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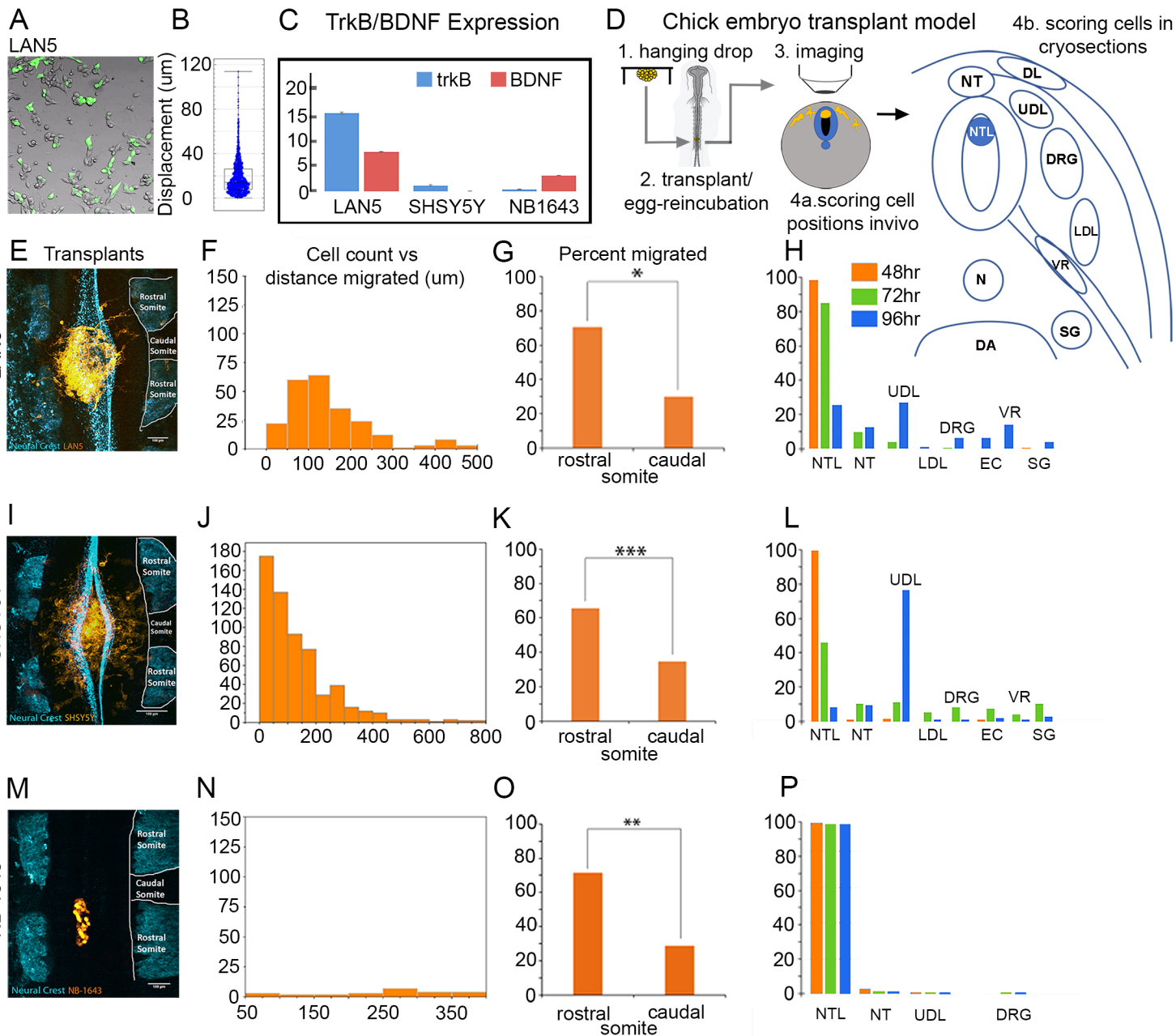
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