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Citation for final published version:

Kasemeier-Kulesa, Jennifer C., Spengler, Jennifer A., Muolo, Connor E., Morrison, Jason A., Woolley, Thomas, Schnell, Santiago and Kulesa, Paul M. 2021. The embryonic trunk neural crest microenvironment regulates the plasticity and invasion of human neuroblastoma via TrkB signaling. *Developmental Biology* 480, pp. 78-90. 10.1016/j.ydbio.2021.08.007

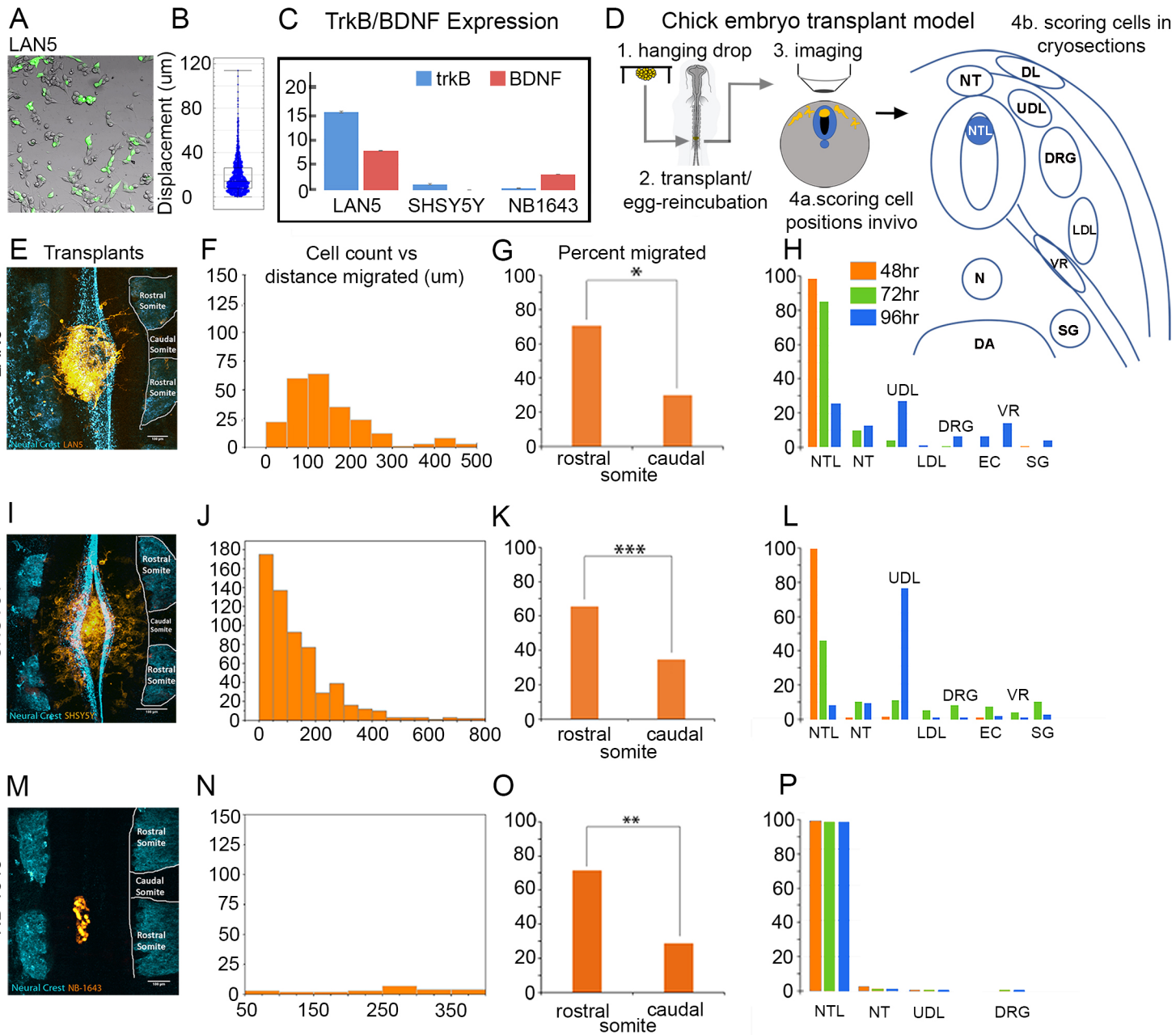
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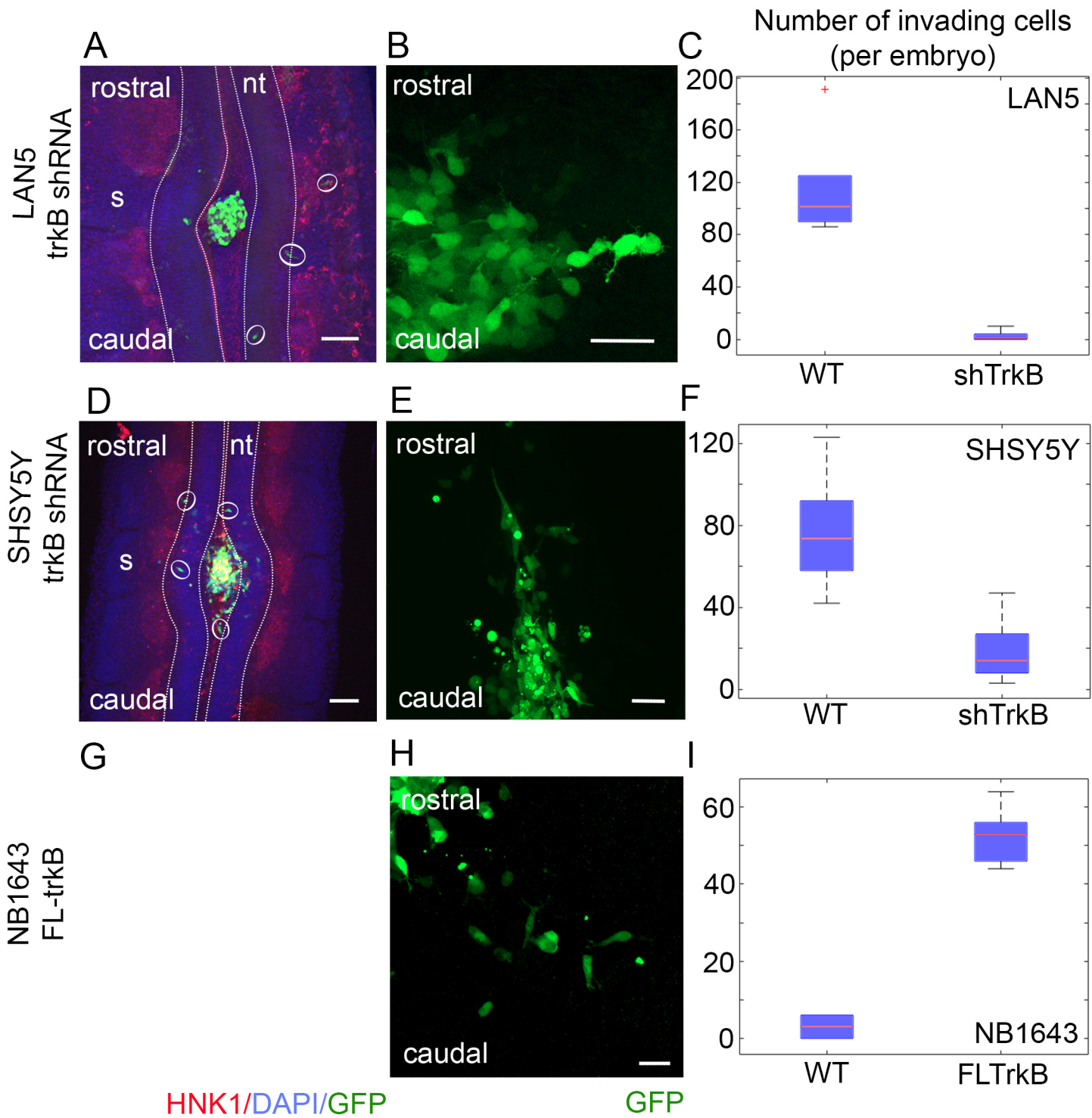
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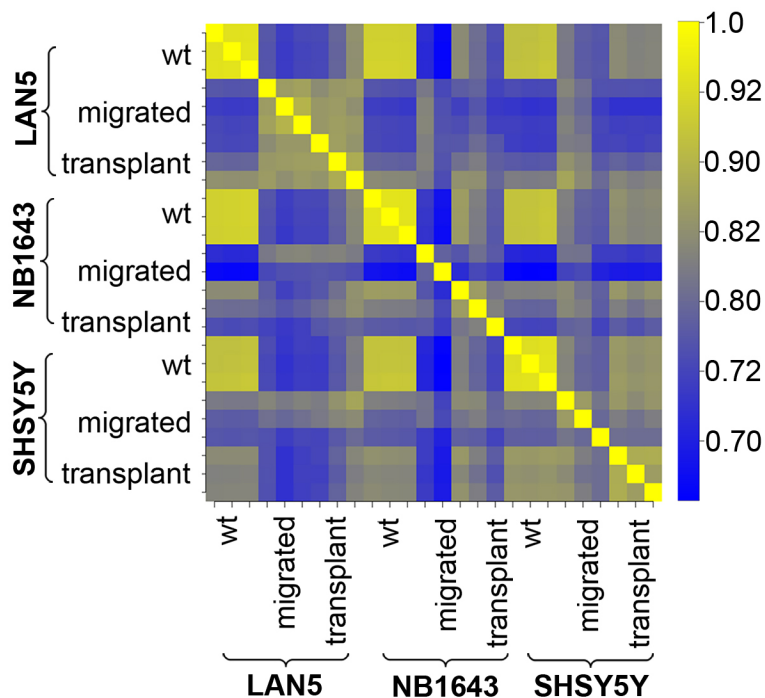
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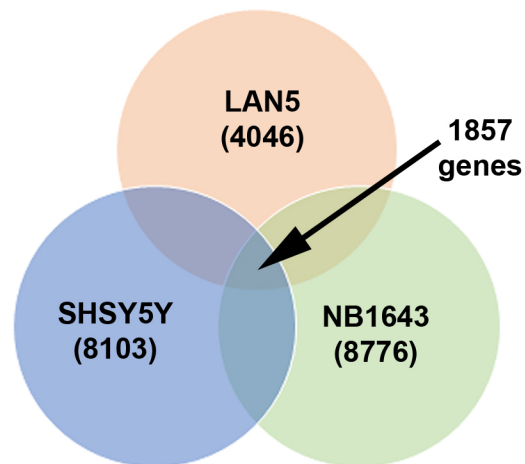




