

UNIVERSITY OF CALIFORNIA, SAN DIEGO

**HUMAN NEURAL STEM CELL THERAPY IN A PRIMATE MODEL FOR  
PARKINSON'S DISEASE**

A dissertation submitted in partial satisfaction of the requirements  
of the degree Doctor of Philosophy

in

Biomedical Sciences

by

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Chair

University of California, San Diego

2010

## **DEDICATION**

To my family, who have always been my biggest supporters, especially my grandparents, Marjorie V. Wakeman, Harrison “Bud” Wakeman, and Wilma Stookey who taught me honesty and integrity in life.

## EPIGRAPH

“There is a road, no simple highway  
Between the dawn and the dark of night  
And if you go, no one may follow  
That path is for your steps alone”

- *Robert Hunter* -

“Who can unlearn all the facts that I've learned  
As I sat in their chairs and my synapses burned  
The torture of chalk dust collects on my tongue  
Thoughts follow my vision and dance in the sun  
All my vasoconstrictors they come slowly undone  
Can't this wait till I'm old? Can't I live while I'm young?”

- *Tom Marshall* -

“Once in awhile you get shown the light  
In the strangest of places if you look at it right”

- *Jerry Garcia* -

“Sometimes you eat the bar and sometimes the bar eats you”

- *Sam Elliott* -

“The Dude Abides”

- *Jeffrey “Duder” Lebowski* -

“It is a paradox of our time that, seduced by the similarities, we often neglect the importance of distinctions. Indeed it should be expected that millions of years of evolution which selected enhanced mental capacity for survival would leave a significant mark on the organization of the cerebral cortex”

- *Pasko Rakic* -

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## LIST OF ABBREVIATIONS

bFGF	basic fibroblast growth factor
BLBP	brain lipid binding protein
BrdU	5-Bromo-2'-Deoxyuridine
BSA	bovine serum albumin
CAPIT	Core Assessment Program for Intracerebral Transplantation
CC	corpus callosum
CDB	cell dissociation buffer
Ch	Chemicon
CM	conditioned media
CNS	central nervous system
CPu	caudate putamen
DA	dopamine; Dopaminergic
DAT	dopamine transporter
DCX	doublecortin
DPBS	Dulbecco's modified phosphate buffered saline
EAE	experimental allergic encephalomyelitis
ECM	extracellular matrix
EM,	Erlenmeyer
EGF	epidermal growth factor
eGFP	enhanced green fluorescent protein
FB	Fast Blue
FBS	fetal bovine serum
FG	fluorogold
GDM	glial defined medium
GDNF	glial cell line-derived neurotrophic factor
GFAP	glial fibrillar acidic protein
GFP	green fluorescent protein
HBSS	Hank's balanced salt solution
HD	Huntington's disease
huNuc	human nuclei,
(h)ESC	(human) embryonic stem cell
(h)NPC	(human) neural precursor cell
(h)NSC	(human) neural stem cell
(h)fNPC	human fetal neural precursor cell
(h)fNSC	human fetal neural stem cell
IGF-1	insulin-like growth factor-1
ir	immunoreactive
iPS	induced pluripotent stem (cell)
LIF	leukemia inhibitory factor
MAN	multilayer adherent network
MAPK	mitogen activated protein kinase
mCherry	monomeric cherry
MCI	mitochondrial complex I
MFB	medial forebrain bundle
MHC	major histocompatibility complex
MMS	MAN membrane system
MPTP	1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine

MS	Multiple Sclerosis
NB-B27	Neurobasal-B-27 Medium
Neurobasal-N	Neurobasal-neuronal medium
NFM	neurofilament
NGF	nerve growth factor
NS	nigrostriatal
NSA	neurosphere assay
NuMa	nuclear mitotic apparatus
OB	olfactory bulb
OL	oligodendrocyte
OLP	oligodendrocyte progenitor
OSM	oligodendrocyte specification medium
PBS	phosphate buffered saline
PD	Parkinson's disease
PDGF	platelet derived growth factor
PFA	paraformaldehyde
PFS	Parkinson's factor score
PSA-NCAM	polysialylated neural cell adhesion molecule
RG	radial glia
RMS	rostral migratory stream
PS	protamine sulfate
SC	Santa Cruz
SCI	spinal cord injury
SPIO	super-paramagnetic iron oxide
SN	substantia nigra
SNpc	substantia nigra pars compacta
SVZ	subventricular zone
TBS	Tris-buffered saline.
TC	tissue culture
TH	tyrosine hydroxylase
VM	ventral mesencephalic
2D	two dimensional
3D	three dimensional
6-OHDA	6-hydroxydopamine

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author was a co-author of this paper. DRW performed transplantations, immunohistochemistry/ cytochemistry, analyzed data, co-prepared manuscript, co-assembled figures, and edited text. DER was principal investigator, performed transplantations and oversaw all animal operations, co-prepared entire manuscript, and managed the project. KBB performed stereotactic counting. BCB, EAM and ZG performed IHC. JRS prepared histological figures. YDT, VO, JO, RG, XHP cultured cells. RHR, JDE, RLS, EYS, SAL, SUK supervised. Together with his coauthors, we thank the staff of St. Kitts Biomedical Research Foundation for their contributions to the *in vivo* primate studies, Csaba Leranth and Robert Makuch for histological and statistical advice, and Marcel Daadi for advice and studies differentiating hNSCs into DA neurons. This work was supported by National Institute of Neurological Disorders and Stroke Grants RO1NS40822, PO1NS44281 (to D.E.R.), and R21NS053935; Veterans Affairs Biomedical Laboratory Research and Development Grant 121F (to Y.D.T.); the National Institutes of Health/National Institute of General Medical Sciences Grant T32GM08666 (to D.R.W.); the Axion Research Foundation; Project ALS; the American Parkinson's Disease Association; the Michael J. Fox Foundation; the International Organization of Glutaric Acidemia; the A-T Children's Project; and an anonymous donor to the Combined Jewish Philanthropies.

