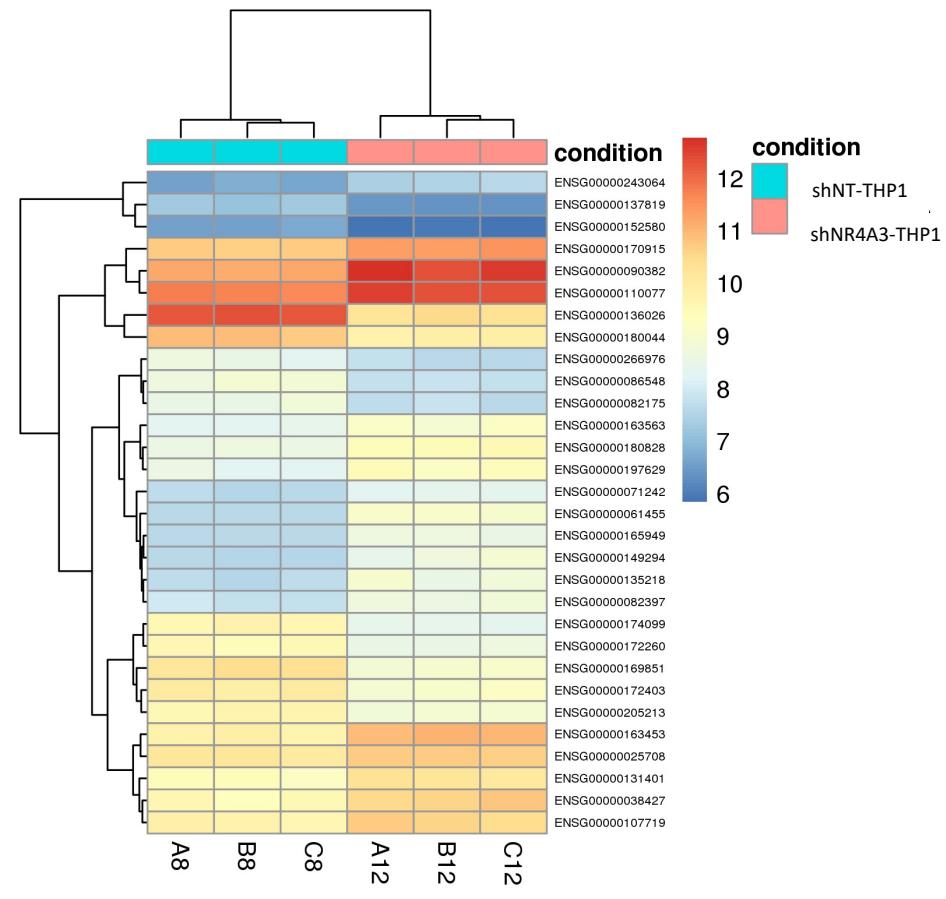
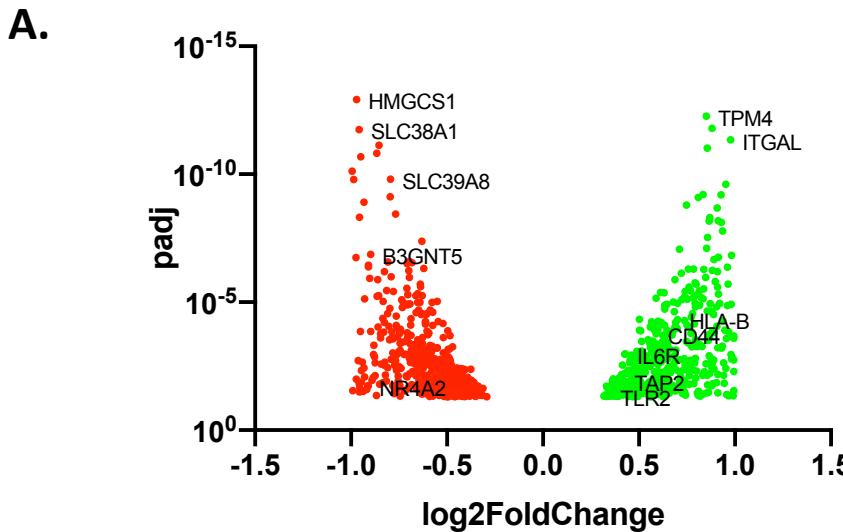


Figure. S3

shNR4A2



B.

shNR4A3

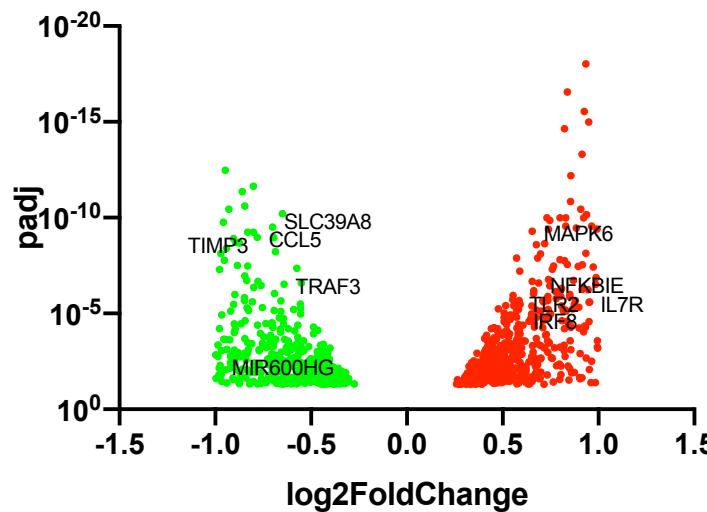
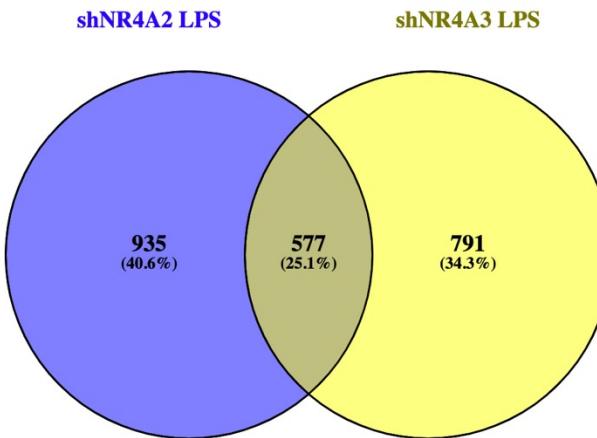


Figure. S4**A.**

4. shNR4A2 only regulated genes in the presence of LPS (935)
5. shNR4A3 only regulated genes in the presence of LPS (791)
6. shNR4A2 & shNR4A3 regulated genes in presence of LPS (577)