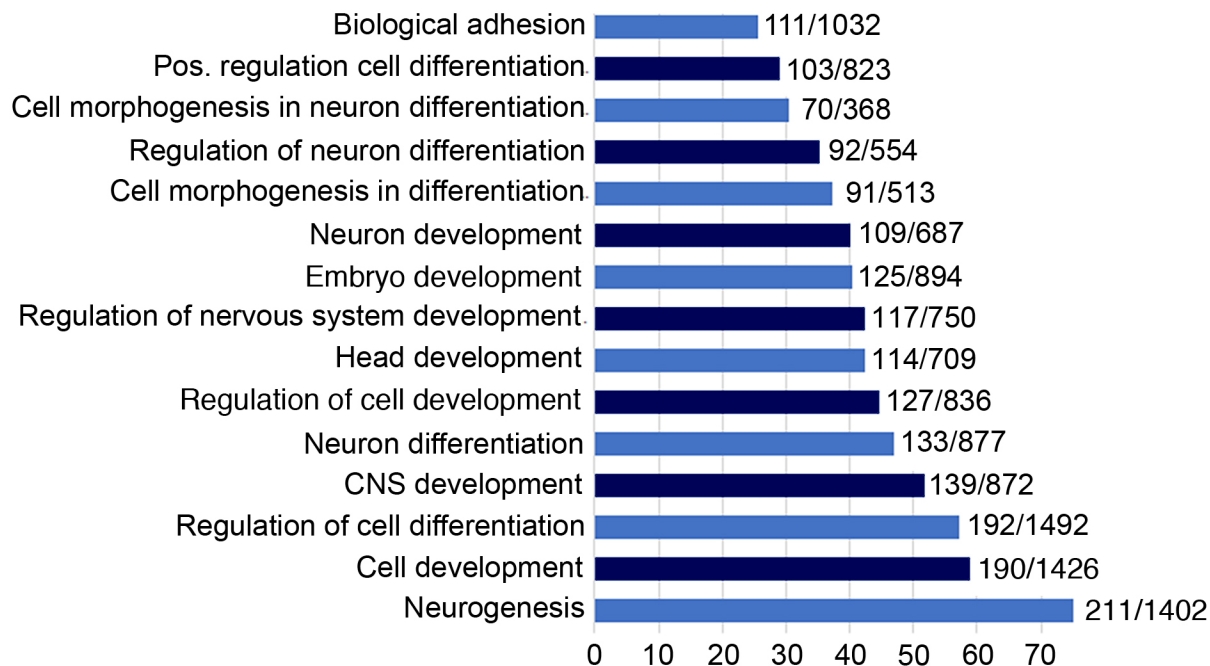
A Gene expression correlation matrix
(after embryo transplantation)
In ovo (migrated vs non-migrated) vs wildtype