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## PUBLICATIONS

### Abstracts:

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**ABSTRACT OF THE DISSERTATION**

**HUMAN NEURAL STEM CELL THERAPY IN A PRIMATE MODEL FOR  
PARKINSON'S DISEASE**

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Neural stem cells derived from various sources have shown considerable promise for the treatment of parkinsonian symptoms in a variety of animal models; however, the long-term potential of human neural stem cells to engraft, differentiate into dopamine neurons, and restore function in the dopamine-depleted non-human primate brain remains unknown. This dissertation describes a clinically relevant paradigm for transplantation of undifferentiated human fetal subventricular-zone derived NSC in the parkinsonian primate, including gene therapy enhancement with GDNF. Chapter 1 reviews literature regarding functional properties of NSC that make them a candidate for cellular transplantation in neurodegenerative disorders, the potent neurotrophic effects of GDNF, as well as the rationale for utilizing large animals to adequately test stem cell therapeutics. Chapter 2 describes a novel culturing system to overcome cellular senescence and efficiently expand large-scale human fetal NSC long-term

without aberration. Included are clinically pertinent techniques to produce and deliver hfNSC into both the rodent and primate brain, as well as new advances in radiological tracking and imaging utilizing FDA-approved supra-magnetic iron-oxide particles. Chapter 3 demonstrates the first reported evidence that undifferentiated hfNSC differentiate into some TH-ir neurons and restore functional deficits in parkinsonian in non-human primates. Normalization of dopamine-levels and nigrostriatal circuitry argue for neuroprotective effects of endogenous cells rather than direct cell replacement. To allow for morphological analysis of donor grafts, reporter hfNSC were created (Ch.2) for long-term studies. Chapter 4 presents a new paradigm in which hfNSC are homotopically transplanted concomitantly with striatal AAV-GDNF to enhance graft survival and promote axonal outgrowth. Transplanted cells engrafted for up to 11-months and differentiated extensively congruent with host circuitry, demonstrating for the first time that undifferentiated hfNSC retain developmentally relevant programs of differentiation and respond to host signals in the dopamine-depleted primate brain. Further, evidence supports the standing argument that fetal subventricular-zone derived NSC do not significantly differentiate into mature A9-subtype midbrain dopamine neurons in-vivo, even in a GDNF-rich environment. Chapter 5 discusses the significance of these findings to cellular transplantation in the adult CNS and the future application of stem cell transplantation with regards to Parkinson's disease.

## CHAPTER 1A:

### INTRODUCTION-

#### THERAPEUTIC RATIONALE FOR HUMAN NEURAL STEM CELL TRANSPLANTATION IN PARKINSON'S DISEASE

##### 1A.1 Background

Parkinson's disease (PD) presently affects 1 to 1.5 million Americans, and this number is expected to increase with the aging of the population. Present treatments are palliative and some possibly contribute to disease progression. The proposed research, as next steps for developing a biologically restorative treatment method from "replacement" cells<sup>1</sup>, could lead to a treatment for patients with early stage as well as untreatable late-stage Parkinson's disease.

Loss of DA neurons in the substantia nigra (SN) and the resulting deficit in DA release in the striatum and other areas appear to be responsible for the characteristic motor and cognitive manifestations of PD<sup>2</sup>. Substantial improvements result from the systemic administration of the DA precursor, L-Dopa, or DA agonists<sup>3</sup>, supporting the idea that DA release provided by transplanted replacement neurons might have therapeutic effects. This seems now to have been verified in numerous studies in rodents<sup>4-8</sup> and non-human primates<sup>9-17</sup> which have shown that grafts of fetal DA neurons can lead to improvements in biochemical and behavioral indices of DA deficiency. However, in clinical studies, the improvements in parkinsonism have been rather modest and variable<sup>18-30</sup>. The main reasons appear to be insufficient restoration of striatal DA tone which, in turn, stems from the large volume of striatum that has to be reinnervated in man, together with the low rate of survival of transplanted fetal DA neurons, (reviewed by Redmond<sup>31</sup> and Wakeman<sup>32</sup>).

It is the goal of this dissertation to test the potential of human neural stem cells (hNSC) to reverse dysfunction in the neurotoxic model of parkinsonism produced by MPTP in monkeys. We have chosen to focus on how best to achieve permanent dopamine (DA) replacement and functional recovery from human stem cells in the most physiologic and functional manner without side effects or cell over-growth. The following dissertation chapters address fundamental unknown questions relevant to stem cell therapeutics and introduce novel methodological advances critical for advancing stem cell transplantation therapies into clinically relevant paradigms. Specifically, the development of new defined culturing systems for long-term mass-production of human fetal neural stem cells (hfNSC), as well as a developmentally defined system to systematically produce oligodendrocyte precursor cells in-vitro. In addition, we developed methods for tracking transplanted hNSC with iron-particles and magnetic resonance imaging and refined neurosurgical techniques to mimic the clinical surgical setting in the most relevant non-human primate model of parkinsonism. With the development of these tools, we sought to determine whether:

- (1)** Undifferentiated hfNSCs will engraft, survive, differentiate, and induce significant recovery of function in the dopamine-depleted striatum and substantia nigra (SN) (Chapters 3 and 4).
- (2)** HfNSCs will promote a return to homeostasis through multiple compensatory mechanisms.
  - i.** Provide neuroprotective support to host DA neurons and circuitry and preserve or increase DA concentrations, DA cell survival, and the endogenous connections that already exist (Chapter 3).
  - ii.** Differentiate into other relevant support cells (astrocytes, oligodendrocytes)

and secrete neurotrophic factors (Chapters 2A, 3, 4, Appendix A)

- (3) AAV-GDNF overexpression in the host will increase survival and distribution of endogenous and hfNSC derived cells, specifically the induction of the A9-nigral midbrain phenotype, and favorably direct their neuritic outgrowth toward striatal target regions (Chapters 3 and 4).