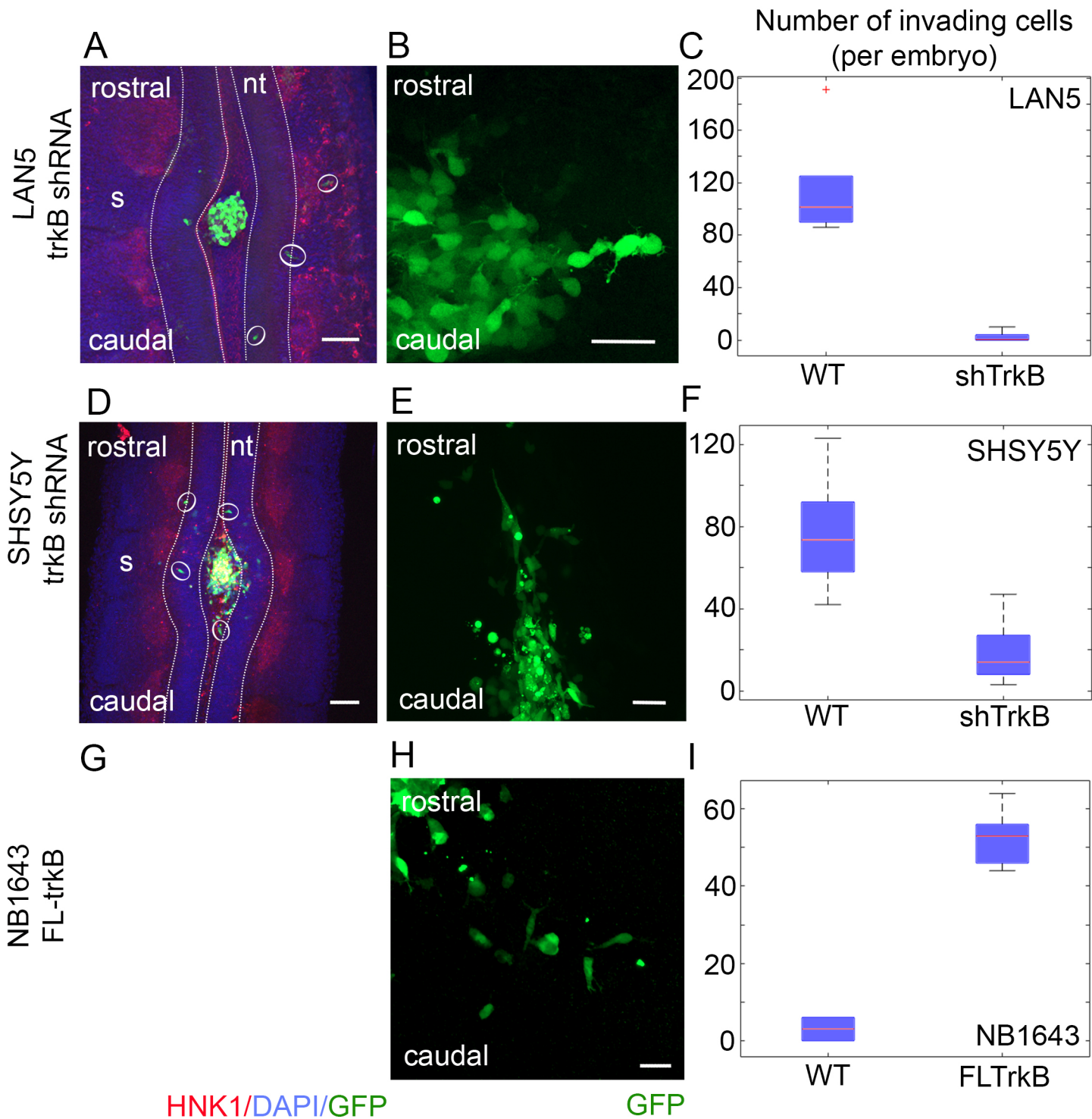
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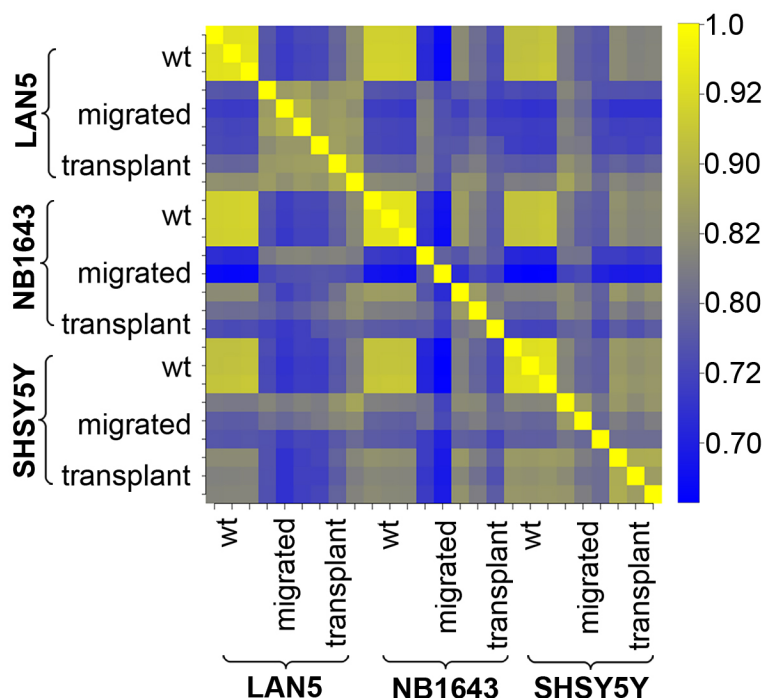