**B.**

	Cluster 4	Cluster 6	Cluster 5
GO Term	<ul style="list-style-type: none"> (i) Antigen processing and presentation of endogenous peptide antigen via MHC class I via ER pathway, TAP-independent (GO:0002486) (ii) Antigen processing and presentation of endogenous peptide antigen via MHC class I via ER pathway (GO:0002484) (iii) Antigen processing and presentation of endogenous peptide antigen via MHC class Ib (GO:0002476) 	<ul style="list-style-type: none"> (i) T cell activation via T cell receptor contact with antigen bound to MHC molecule on antigen presenting cell (GO:0002291) (ii) Regulation of guanyl-nucleotide exchange factor activity (GO:1905097) (iii) Regulation of chemokine-mediated signaling pathway (GO:0070099) 	<ul style="list-style-type: none"> (i) Positive regulation of monocyte aggregation (GO:1900625) (ii) p38MAPK cascade (GO:0038066) (iii) Cellular response to interferon-alpha (GO:0035457)
Genes	<ul style="list-style-type: none"> (i) HLA-C, HLA-E, HLA-B, HLA-F, HLA-A, HLA-G, HLA-H (ii) HLA-C, HLA-E, HLA-B, HLA-F, HLA-A, HLA-G, HLA-H (iii) HLA-E, HLA-F, HLA-G, HLA-H, TAP2 	<ul style="list-style-type: none"> (i) ITGAL, FGL2, ICAM1 (ii) GPSM1, MET, CCPG1, PBXIP1 (iii) EDN1, ROBO1, PADI2, SH2B3 	<ul style="list-style-type: none"> (i) NR4A3, HAS2, AL133330 (ii) ZFP36, ZFP36L1, LGALS9 (iii) IFIT2, OAS1, IFIT3, AXL, GAS6PHYSIP