### 1A.2 Beneficial Characteristics of hNSC

Stem cells may provide a solution to many of the known and suspected problems of neural transplantation with fetal DA precursor cells. Multiple classes of human "stem cells," have been defined, including hNSCs, the most primordial, uncommitted and multipotent cells of the nervous system, which give rise to the vast array of more specialized cells of the CNS<sup>33-53</sup>. HfNSCs have been most effectively obtained from neuroectodermal-derived germinal zones, e.g., the ventricular zone of the fetal telencephalon<sup>54</sup> and are operationally defined by their ability to differentiate into most (if not all) neuronal and glial lineages in multiple anatomical and developmental contexts, and to populate developing and/or degenerating CNS regions. They can be readily isolated from the brain and *efficiently expanded* in vitro by multiple safe and effective means, propagated, characterized, manipulated, and tested for efficacy and lack of pathogenicity. In addition, NSCs have been reported to have a number of properties that might make them useful for brain repair, including (1) Facile engraftability (particularly into germinal zones), (2) Permeability within the blood brain barrier, (3) Genetic manipulability (easily transduced ex-vivo by most viral and non-viral gene transfer methods), (4) Migratory behavior enabling replacement of cells and genes at disseminated lesion sites,<sup>55</sup> (5) Plasticity, (6) Tropism and trophism within regions of active CNS degeneration<sup>56-58</sup>, and (7) Low immunogenicity despite high MHC expression<sup>59</sup> (Reviewed by Park<sup>60,61</sup>). Prior data in rodents (by many investigators) and progress with hNSCs obtained from human fetal brain

(reported by our group) suggest that stem cells may provide both cellular replacement and neurorestorative effects in the host that may be applicable to the treatment of PD.

### **1A.3 Non-Human Primates are Critical for Pre-clinical Evaluation**

There are numerous dissimilarities between the brains of rodents and primates that limit the clinical predictive power and experimental utility of rodent experiments alone. The most relevant differences begin with genomics but are manifested as differences in (a) biochemistry, (b) pharmacological responsiveness, (c) development, (d) brain size and organization, and (e) behavior. In all of these areas, Old World non-human primates are far more like humans. In addition, recent genetic studies showing that an alpha-synuclein substitution/mutation at P53 (an alanine for a threonine), which is associated with neurodegenerative diseases, including PD, occurred about 38 million years ago at the divergence of Old World from New World primates<sup>62</sup>. This difference may help to explain the unique vulnerabilities of Old World primates to neurotoxins such as MPTP and, perhaps, with additional genetic alterations, the unique vulnerability of humans to PD. Significant other biochemical and pharmacological differences exist between rodent and primate brains in aspects of the monoamine systems relevant to PD<sup>63</sup>; there are fewer differences between monkey and human. The existence *only* in primates of the excellent symptomatic model of human parkinsonism after systemic MPTP exposure raises powerful ethical arguments in *favor* of using it to develop the most effective clinical interventions possible. Misinformation from rat studies has led to erroneous but tenacious interpretations of human clinical trials. For example, despite the large number of rodent PD "model" studies using fetal tissue, they failed to anticipate the dyskinetic side effects ultimately observed following the use of such graft material in actual PD patients and overestimated its chances for success. Work with primates, while having the advantage of being much more directly connected to the human condition, is

extremely labor-intensive and requires longer periods of time to prepare animals, carry out experiments, and analyze post mortem specimens. However, we believe that the value of the data ultimately acquired provides information that cannot be obtained or predicted from studies in rodents or even from post mortem outcomes in human patients (See also: Chapter 1B for a detailed description)

#### **1A.4 MPTP Model of PD in Non-Human Primates**

For functional studies to have relevance to PD, it is important to study the best available models. MPTP was inadvertently discovered to induce Parkinson's disease in humans as the toxic by-product of a synthetic street drug<sup>64,65</sup>. MPTP is a potent, irreversible inhibitor of mitochondrial complex I (MCI) when converted by astrocytes to MPP<sup>+</sup> and taken into DA neurons by the dopamine transporter (DAT)<sup>66-71</sup>, causing selective degeneration of A9 DA neurons within the midbrain structure, substantia nigra pars compacta (SNpc), thus mimicking human PD. The range of variability, course, and duration of the systemic MPTP syndrome in the African green monkey has been studied for periods up to 14 years and shown that this species shows most of the characteristics of PD. The use of highly symptomatic parkinsonian animals in order to improve the parallels with PD and to assure the stability (or deterioration) of the condition over time, also has drawbacks due to the care required to maintain life and to minimize stress and discomfort. Unfortunately, no other model of PD (not unilateral intra-carotid MPTP infusion or unilateral or bilateral 6-OHDA administration) in non-human primates mimics as closely the full behavioral syndrome or neuropathology<sup>66-68</sup>. Fortuitously, the St. Kitts green monkey appears to show a more complete model of parkinsonism than any other species of monkey<sup>69</sup>. In addition, our behavioral assessment method for parkinsonism correlates extremely well with measures of DA transporters *in vivo* and DA concentrations post mortem. Our studies of DA transporters *in vivo* using SPECT

imaging with [<sup>123</sup>I]β-CIT in the striatum of MPTP monkeys have included determination of the reproducibility of the imaging measures, optimization of the SPECT outcome measure, and longitudinal assessment of MPTP animals with SPECT. Mean DA punch biopsy measures were significantly correlated with SPECT striatal uptake ( $r=0.80$ ,  $n=18$ ,  $p<0.015$ ). DA concentrations in post-mortem biochemical punches highly correlate with both the Parkinson Score and Healthy Behavior Score. This finding is similar to the finding that UPDRS correlates with SPECT beta-CIT in patients<sup>70,71</sup>.

### 1A.5 hNSCs Effect on the Parkinsonian Brain

A major aim of the proposed studies will be to determine the adequacy of hNSC to (1) provide properly distributed, functionally integrated dopamine neurons to restore normal function and (2) to provide neuroprotective support to host DA neurons and circuitry in order to preserve the connections that already exist. Though multiple studies have now demonstrated that implanted neural progenitors, when transplanted in *non*-neurogenic regions of the intact adult brain do not yield neurons<sup>41,72,73</sup>, more recent evidence suggests that, within neurodegenerative environments, neurogenic signals are recapitulated (at least transiently) such that new, donor NSC-derived neurons now emerge<sup>37,74,75</sup> successfully engraft and assume legitimate neural phenotypes. It appears that such "self-repair" signals to which NSCs are responsive may be elaborated constitutively in the adult DA-depleted primate brain and now confirmed by our studies in monkeys. However, they may not be of sufficient magnitude to yield optimal functional recovery. Hence, an additional strategy that must be compared with unmodified hNSCs would be to *increase* the proportion of hNSCs that differentiate into DA neurons. The most obvious method for increasing these numbers is to inject more hNSCs. But based upon some on-going and published studies in the St. Kitts green monkey, we failed to increase the number of surviving TH+ cells by doubling the amount of primary fetal tissue

injected, suggesting that the host environment may limit this approach for stem cells also. In our recent studies, there also appeared to be no clear relationship between the number of hNSC injected and the number that were found post-mortem.