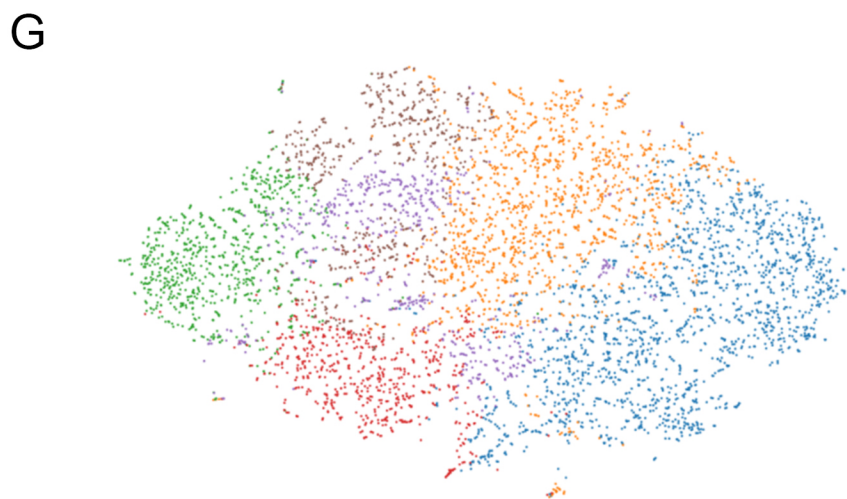
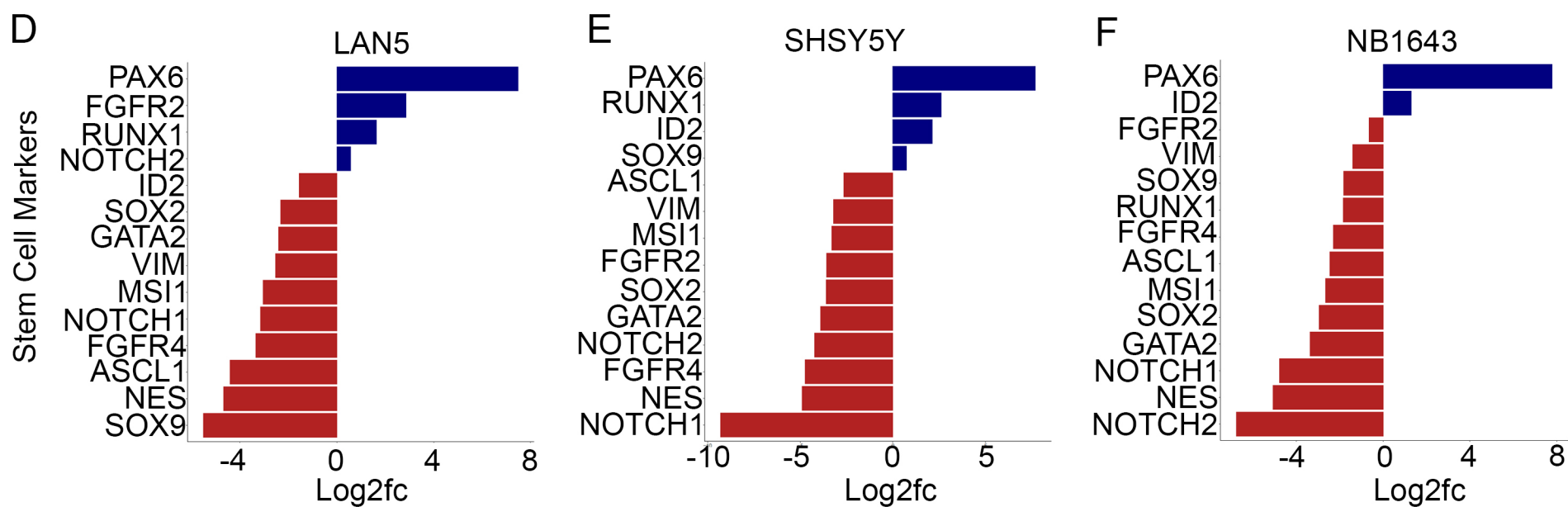
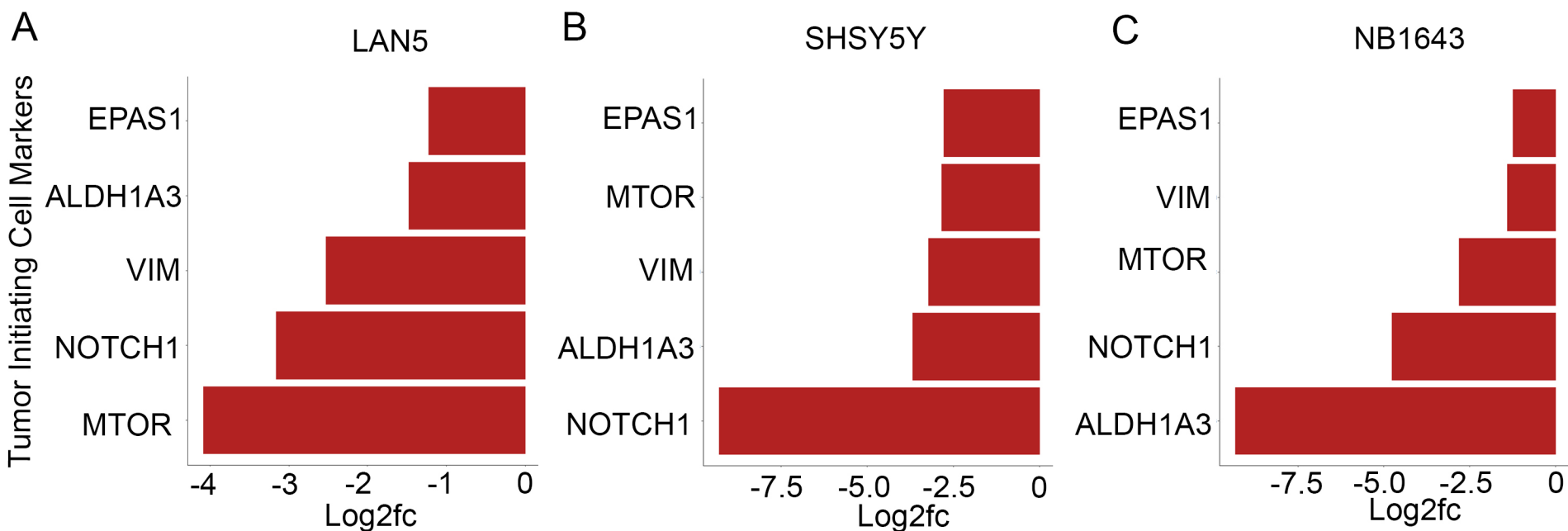