Figure. S5

7. Genes that appear to be equally regulated by shNR4A2 and shNR4A3 in both the basal and stimulated state (276)

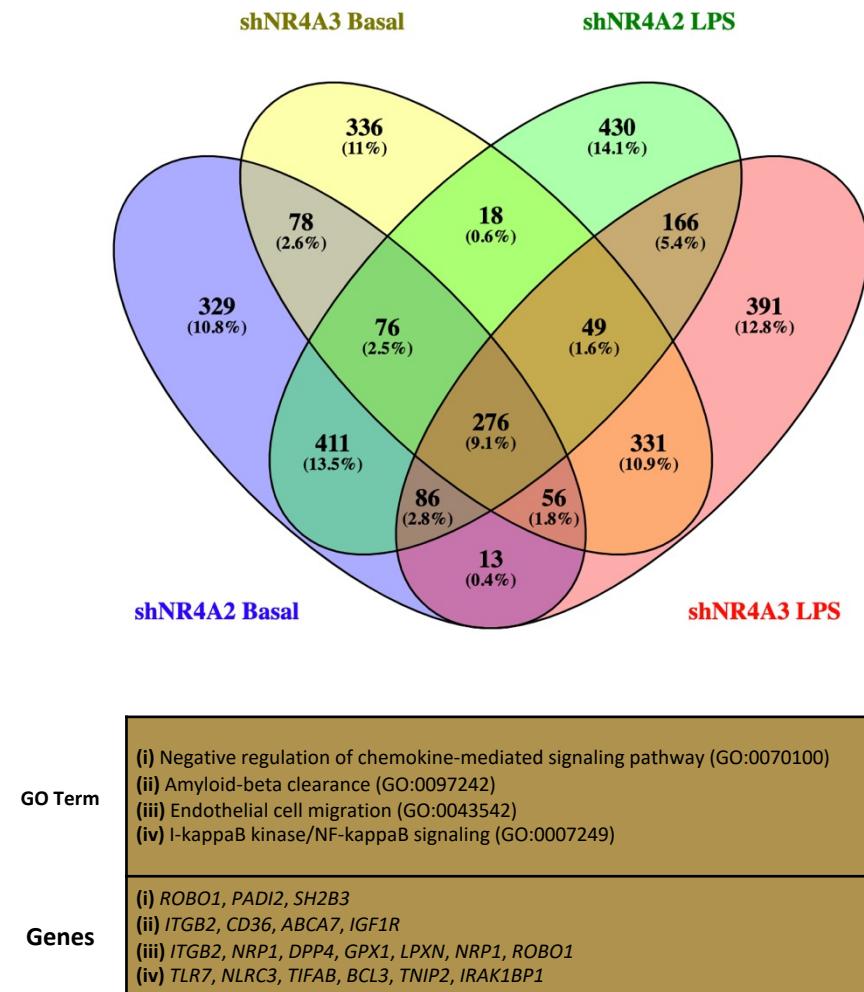


Figure. S6**Fibroblast activation genes GO:0072537**

shNR4A2 selective – Cluster 1

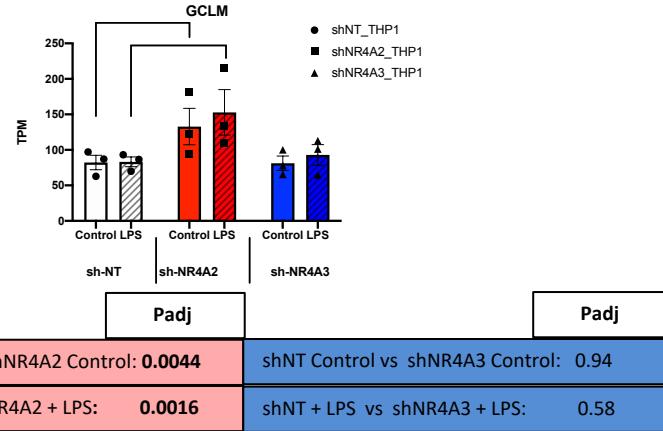
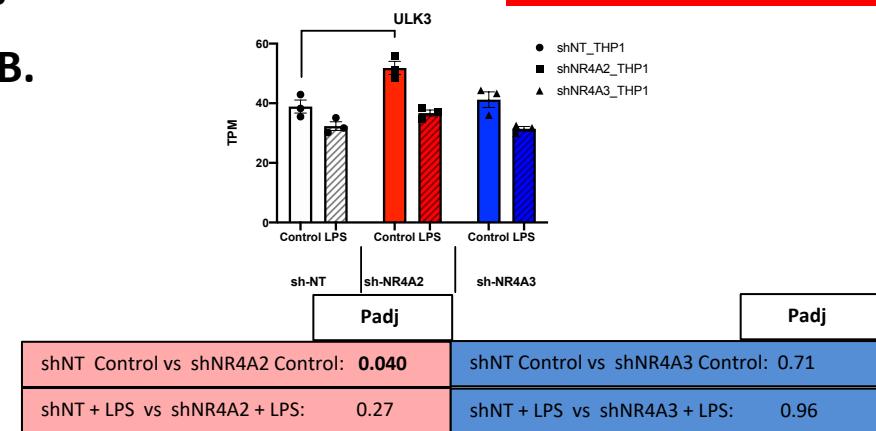
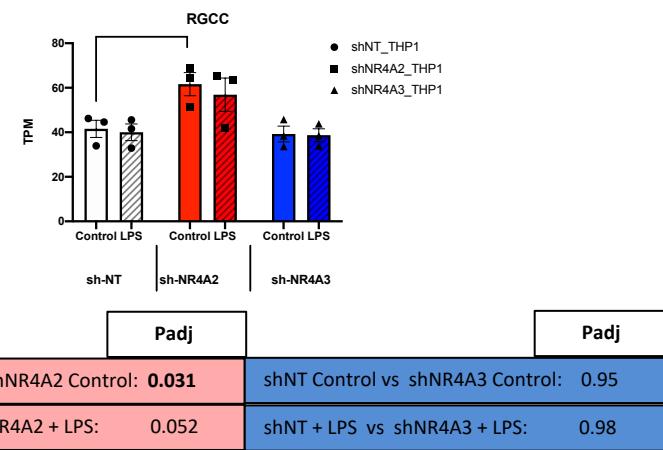
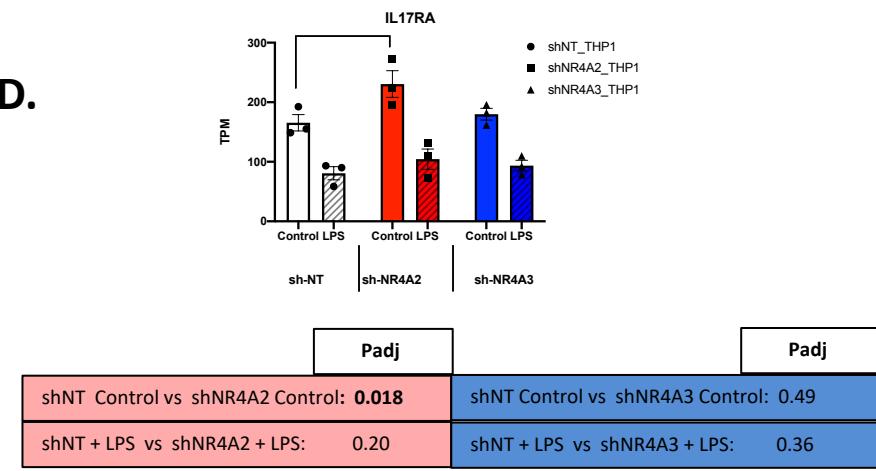
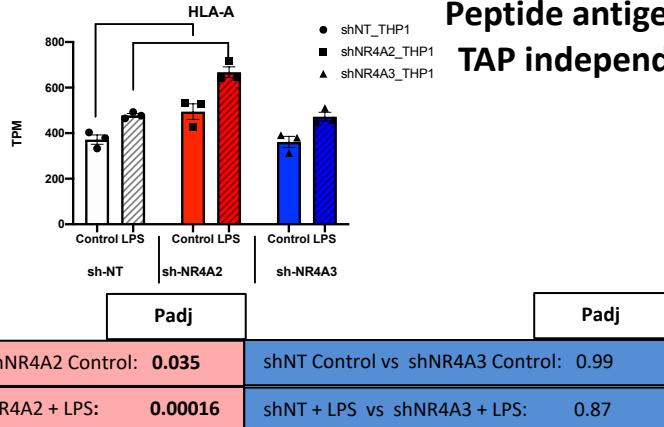
A.**B.****C.****D.**

Figure. S7

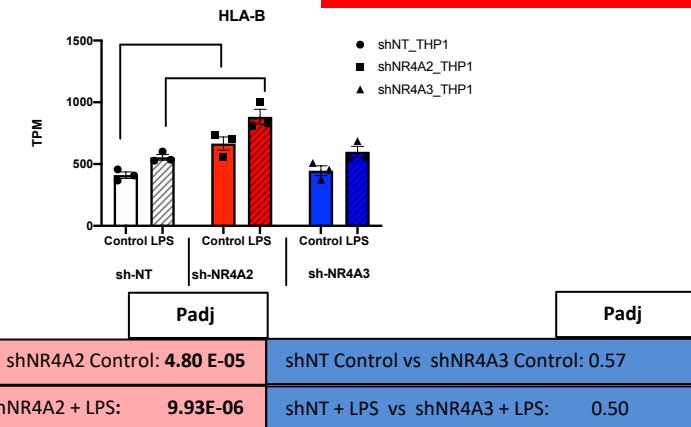
**Antigen processing and presentation of exogenous
Peptide antigen via MHC class I,
TAP independent GO:0002480**

shNR4A2 selective – Cluster 1

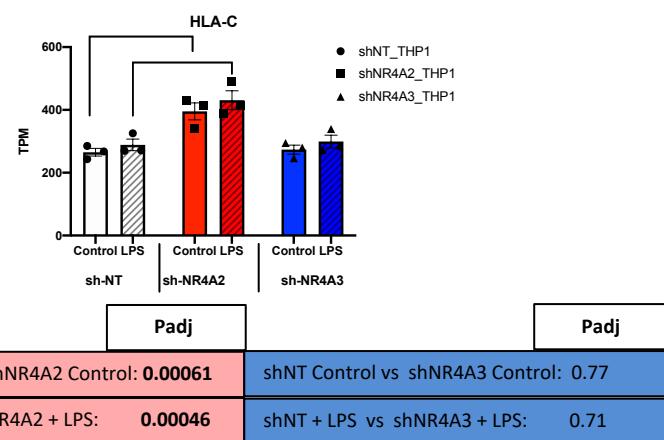
A.



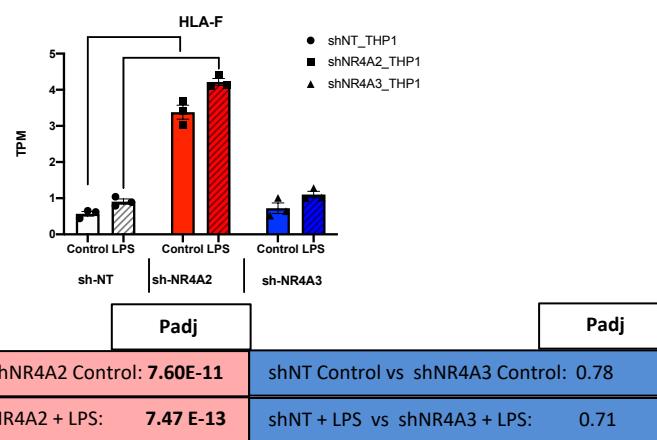
B.



C.



D.



E.