#### **1A.6 GDNF enhances survival and outgrowth of immature DA neurons:**

A recognized problem with conventional primary fetal neural tissue transplantation is the high percentage of transplanted cells that die. It appears that most DA neurons are, in fact, apoptotic within the first 24 hours of grafting<sup>76</sup>. Thus, early loss of DA neurons probably accounts in large part for the observation that 90-99% of transplanted neurons are lost by the end of a study, typically several weeks after grafting occurred<sup>77-81</sup>. The relatively impoverished neurotrophic environment of the adult brain is one factor that may trigger death of grafted DA neurons<sup>82-87</sup>. There have been attempts to counter the effects of an experimentally induced DA loss with transplants to the striatum of primate stem cells converted *ex vivo* to a DA phenotype. Success has been mixed with regard to survival of transplanted cells or loss of phenotype and hence functional impact<sup>95</sup>.

GDNF has been found to be a potent neurotrophic factor necessary for DA neuron survival in the developmental and adult stages<sup>88</sup>, and its actions on the VM are rather selective for DA neurons<sup>89,90</sup>. When administered exogenously to rodents and humans with Parkinsonism, it appears to protect endangered DA neurons<sup>91-94</sup>. Indeed, GDNF supplied via viral vectors in non-transplant paradigms to parkinsonian animal models has been shown to be a potent neuroprotective agent. Overexpression of GDNF induced by a lentiviral vector provides an additional means of trophic support during possibly critical periods<sup>91</sup>. An added benefit of GDNF may not only be the autocrine/paracrine effect such a neurotrophin may have on donor-derived cells, but the protective effect this factor may have on *host* cells that may be in the process of undergoing apoptotic degeneration<sup>95</sup>. Recent work from our collaborators<sup>96</sup>

demonstrated that injection of AAV2-GDNF into the monkey striatum lead to stable and long-lasting (12 month) overexpression of GDNF in that region of the brain. We have also shown that AAV2-mediated gene transfer of GDNF to the striatum of MPTP-treated monkeys results in a marked increase in survival of primary fetal VM DA-precursor cells by several fold and elicits directional outgrowth of axons<sup>97-99</sup>. Furthermore, recent data indicates that GDNF can act as a chemoattractant in stem cell migration in the brain<sup>100-102</sup>. These data provide solid support for the hypothesis that injections of AAV-GDNF in combination with hNSC transplantation may also enhance the survival and outgrowth of DA neurons derived from stem cells.

#### **1A.7 EGFP labeling improves viability of donor cells**

Traditional methods for pre-labeling NSCs prior to transplantation require harsh DNA intercalating thymidine analogs such as BrdU and CldU, which have been shown to greatly underestimate overall engraftment success and create a variety of false positives when administered for extensive periods of time. Generation of an independently labeled fluorescent hNSC line eliminates the need for these toxic compounds, while increasing both cell viability and engraftment efficiency. In addition, fluorescently labeled donor cells can be easily identified among their host counterparts allowing for enhanced visualization of axonal processes and their dendritic counterparts. These optical properties allow us to easily assess the overall multi-potentiality of each subline. The stem cell community now demands extensive validation of donor cell origin to effectively prove that exogenous benefits were directly derived from transplanted donor hNSCs, therefore, extensive characterization of these reporter hNSC sublines is pertinent to the progression of NSC therapy in neurodegenerative disease.

### 1A.8 Significance

Considerable data derived from studies of rodents and now our initial progress in primates indicate that stem cells can be transplanted, survive, migrate, differentiate into TH+ cells, have other normalizing effects on the host, and perform at least some of the functions for which they are specialized. There is a clear possibility that the functional changes and motor deficits produced by MPTP, similar to those in Parkinson's disease, could be reversed, and this possibility is supported by our studies in primates to date. It is important also that the desirable and undesirable consequences of stem cell engraftment be determined in animals which can be studied over long periods of time and whose brains can be examined directly to determine morphological, biochemical, and behavioral consequences of grafting. Such studies may lead to new understanding of the disease processes affecting DA systems, as well as adding to our understanding of brain function and plasticity.

