

¹Biology Department, ²Texas A&M University, ³Texas A&M Institute for Neuroscience; Texas A&M University, College Station, Texas

Background

Neural progenitor cell (NPC) transplantation is a promising therapeutic strategy following spinal cord injury (SCI). NPCs transplanted into the injured spinal cord undergo specification through normal developmental processes, and give rise to diverse neuronal subtypes (Fig. 1A) & glia. Studies suggest that the graft cell identity may significantly affect circuit-appropriate integration and recovery of function, highlighting the importance of graft composition.

Spinal cord neurogenesis occurs over multiple days with different progenitor classes exhibiting different temporal patterns (Fig.1B).

In this study, we assessed the effects of developmental restriction on NPC graft composition.



Figure 1. (a) Morphogen gradients in embryonic neural tube lead to development of distinct neuronal domains in the adult spinal cord; image adapted from Lai, et al., 2016 [1]. (b) Distribution of BrdU+ cells in embryonic spinal cord after one pulse of BrdU. Image shows differing areas of proliferating cells at various embryonic time points; image adapted from Petracca, et al. 2016 [2].

Experimental Approach

Donor neural progenitor cells (NPCs) were isolated from GFP+ mouse embryonic spinal cords at time points E11.5, E12.5, and E13.5.

NPCs were transplanted into wild type host mice following a C4 dorsal column lesion (1 million cells per subject).

Four weeks post-transplantation, tissue was collected and sectioned at 20µm for immunohistochemical analysis.

Graft composition was characterized using cell markers including NeuN, Sox9, Lbx1, Tlx3, Calbindin, and ChAT.

Quantification was performed with ImageJ and an automated cell counting plugin created in our lab (Chen et al., *in prep*).



Figure 2. NPC transplantation into the lesion site in a mouse model of spinal cord injury. Image adapted from Dulin et al., 2018 [2].

Developmental restriction alters the composition of neural progenitor grafts after spinal cord injury Aleena Lukose¹, Joseph Chen², Miriam Aceves³, Ashley Tucker⁴, Jennifer N. Dulin⁵





these cells present outside of the graft. (c) Quantification of total graft-derived astrocytes (d) Quantification of graftderived astrocytes within the boundary of the graft. (e) Ratio of graft-derived neurons to astrocytes across embryonic time points. (f) Quantification of graft-derived astrocytes found outside of the graft and in the host tissue.

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laboratory

Discussion

Our data suggests that the developmental stage during NPC harvesting may significantly affect graft composition.

The E12.5 NPCs, considered the "golden standard" in cell transplantation studies, showed a tendency for smaller graft volume, and lower neuron and astrocyte numbers.

We are currently characterizing the effects of developmental restriction on "phenotypic identity." Preliminary data suggests that earlier-stage grafts show a more ventral/motor profile, while later-stage grafts show a more dorsal/sensory profile.

Follow-up experiments will be necessary to evaluate how different graft-derived neuron subtypes mediate graft/host integration and recovery of function.

Importantly, these experiments could inform engineering of improved and optimized grafts for translation into the clinic.

References

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Acknowledgments