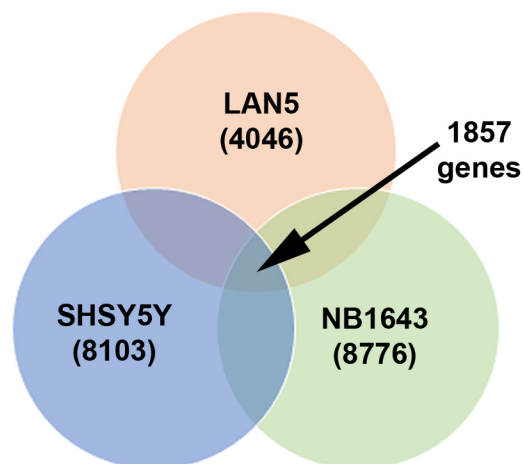




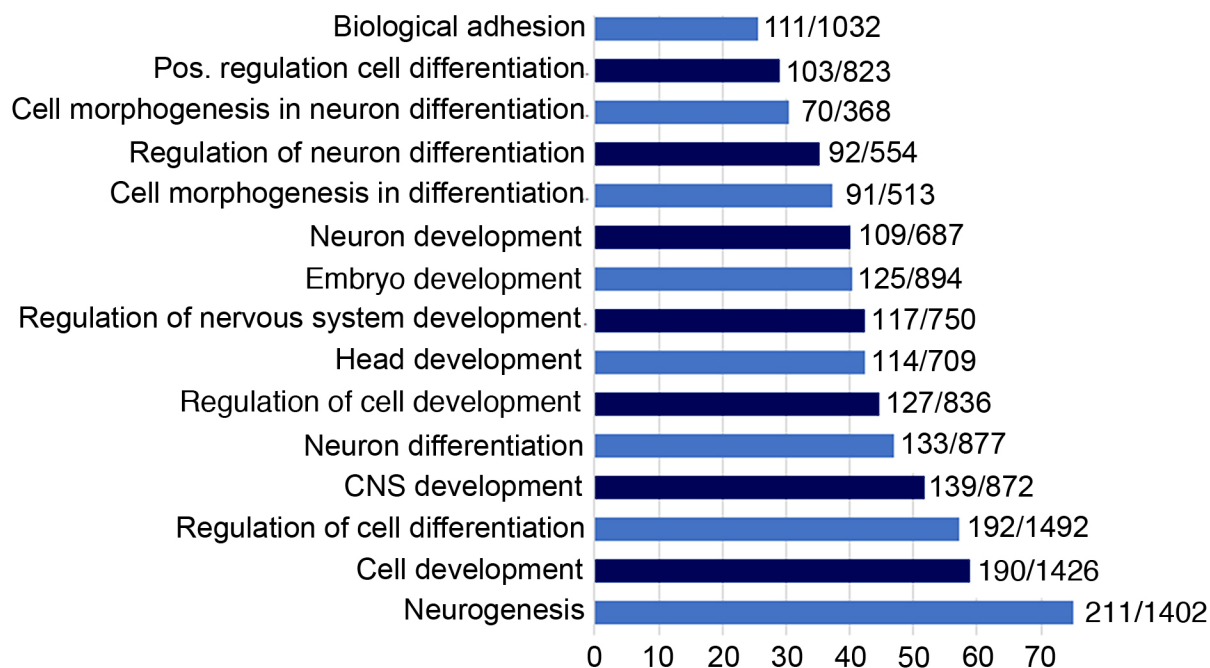
**A** Gene expression correlation matrix  
(after embryo transplantation)  
In ovo (migrated vs non-migrated) vs wildtype