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## **CHAPTER 1B:**

### **INTRODUCTION-**

#### **LARGE ANIMAL MODELS ARE CRITICAL FOR RATIONALLY ADVANCING REGENERATIVE THERAPIES**

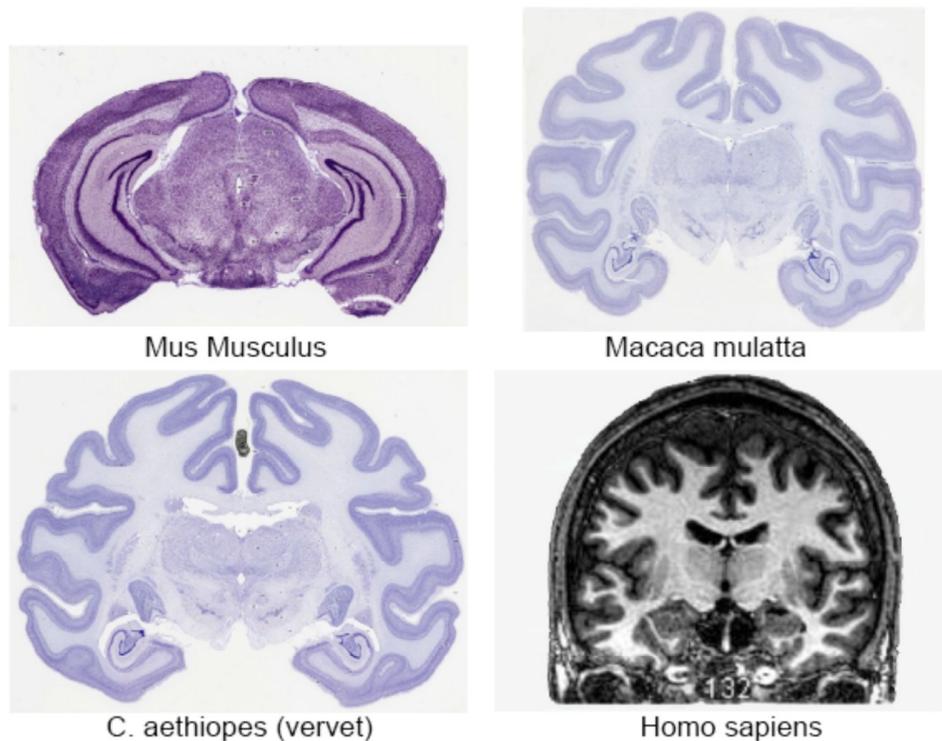
##### **1B.1 Introduction**

Enthusiasm for therapies based on the transplantation of exogenous cells or the transfer of genes by viral vectors has burgeoned over the past thirty years, accompanied by a predictable exhortation to launch clinical trials as soon as possible. The majority of data regarding safety, efficacy, and mechanisms of these therapies have been derived from studies in rodents alone. While such “small animal” systems offer invaluable insights into fundamental biological questions, it is often misleading and perilous to unquestionably equate the higher order motor, sensory, and cognitive processes that characterize human disease with that gleaned from a mouse or rat. Indeed, the literature is littered with clinical trials that failed and, in some cases, led to unforeseen adverse outcomes because the field had leap-frogged over the requisite large animal model. Large animals often provide an essential bridge between insights into fundamental biology and pathophysiology gleaned from simple systems and the realities of treating a human disease. This is often especially true for neurological disorders where not only do differences in size and scale pertain, but also in neuroanatomic connections and organization, cognitive capacities, signaling pathways, genetic redundancy, or the etiology of the disease.

While the gene therapy field has increased their use of non-human primates prior to the application of viral vectors in clinical trials, the cellular therapy field – represented most conspicuously of late by the stem cell field – has only recently begun to properly address this

need. Monkeys and the mini-pig may prove to be excellent preclinical models because of their similar comparative anatomy, pharmacokinetics, and physiologic and metabolic interactions. These models have proven to be extremely useful for studying endocrinological diseases such as type-1 diabetes and neurological disorders including Parkinson's disease (PD), spinal cord injury (SCI), and multiple sclerosis (MS). Prudence would argue that clinical trials for diseases in which differences between rodent and human transcend size and scale should require clear and definitive proof-of-concept in at least one relevant large animal model, in order to safeguard patients. This need, of course is counterbalanced by considerations of the substantial cost, time, and ethical circumspection that typically accompanies such research. In this editorial, we will attempt to help researchers reason through the potential need and advisability of using a large animal model for their particular biological question.

There are numerous differences between rodents and humans that make the unqualified translation of rodent data inadequate. The most relevant differences start at the genomic level and are manifested as differences in pharmacological, biochemical, developmental, behavioral, and functional responses to perturbations and interventions (1). Anatomical and cytoarchitectural differences are particularly profound in the brain, where innervation of specific regions is critical to addressing a neurological deficit (Figure 1A-1). In PD for example, it has been estimated that for each volume of tissue innervated by a rat dopaminergic (DA) neuron, a monkey neuron must innervate 20 times that volume, while humans would need a staggering volume of 200 units to mimic similar reinnervation of nigrostriatal projections (2).



**Figure 1B-1. Coronal images of Nissl stained sections** from *Mus musculus*, *Chlorocebus aethiops*, *Macaca mulatta*, and MRI of human cortex (135).

In addition, differences in monoamine biochemistry (3-7) and developmental life span make extrapolating rodent data to human therapies risky and insufficient. Behavioral differences between rodents and humans are also of great significance (8-10). It is likely that only a large animal model can adequately mirror the complexities of studying and restoring bipedal gait, balance, tremor, hand preference, fine motor coordination, spontaneous blink rate and cognitive deficits (11-16).

## 1B.2 Lesson's from Parkinson's Disease

PD is characterized by degeneration of dopaminergic (DA) neurons in the substantia nigra pars compacta (SNpc) and subsequent loss of striatal DA release. Cell based strategies

for replacement of damaged DA neurons have been a promising strategy since the early 1970's, when Olsen, Stenevi, Bjorkland, and colleagues performed their initial transplantation experiments in rat brains (59-63). Several subsequent studies in rats provided extended evidence for fetal cell transplants repairing DA deficiency (64, 65), however, despite numerous attempts by many groups, these results could never be recapitulated in primates (1). Coupled with mounting ethical considerations, researchers turned to adrenal allografts as a new source of tissue, but these also failed, and subsequent clinical trials were abandoned (66-68). New hope came again after 1985 when human fetal neural cells were derived and successfully implanted into the rat and monkey brain (69-76), however ethical opponents argued that the large amount of fetal tissue required made these procedures for routine regenerative therapy unfeasible and, therefore, unfundable.

Based on the varying methods published in the mid 1970's, several groups began preliminary efficacy and safety clinical studies of fetal ventral mesencephalic grafts into PD patients. Initial studies in Sweden (77), England (78), and Mexico (79) produced variable results, elevating the ethical debate and political controversy, eventually leading to a funding moratorium in the United States (80, 81). Subsequent studies seemed promising but the overall consensus was that the results were inconclusive due to variability within small studies, limited short-term functional improvements, and overall difficulty in comparing independent procedures (1). To investigate this problem and improve the ability to compare studies, a panel of experts published a recommended protocol for future investigations termed, Core Assessment Program for Intracerebral Transplantation (CAPIT) (82).

In the same year, the first randomized controlled clinical trial, headed by Curt Freed and Stanley Fahn, was initiated studying 40 PD patients over seven years, half of which underwent bilateral stereotactic transplantation of human embryonic mesencephalic tissue into the putamen (83). Although initial results appeared promising, it was determined that no

benefit in primary outcome measure had been attained after 12 months post-op. However, a small subset of patients < 60 years old did show small double blind improvements on a validated PD rating scale. More importantly, predictions made from animal models failed to predict the relatively high occurrence of dystonia and dyskinesia associated with long-term survival (84). These gross abnormalities occurred in 5 of 33 patients that lived 3 years post-transplant, of whom all five were less than 60 years of age and had displayed improvements in PD score after 12 months post surgery.