B Changes in the number of genes after embryonic exposure

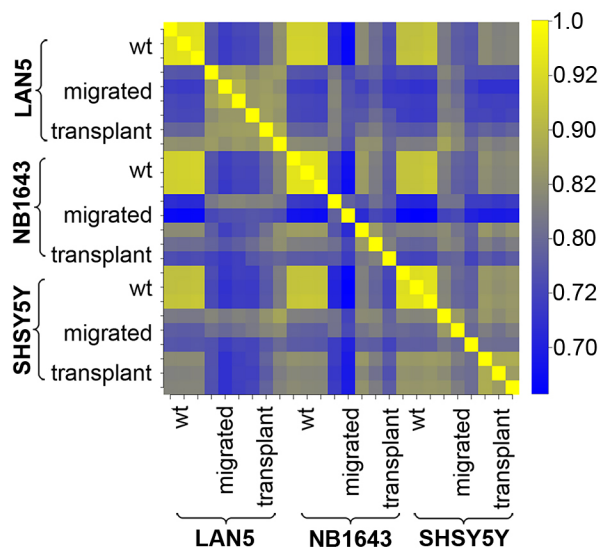


C Go Term Enrichment of neuroblastoma cells after embryonic exposure



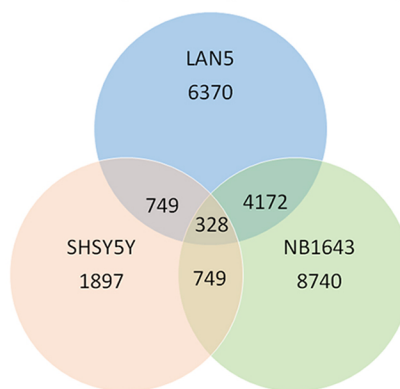


A Gene expression correlation matrix
(after embryo transplantation)
In ovo (migrated vs non-migrated) vs wildtype