**B** Changes in the number of genes after embryonic exposure



**C** Go Term Enrichment of neuroblastoma cells after embryonic exposure

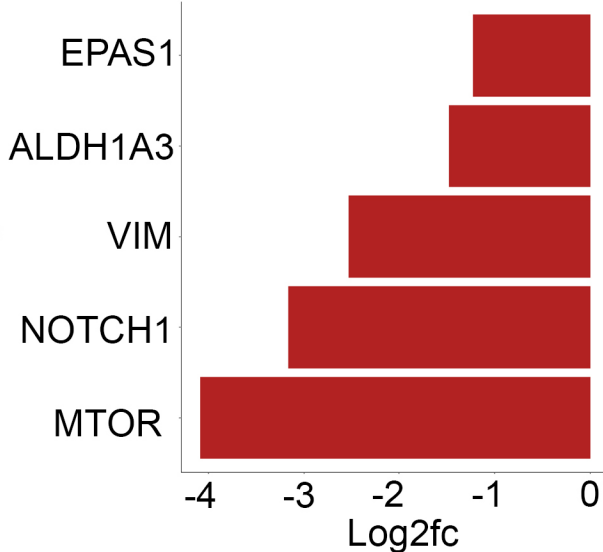




A

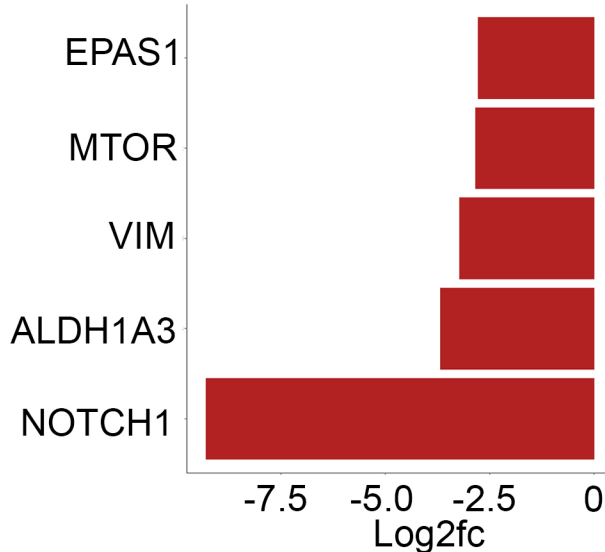
Tumor Initiating Cell Markers

LAN5



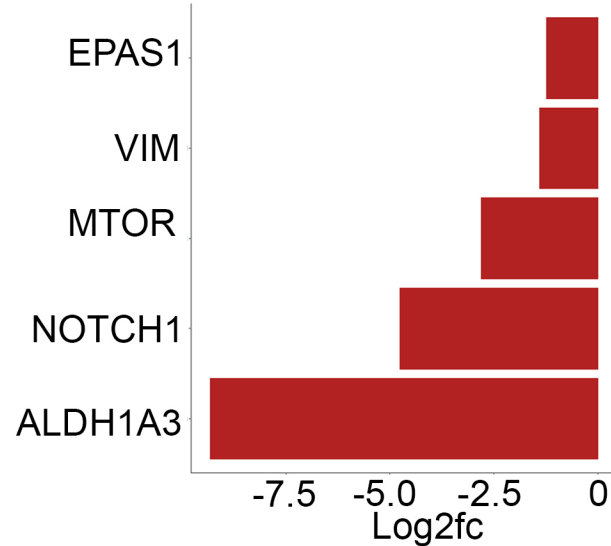
B

SHSY5Y



C

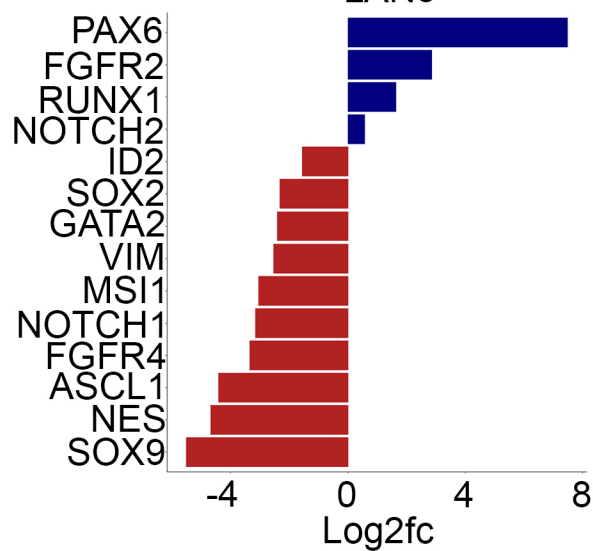
NB1643



D

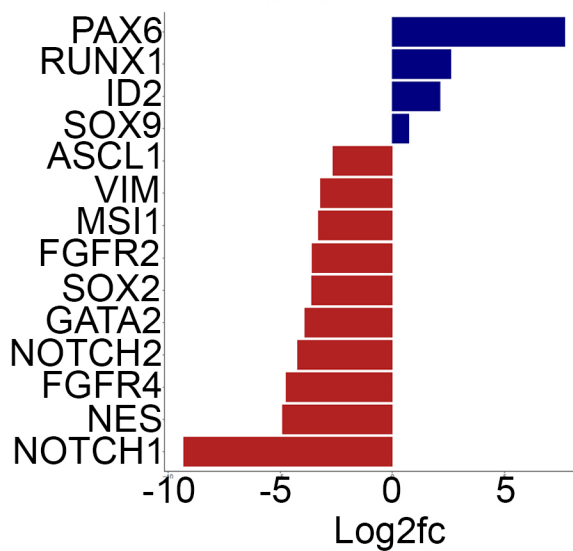
Stem Cell Markers

LAN5