The human trials appeared to have a solid rationale based on a decade of rodent studies. Hundreds of investigators had reported survival and functional recovery from fetal ventral mesencephalic derived neural tissue transplanted into the striatum of rats (85). The grafts were shown to express tyrosine hydroxylase (TH), increase DA concentrations, project long outgrowths, form synaptic connections, and respond to afferent stimulation (86-90). However, the rat models of PD overestimated the likely success of fetal precursor transplants and failed to anticipate the dyskinetic side effects in human trials (84). And, although large animal models were tested, it is possible that they were not followed for a long enough time, simply because the vast difference between rodents and primates (of which human is an example) was not appreciated. Furthermore, while the Freed et al. study made extremely valuable contributions to our understanding of PD, small design flaws such as method of tissue preparation, limited target area, and lack of immunosuppression may have inherently impeded the overall success of the study (1). Many of these details may have been worked out in large animals before reaching the clinic, potentially predicting adverse effects. Unfortunately, cell replacement therapy received negative media coverage (91-94), creating an impression of defeat in the public's mind and even creating dissension among PD investigators (84, 92, 95-102). Much of this may have been avoided had a higher priority been placed on more extensive primate studies with larger numbers and longer follow-up.

### **1B.3 Going Forward in PD- A Case-in-Point for Choosing the Correct Model**

Having detailed above the pitfalls into which the PD field slipped by relying upon non-representative data for its first cell-based clinical trial, how might it recover its momentum – and can this serve as a lesson for future trials for other diseases?

As noted above, most of the early transplantation studies in PD were performed in rodents that had received a unilateral injection of the dopaminergic toxin 6-hydroxydopamine (6-OHDA) into the medial forebrain bundle or nearby (103, 104). Although transplantation appeared to promote recovery from the resultant PD-like symptoms, these behaviors actually bore little resemblance to human PD. If therapies were to be tested solely on such a rodent model, the results would likely have poor predictive value for actual patients. On the other hand, through serendipity in the 60's and 70's, it was discovered that the complex I inhibitor, 1-methyl-4-phenyl-1, 2,3,6-tetrahydropyridine (MPTP), created pathology quite similar to human idiopathic PD in both humans (105-107) and monkeys (70, 108-112), especially the vervet (113, 114). MPTP depletes SN TH-ir and striatal DA to levels similar to PD, as well as producing characteristic Parkinsonian symptomatology including akinesia, rigidity, postural alterations, and tremor (9, 10). Overexpressing alpha-synuclein in conjunction with MPTP may create Lewy bodies (115, 116). Hence, systemically administered MPTP in African green monkeys has become acknowledged as the most authentic model for actual human idiopathic PD and the model in which the safety and efficacy of viral vector- and cell-based therapies in humans can most reliably be judged (expert reviews- 1, 8-10, 117). Indeed, the potential use of lentiviral and AAV-based vectors for delivering neurotrophic factors in PD patients has been and is currently being tested in monkey first (118-120). Such studies, in turn, did set the stage for some phase-1 clinical trials (121-131). It is disheartening that very few investigators to date in the cell therapy field (including the stem cell field) have taken advantage of this model

for work in PD despite widespread recognition of its power. Most likely this reticence is due to the huge expense entailed in performing such complex studies with a sufficiently large sample size and with suitably objective and predictive readouts over a long enough period of observation. Nevertheless, that expense is vastly smaller than the cost to the health care system, to science, and to patient well being when an ineffective or injurious clinical intervention becomes misguidedly launched. Funding agencies and scientists must begin to change their thinking and view adequately powered large animal studies for some diseases not as a luxury, but as a necessity.

Another source of hesitancy in the use of large animals, particularly monkeys, is not solely financial but also ethical. There is no question that experimentation on “higher order” animals provoke, more than any other type of scientific study, circumspection regarding the appropriateness of animal experimentation. Although we have outlined above its scientific justification, there is no question that investigators must be acutely sensitive (as they should be for all animal work) to using the minimal number of monkeys required to statistically overcome type 1 and type 2 error rates (1), to treating the animals humanely, and to using animals only in the most judicious manner, for only exquisitely well-conceived and rigorously designed experiments.

Another level of ethical consideration that has recently emerged as a consequence of the burgeoning human stem cell field is whether integrating human cells into the brains of non-human primates (particularly prenatally) somehow “humanizes” them. In other words, might a human-primate chimeric brain develop consciousness and hence merit consideration beyond that of just an experimental laboratory animal (131-134). While an interesting metaphysical consideration, chimerism in our hands, even under the most optimal transplantation conditions, yields only a relatively small human cell contribution to an

overwhelmingly primate CNS structure. Furthermore, we have never observed anything but typical primate behaviors, emotions, and skills in our animals.

#### **1B.4 Summary and Conclusions**

Appropriately assessing the safety and efficacy of many future interventions in regenerative medicine will require large animal pre-clinical testing in order to avoid the failures of previous studies in patients. Such recognition requires scientists, investors, funding agencies, and the public to recalibrate their thinking. Simpler rodent models, while useful for unraveling certain biological conundrums often fall short in their predictive value for human disease or their appropriateness for devising effective delivery methods. A greater number of multi center collaborations may be required to insure adequate funding and expertise for large animal studies. Such an approach requires a less egocentric approach to scientific accomplishment. The ultimate cost of *not* including the rationale use of large animal models in preclinical studies – the price of unsuccessful trials, injury to patients, and failure to bring potentially groundbreaking therapies to the bedside – far outweighs the short-term expense of the studies themselves.