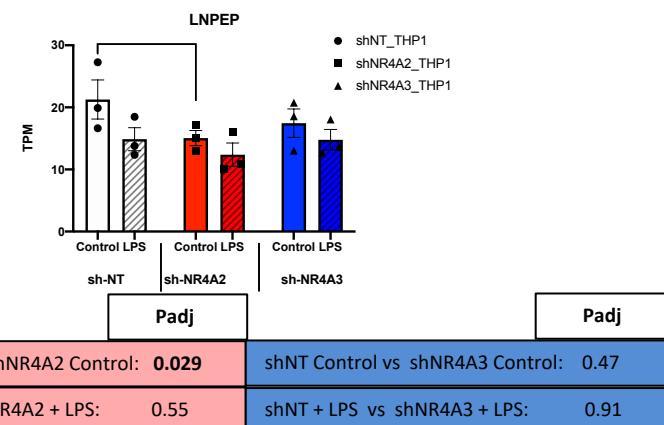
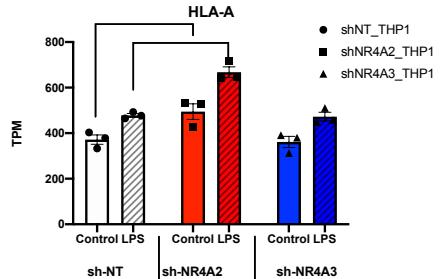
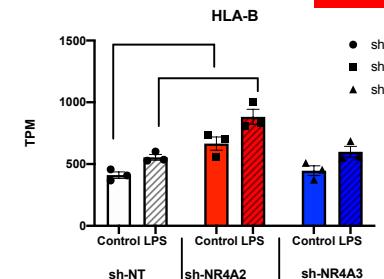
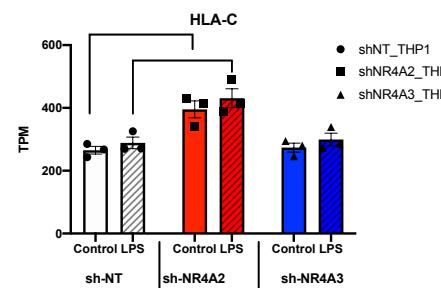
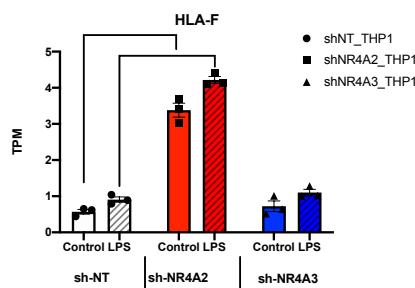
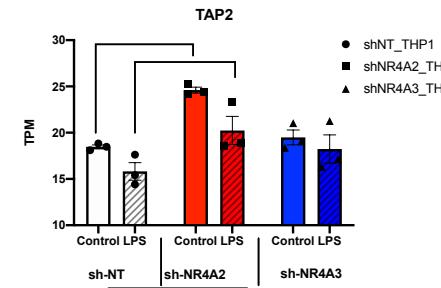
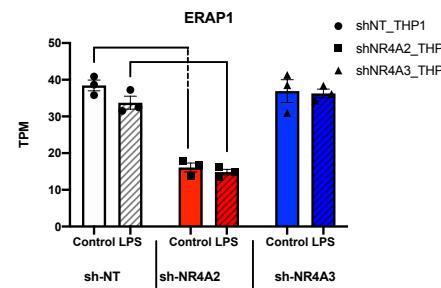


Figure. S8

Antigen processing and presentation of endogenous peptide antigen GO:0002483

shNR4A2 selective – Cluster 1

A.**B.****C.****D.****E.****F.**

shNT Control vs shNR4A2 Control: **0.012**
shNT + LPS vs shNR4A2 + LPS: **0.018**

Padj

shNT Control vs shNR4A3 Control: **0.66**
shNT + LPS vs shNR4A3 + LPS: **0.21**

Padj

shNT Control vs shNR4A2 Control: **8.34E-21**
shNT + LPS vs shNR4A2 + LPS: **1.82E-20**

Padj

shNT Control vs shNR4A3 Control: **0.98**
shNT + LPS vs shNR4A3 + LPS: **0.36**

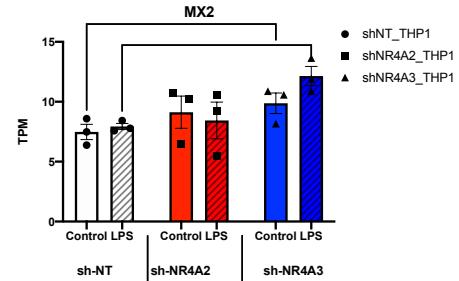
Padj

Response to interferon alpha GO:035455

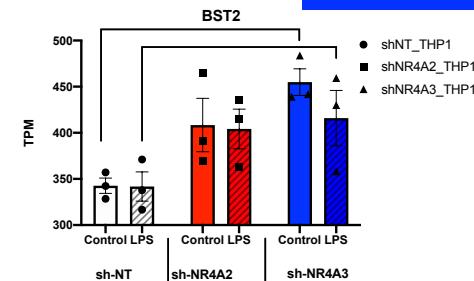
shNR4A3 selective – Cluster 2

Figure. S9

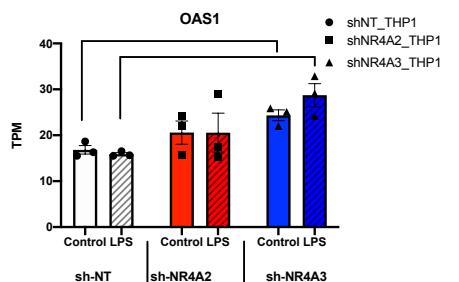
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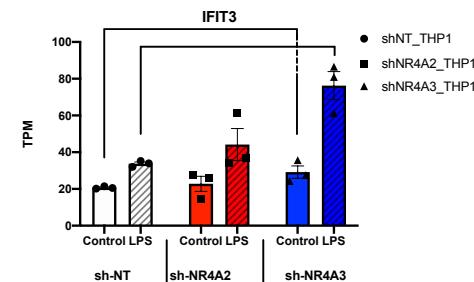
B.



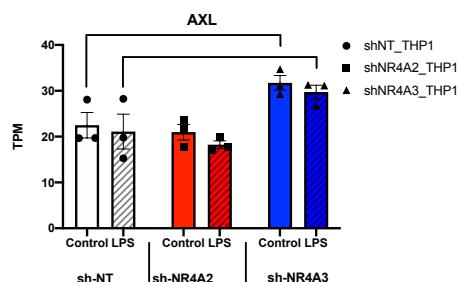
C.



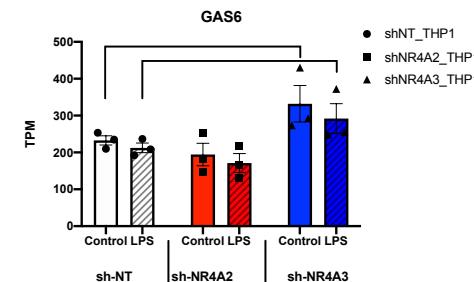
D.



E.