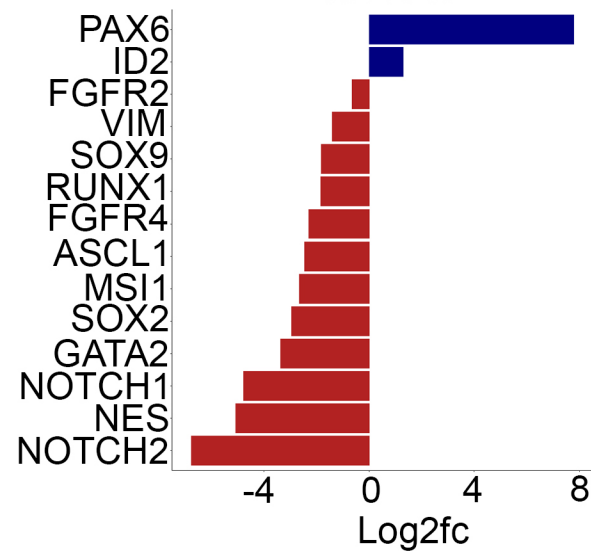
E

SHSY5Y



F

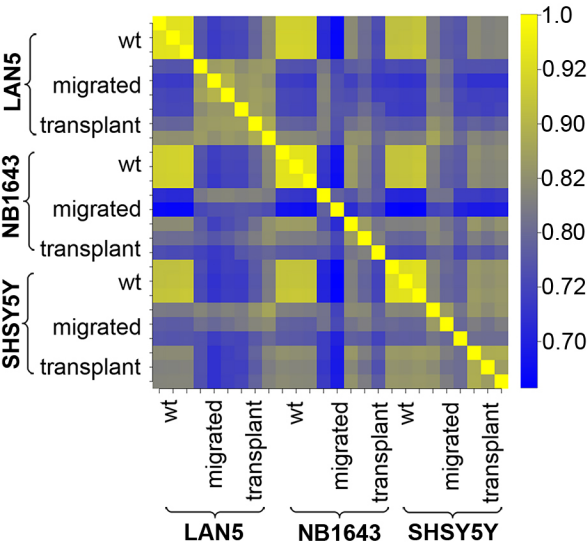
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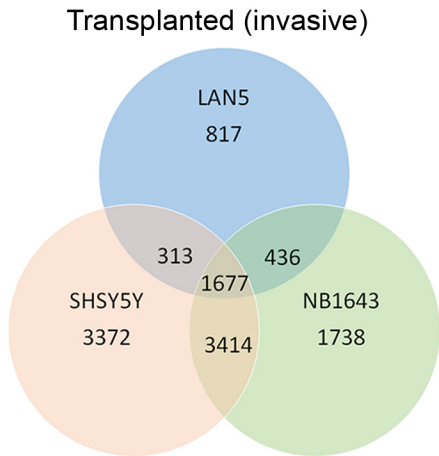
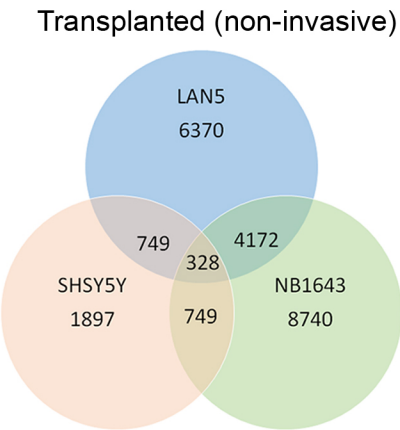
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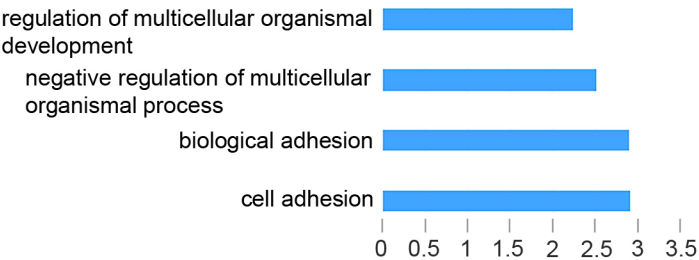
# A Gene expression correlation matrix (after embryo transplantation) In ovo (migrated vs non-migrated) vs wildtype