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## CHAPTER 2:

# LONG-TERM MULTILAYER ADHERENT NETWORK (MAN) EXPANSION, MAINTENANCE, AND CHARACTERIZATION, CHEMICAL AND GENETIC, MANIPULATION, AND TRANSPLANTATION OF HUMAN FETAL FOREBRAIN NEURAL STEM CELLS

### 2.1 Summary:

Human neural stem/precursor cells (hNSC/hNPC) have been targeted for application in a variety of research models and as prospective candidates for cell-based therapeutic modalities in central nervous system (CNS) disorders. To this end, the successful derivation, expansion, and sustained maintenance of undifferentiated hNSC/hNPC in vitro, as artificial expandable neurogenic micro-niches, promises a diversity of applications as well as future potential for a variety of experimental paradigms modeling early human neurogenesis, neuronal migration, and neurogenetic disorders, and could also serve as a platform for small-molecule drug screening in the CNS. Furthermore, hNPC transplants provide an alternative substrate for cellular regeneration and restoration of damaged tissue in neurodegenerative disorders such as Parkinson's disease and Alzheimer's disease. Human somatic neural stem/progenitor cells (NSC/NPC) have been derived from a variety of cadaveric sources and proven engraftable in a cytoarchitecturally appropriate manner into the developing and adult rodent and monkey brain while maintaining both functional and migratory capabilities in pathological models of disease. In the following unit, we describe a new procedure that we have successfully employed to maintain operationally defined human somatic NSC/NPC from developing fetal, pre-term postnatal, and adult cadaveric forebrain. Specifically, we outline the

detailed methodology for in vitro expansion, long-term maintenance, manipulation, and transplantation of these multipotent precursors.

## **2.2 Introduction**

A number of techniques have been devised to attempt to identify and isolate rodent and human neural stem/precursor cells (NSCs/NPCs). Some have relied on the aggregation of cells in suspension cultures—termed “neurospheres” and giving rise to the “neurosphere forming assay” (NSA; Reynolds and Weiss, 1992; Reynolds et al., 1992; Rietze and Reynolds, 2006)—for artificially expanding nonclonal NSC/NPC populations in vitro (Singec et al., 2006) in serum-free medium. However, other techniques have been employed prior to (Ryder et al., 1990; Redies et al., 1991; Renfranz et al., 1991; Snyder et al., 1992) and since (Flax et al., 1998; Shihabuddin et al., 1996; Rubio et al., 2000) popularization of the NSA which, in fact, have been found to have beneficial properties compared to the NSA. It is these techniques that our group has long employed to great advantage and success—particularly when interested in using NSCs/NPCs for transplantation, genetic manipulation, rigorous clonal analyses, and developmental studies—and which will be described in this unit. Human embryonic, fetal, newborn, and adult cadaveric CNS precursors have been shown to thrive when derived and maintained as two-dimensional (2-D) adherent cultures. This technique offers many growth and culture advantages over the NSA and, in fact, has come to supplant the NSA in many neurobiological laboratories.

Over the past two decades, numerous techniques have been described for the derivation and expansion of suspension of human neural precursors either in suspension or as adherent monolayers (Ray et al., 1995; Svendsen et al., 1999; Wu et al., 2002; Walsh et al., 2005; Rajan and Snyder, 2006; Ray and Gage, 2006; Pollard et al., 2006a,b), utilizing an assortment of growth factors (Buc-Caron, 1995; Chalmers-Redman et al., 1997; Moyer et al.,

1997; Sah et al., 1997; Svendsen et al., 1998, 1999; Carpenter et al., 1999; Kukekov et al., 1999; Vescovi et al., 1999a,b; Roy et al., 2000; Uchida et al., 2000; Villa et al., 2000; Piper et al., 2000, 2001; Arsenijevic et al., 2001a,b; Keyoung et al., 2001; Palmer et al., 2001; Cai et al., 2002; Laywell et al., 2002; Nunes et al., 2003; Schwartz et al., 2003; Zhang et al., 2005; Conti et al., 2005; Li et al., 2005; Pollard et al., 2006a,b; Yin et al., 2006; Ray, 2008).

In this unit, we outline methodology for the expansion, long-term maintenance, manipulation, and transplantation of human fetal (10- to 25-week) neural precursor cells (hNPC). Specifically, we describe a new method for long-term expansion of karyotypically stable hNPC, termed the Multilayer Adherent Network (MAN), to generate large scale self-renewing multipotent hNPC populations, amenable to in vitro manipulation and transplantation in vivo. We describe in detail the methods we have successfully utilized to prepare and transplant hNPC into the neonatal mouse and adult nonhuman primate. In addition, we provide basic procedures for characterization of undifferentiated and differentiated hNPC, as well as the processing of engrafted brains. Furthermore, we illustrate techniques for the efficient labeling of hNPC, including lentivirus infection and noninvasive super-paramagnetic iron oxide (SPIO) particle transfection. For simplicity's sake, we will refrain from the operational NSC debate and simply refer to both neural stem and progenitor cells as NPCs from here forward.

## **2.3 STRATEGIC PLANNING**

### **2.3.1 Growth Factor Signaling**

Long-term expansion and maintenance of self-renewing NPC in serum-free media (Reynolds et al., 1992; Reynolds and Weiss, 1992; Svendsen et al., 1996; Rosser et al., 1997) requires mitogenic support from either epidermal growth factor (EGF) or basic fibroblast

growth factor (bFGF) to activate mitogen-activated-protein-kinase (MAPK) signaling and support hNPC division (Gensburger et al., 1987; Walicke, 1988; Kornblum et al., 1990; Murphy et al., 1990; Drago et al., 1991a,b; Ray et al., 1993; Vescovi et al., 1993a,b; Bartlett et al., 1994; Kitchens et al., 1994; Ray and Gage, 1994; Ghosh and Greenberg, 1995; Kilpatrick and Bartlett, 1993, 1995; Kilpatrick et al., 1995; Palmer et al., 1995; Vicario-Abejon et al., 1995; Gritti et al., 1996; Kuhn et al., 1997; Qian et al., 1997; Shihabuddin et al., 1997; Caldwell and Svendsen, 1998; Ciccolini and Svendsen, 1998; Gritti et al., 1999; Palmer et al., 1999; Arsenijevic et al., 2001a,b; Caldwell et al., 2001; Temple, 2001; Ostefeld and Svendsen, 2004; Tarasenko et al., 2004; Kelly et al., 2005; Ray and Gage, 2006). In addition, the neurotrophic leukemia inhibitory factor (LIF) has been shown to enhance telomerase expression, improve viability, and extend the time until terminal senescence of hNPC when used in combination with bFGF and/or EGF (Galli et al., 2000; Molne et al., 2000; Shimazaki et al., 2001; Wright et al., 2003; Bonaguidi et al., 2005; Gregg and Weiss, 2005). Although LIF signaling appears to induce gliogenesis in rodent NPC, in our experience, LIF not only enhances survival and doubling time of human NPC but is absolutely essential for the sustained maintenance of symmetric cell divisions in long-term multilayer adherent network cultures. Direct comparisons of NPC derived from different species or by alternate techniques have shown that NPC characteristics are drastically altered by their environmental inputs and retain these intrinsic cellular properties in direct relation to how they are manipulated in vitro (Ray and Gage, 2006). We have empirically determined the specific regimen of growth factors that best supports growth of human fetal forebrain NPC. As a result, we have adopted a strategy for sustained proliferative expansion of karyotypically normal undifferentiated hNPC in basal growth medium consisting of bFGF and LIF (without EGF).