F.



shNT Control vs shNR4A2 Control: 0.829

Padj

shNT + LPS vs shNR4A2 + LPS: 0.747

Padj

shNT Control vs shNR4A3 Control: 0.0076

shNT + LPS vs shNR4A3 + LPS: 0.039

shNT Control vs shNR4A2 Control: 0.439

shNT Control vs shNR4A3 Control: 0.0068

shNT + LPS vs shNR4A2 + LPS: 0.494

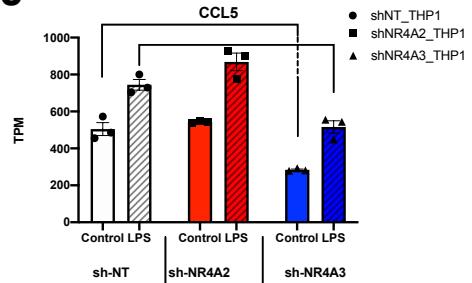
shNT + LPS vs shNR4A3 + LPS: 0.026

Protein Kinase B Signaling GO:0043491

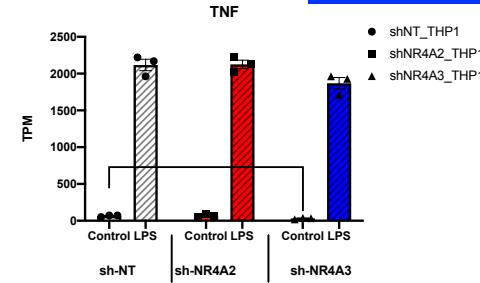
shNR4A3 selective – Cluster 2

Figure. S10

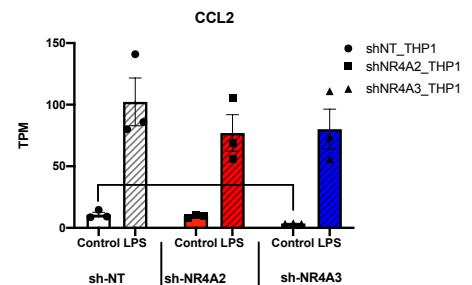
A.



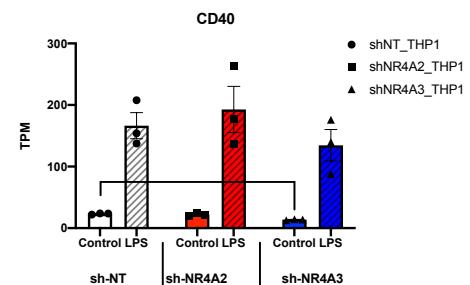
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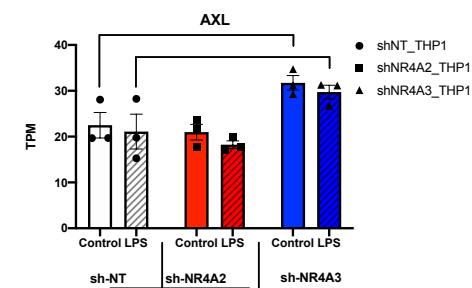
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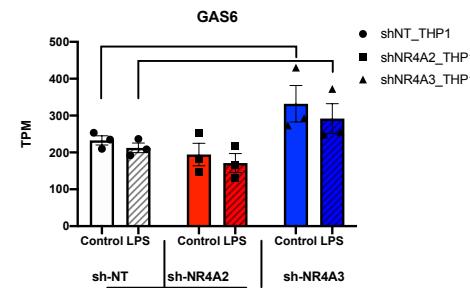
D.



E.



F.



shNT Control vs shNR4A2 Control: 0.83

shNT + LPS vs shNR4A2 + LPS: 0.75

Padj

Padj

shNT Control vs shNR4A3 Control: 0.0076

shNT + LPS vs shNR4A3 + LPS: 0.039

shNT Control vs shNR4A2 Control: 0.44

shNT + LPS vs shNR4A2 + LPS: 0.49

Padj

Padj

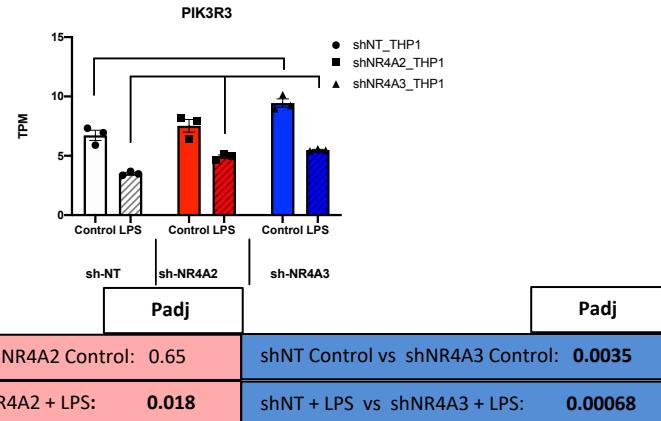
shNT Control vs shNR4A3 Control: 0.0068

shNT + LPS vs shNR4A3 + LPS: 0.026

Protein Kinase B Signaling GO:0043491

Figure. S10

G.



H.

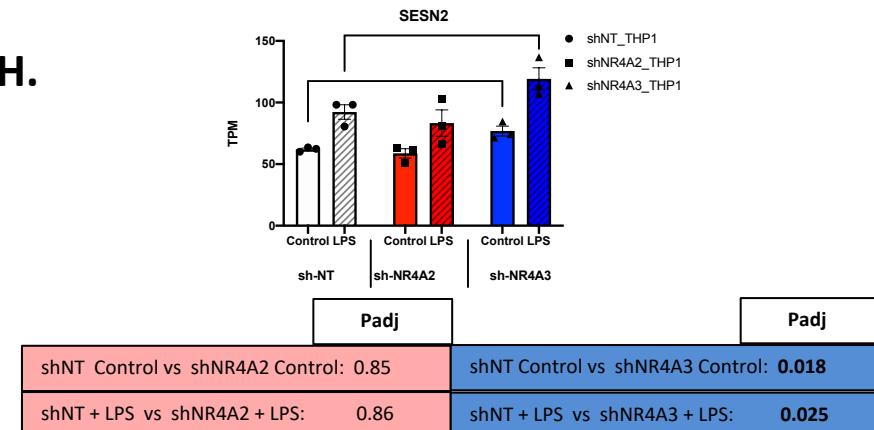
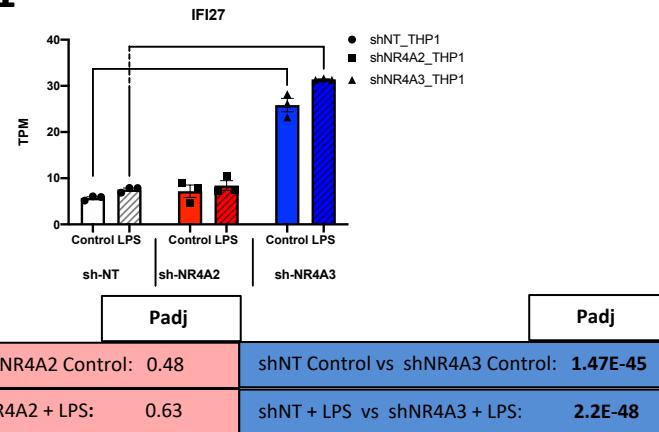