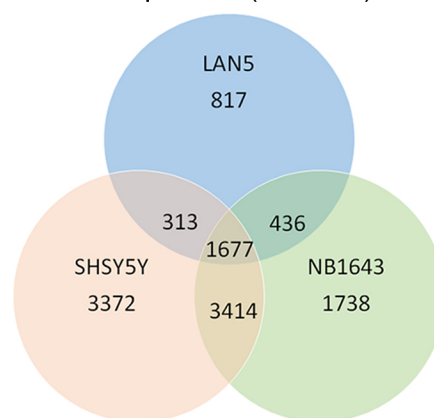


B Changes in the number of genes after embryonic exposure

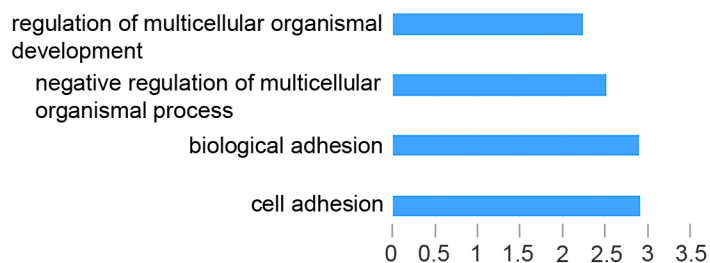
Transplanted (non-invasive)



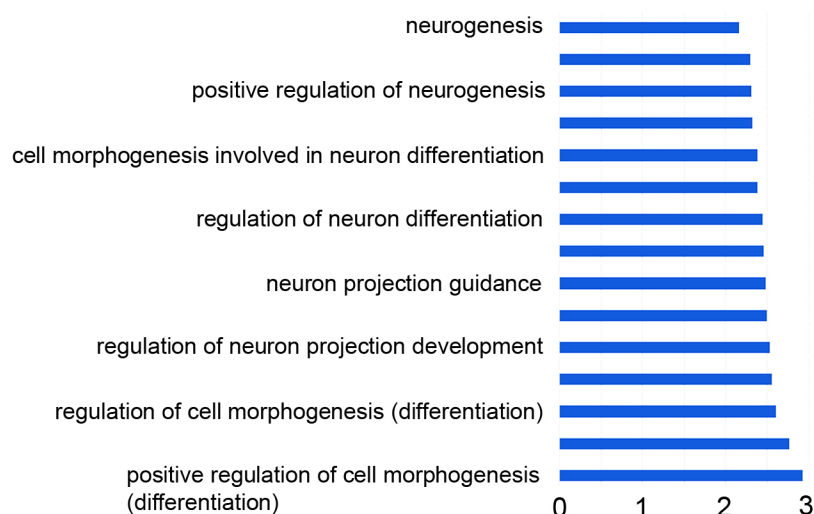
Transplanted (invasive)



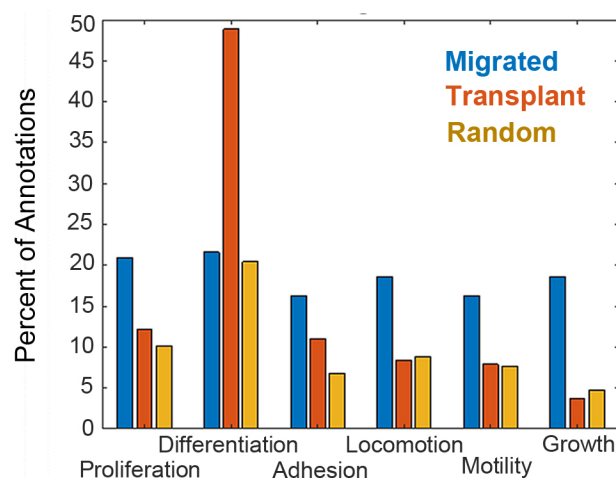
C Go Term Fold Enrichment (328 genes)



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