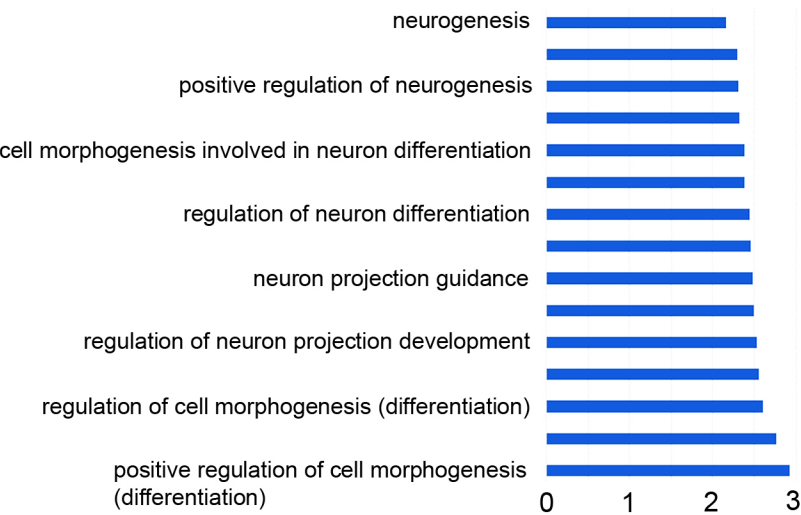
# B Changes in the number of genes after embryonic exposure



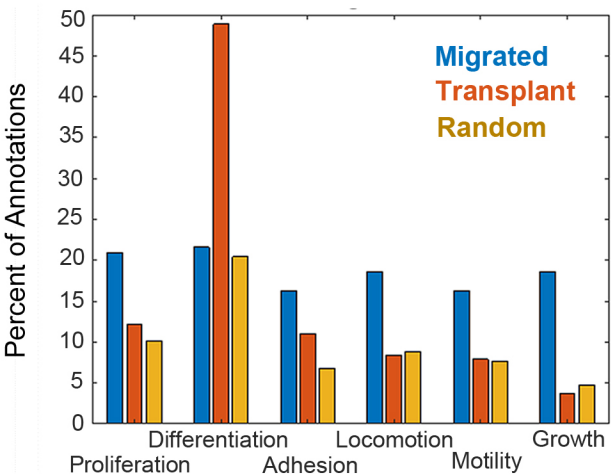
# C Go Term Fold Enrichment (328 genes)



# D Go Term Fold Enrichment (1677 genes)



# E Go Term Alignment



# A Model Input (Gene expression ON/OFF)

# B Model Output (Cell State ON/OFF)

# C Predicted Outcome

LAN5

	MDK	Alk	BDNF	trkA	trkB	NGF
WT						
Transplant						
Migrated						

	Diff	Apop	Angio	Prolif
Ss45				
Ss21				
Ss25				

Intermediate
Intermediate
Poor

SHSY5Y

	MDK	Alk	BDNF	trkA	trkB	NGF
WT						
Transplant						
Migrated						

	Diff	Apop	Angio	Prolif
Ss38				
Ss28				
Ss44				

Intermediate
Good
Poor

NB-1643

	MDK	Alk	BDNF	trkA	trkB	NGF
WT						
Transplant						
Migrated						

	Diff	Apop	Angio	Prolif
Ss45				
Ss25				
Ss25				

Intermediate
Good
Good