Figure. S11

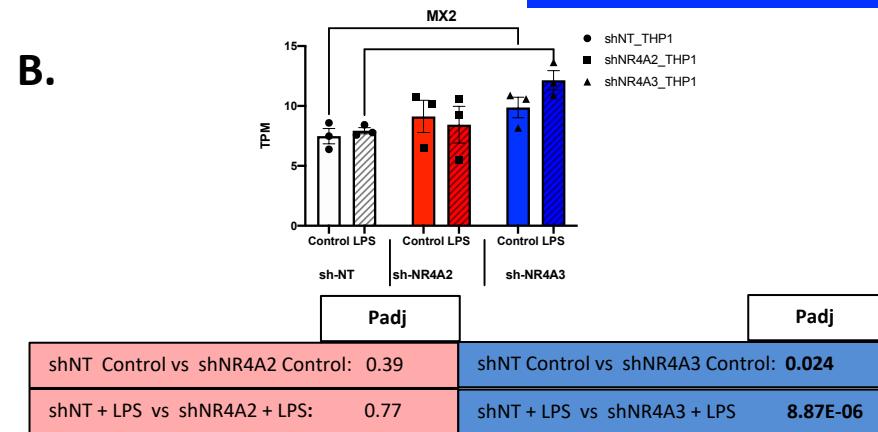
Type 1 interferon signaling pathway GO:0060337

shNR4A3 selective – Cluster 2

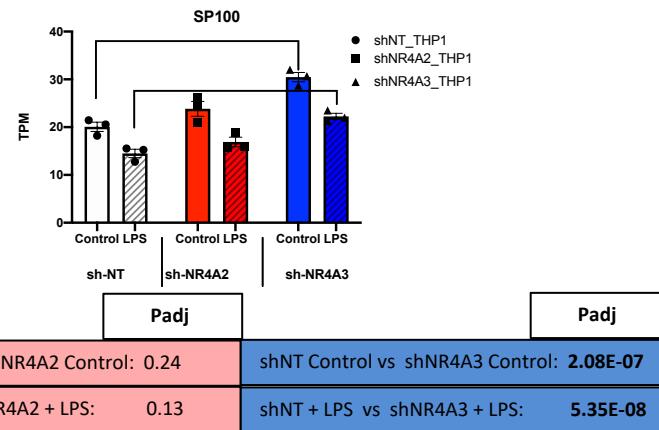
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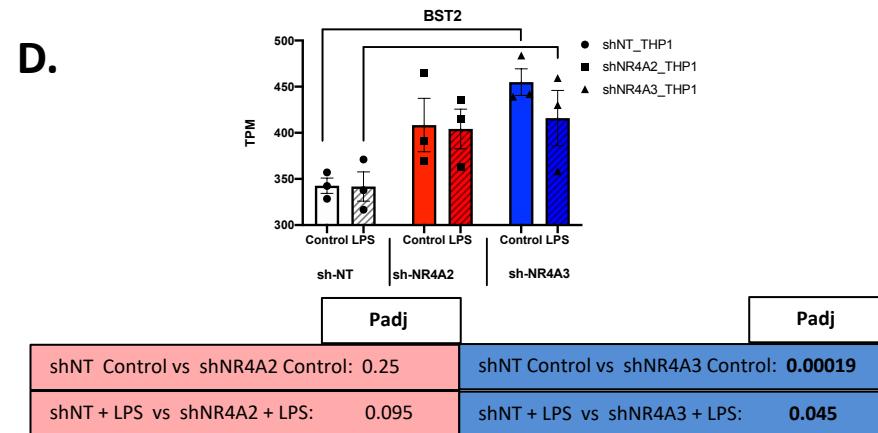
B.



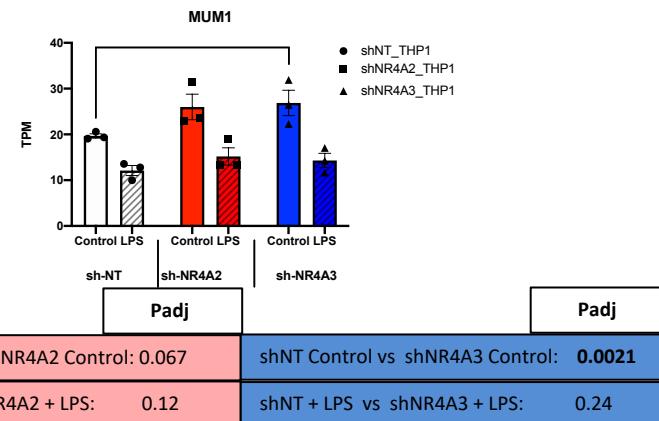
C.



D.



E.



F.

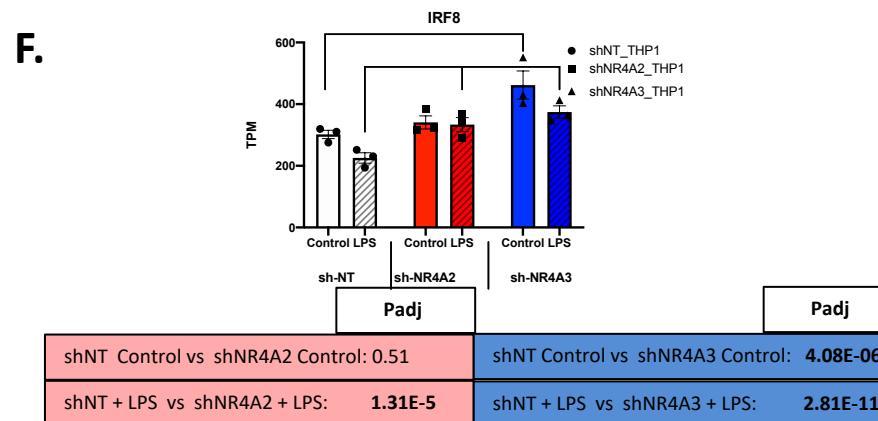
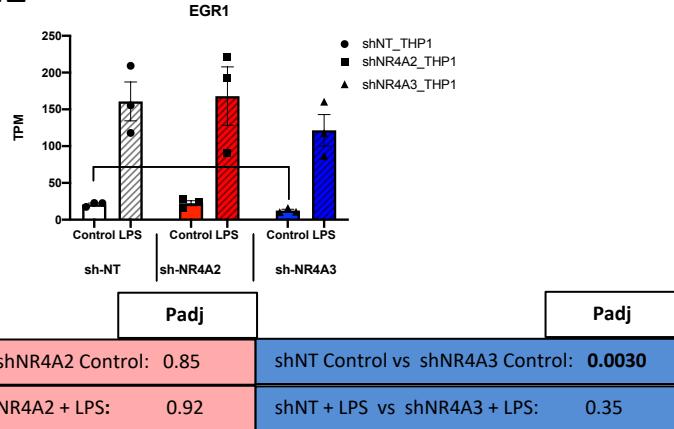


Figure. S11

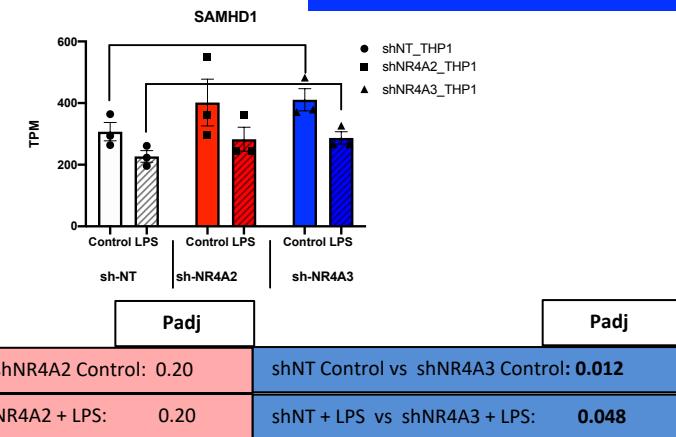
Type 1 interferon signaling pathway GO:0060337

shNR4A3 selective – Cluster 2

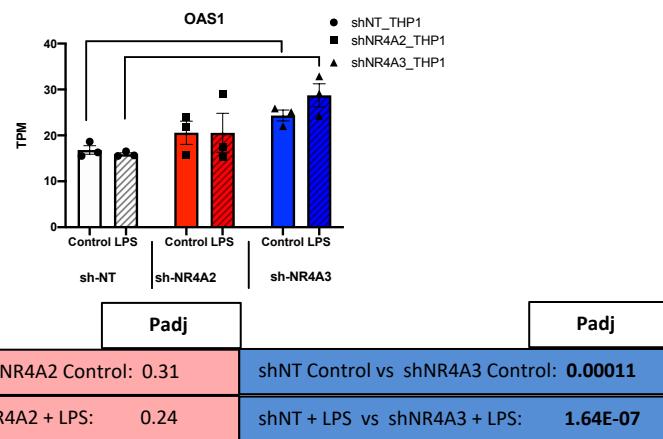
G.



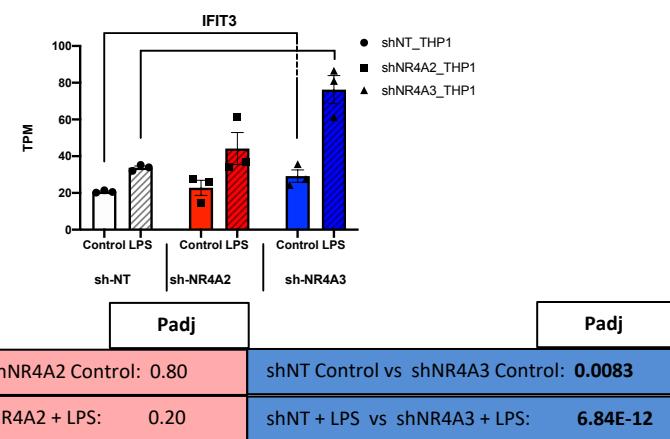
H.



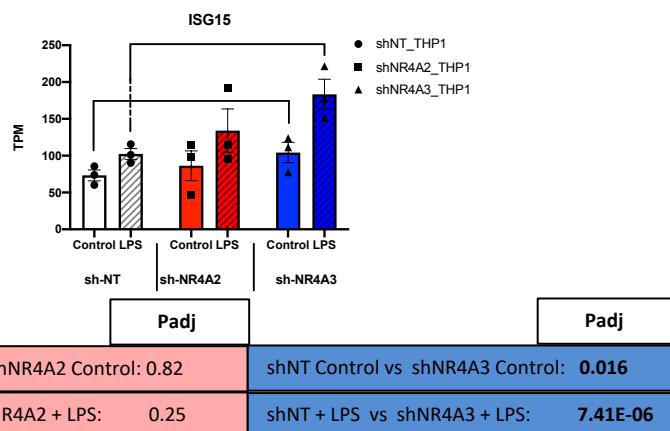
I.



J.



K.



L.

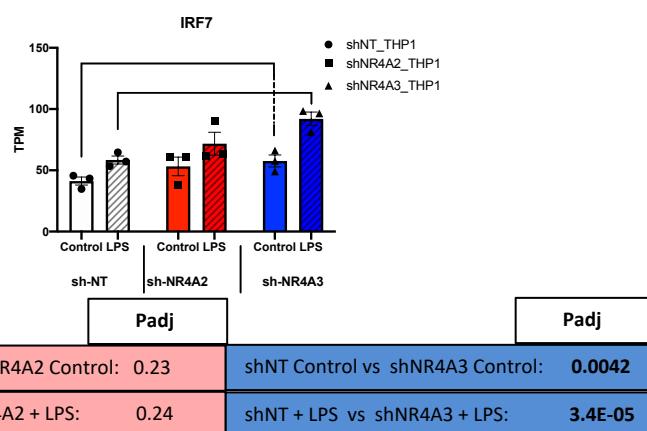


Figure. S11**Type 1 interferon signaling pathway GO:0060337**

shNR4A3 selective – Cluster 2

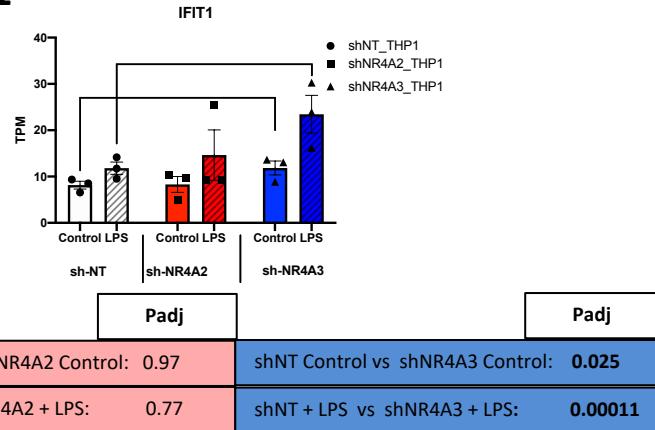
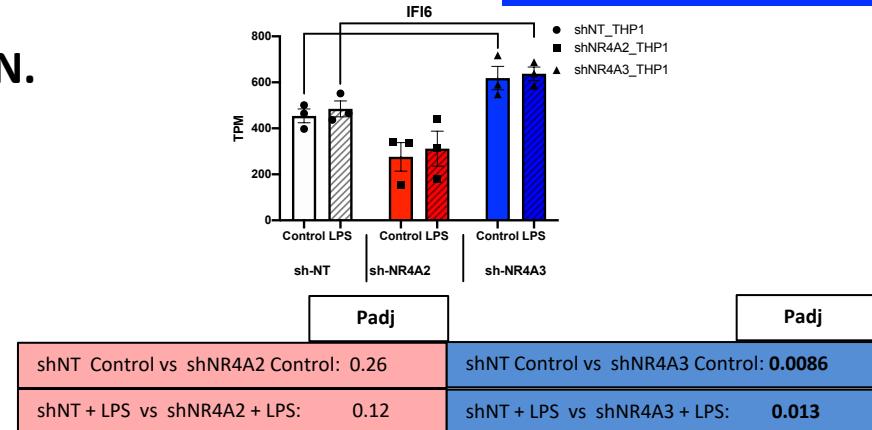
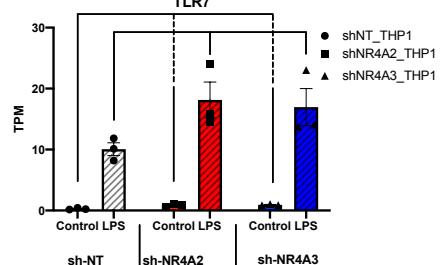
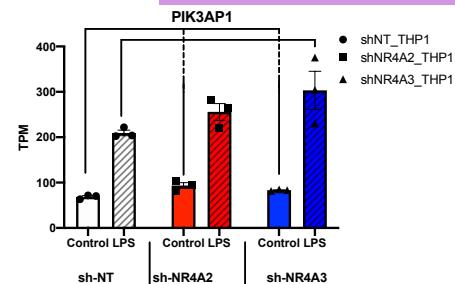
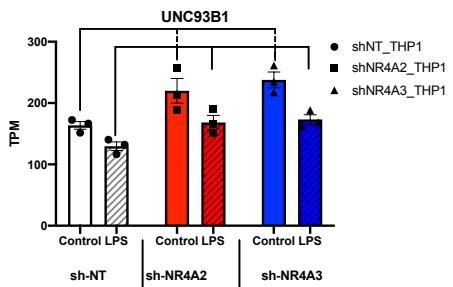
M.**N.**

Figure. S12

Toll-like receptor 7 signaling pathway GO:0034154

shNR4A2/A3 sensitive – Cluster 3

A.**B.****C.**

Padj	Padj
shNT Control vs shNR4A2 Control: 0.042	shNT Control vs shNR4A3 Control: 0.013
shNT + LPS vs shNR4A2 + LPS: 0.0011	shNT + LPS vs shNR4A3 + LPS: 0.0034

Padj	Padj
shNT Control vs shNR4A2 Control: 0.0047	shNT Control vs shNR4A3 Control: 0.011
shNT + LPS vs shNR4A2 + LPS: 0.12	shNT + LPS vs shNR4A3 + LPS: 0.0094

Figure. S13**Toll-like receptor 2 signaling pathway GO:0034134**

shNR4A2/A3 sensitive – Cluster 3

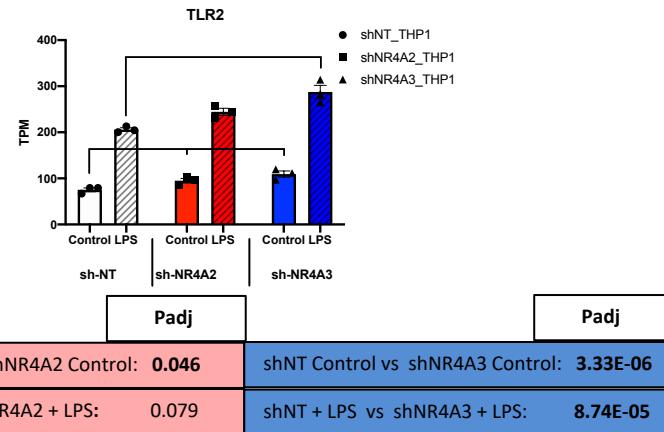
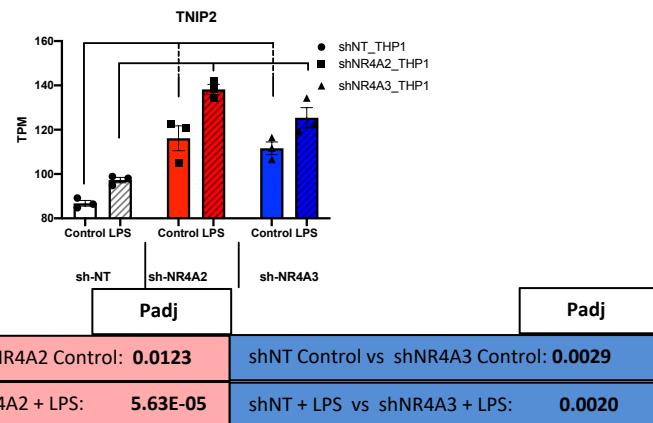
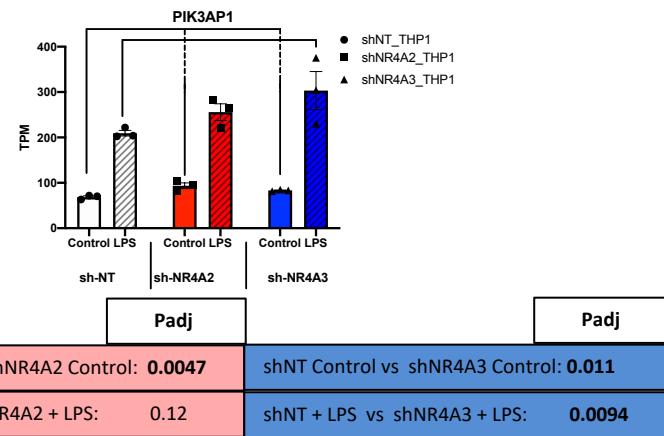
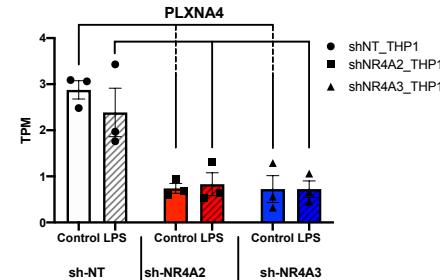
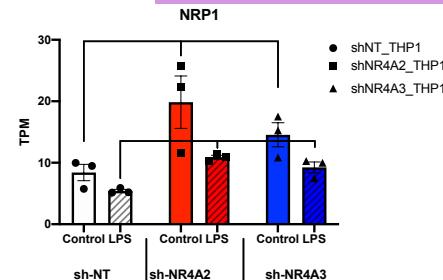
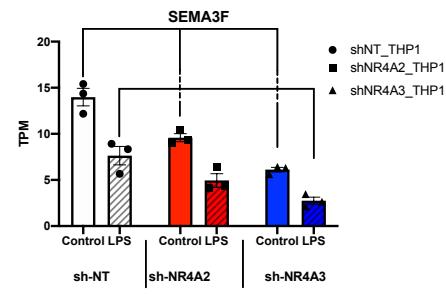
A.**B.****C.**

Figure. S14

Trigeminal Nerve Structure Gene GO:0021637

shNR4A2/A3 sensitive – Cluster 3

A.**B.****C.**

	Padj	Padj
shNT Control vs shNR4A2 Control:	1.39E-16	shNT Control vs shNR4A3 Control: 0.0054
shNT + LPS vs shNR4A2 + LPS:	0.00015	shNT + LPS vs shNR4A3 + LPS: 6.77E-07

	Padj	Padj
shNT Control vs shNR4A2 Control:	0.014	shNT Control vs shNR4A3 Control: 8.87E-10
shNT + LPS vs shNR4A2 + LPS:	0.081	shNT + LPS vs shNR4A3 + LPS: 6.74E-08