### **2.3.2 Media Formulations**

Although traditional serum-free rodent NPC culture has generally utilized DMEM/F12 supplemented with N2, we have adjusted the recipe to accommodate hNPC by utilizing Neurobasal medium (Invitrogen) with B-27 supplement (without vitamin A) to support long-term proliferation of hNPC in vitro (Brewer et al., 1993; Brewer, 1995, 1997; Svendsen et al., 1995; Brewer and Price, 1996; Brewer and Torricelli, 2007). In addition, heparin is added to stabilize the binding of the bFGF heparin-sulfate proteoglycan to its FGFR-1 receptor (Balaci et al., 1994; Caldwell et al., 2004), potentiating cell-cell attachments that favor adherent monolayer hNPC growth (Richard et al., 1995, 2000). On the day of use, prepare fresh hNPC growth medium plus 20 ng/ml bFGF plus 10 ng/ml LIF (see Reagents and Solutions). Growth factors are added fresh on the day of use due to their relative instability (Kanemura et al., 2005). Contamination is possible and thus Normocin (InvivoGen) is supplemented regularly (48-hr half-life) as an antipathogenic agent (replaces penicillin/streptomycin/amphotericin B to deter mycoplasma, Gram-positive and -negative bacteria, and fungal contamination). Normocin and any other antibiotics employed may be gradually withdrawn from the culture after an adequate period of time as desired. Due to the relatively large amount of time and resources involved in hNPC culture, we highly recommend the use of pathogen-control agents. Normocin has remained the most gentle yet potent and comprehensive single treatment application we have tested thus far.

#### **2.4 LONG-TERM EXPANSION AND MAINTENANCE OF hNPC**

Throughout the expansion process, cryopreservation and functional testing of hNPC lines is necessary for the continued long-term maintenance of healthy proliferative progenitors. Cultures are monitored superficially under the light microscope for morphological aberrations that may occur in artificial culture. Once sufficient cell numbers have been established, a more intensive battery of screens for in vitro and in vivo multipotency should be

employed, particularly when hNPC reach high passage number or whenever a new vial of early passage progenitors are thawed from cryopreservation for mass expansion, to ensure hNPC cultures do not change phenotypically or become lineage restricted with time. To test functionality, several vials are reconstituted to assess the overall freeze/thaw success, cell viability, and sustained multipotency. Throughout culture, the genetic stability of hNPC should be confirmed periodically through spectral karyotyping, microarray fingerprinting, and transcriptome and proteomic analysis to demonstrate a normal chromosomal complement and sustained expression profile of all classical stemness-associated genes (Cai et al., 2006; Chang et al., 2006; Luo et al., 2006a,b; Maurer and Kuscinsky, 2006; Shin and Rao, 2006; Anisimov et al., 2007; Shin et al., 2007). In an effort to reduce time and costly resources, hNPC lines should be regularly tested for these attributes before proceeding with any large animal transplantation studies.

*NOTE:* All incubations are performed in a 37°C, 5% CO<sub>2</sub> humidified incubator, unless otherwise noted.

*NOTE:* All reagents and equipment coming into contact with living cells must be sterile, and aseptic technique should be used accordingly.

*NOTE:* Numerous different types and sizes of tissue culture vessels are described in this unit; the plating volumes for common tissue culture petri dishes, multiwell plates, slides, and flasks are listed in Table 2-1.

**Table 2-1.** Plating Volumes for Different Culture Vessels

Vessel	Area (cm <sup>2</sup> /well)	Volume
<i>Petri dishes</i>		
20 mm	3	1 ml
25 mm	8	2.5 ml
60 mm	25	6 ml
100 mm	78.5	18 ml
<i>Multiwell plates</i>		
6 well	9.6	3.5 ml
12 well	3.8	2 ml
24 well	2	1 ml
48 well	0.75	500 $\mu$ l
96 well	0.32	250 $\mu$ l
<i>Slides</i>		
1 well	9.4	3 ml
2 well	4.2	2 ml
4 well	1.8	1 ml
8 well	0.8	250 $\mu$ l
<i>Flask</i>		
25-cm <sup>2</sup>	25	6-8 ml
75-cm <sup>2</sup>	75	16-20 ml
225-cm <sup>2</sup>	225	40-50 ml

#### **2.4.1 ESTABLISHING AND MAINTAINING MULTILAYER ADHERENT NETWORK (MAN) CULTURES**

Traditionally, we have thawed and grown hNSC as small, slightly adherent aggregates for the first 2 to 3 weeks of culture post-thaw. More recently, however, we have developed a new method for expansion of newly thawed or freshly dissociated undifferentiated hNPC on noncoated flasks free of extracellular matrix (ECM). Establishment of these high-density multilayer adherent networks (MAN) is founded on the basic theory of aggregate formation, but is adapted into a novel adherent system that offers many growth advantages for both the progenitor population and the researcher. As a whole, the MAN assay relies on a combination of the inherent hNPC property of forming fusion aggregates at high density, coupled with the

intrinsic behavior of resting hNPC aggregates to attach and migrate over time. The end result is a highly dynamic, proliferative population of undifferentiated hNPC displaying a variety of advantageous growth parameters. In general, we find that mature MAN hNPC cultures proliferate and expand at an elevated doubling rate (3 to 5 days) compared to their neurosphere counterparts (4 to 7 days; Kanemura et al., 2002, 2005; Mori et al., 2006). In addition, feeding MAN cultures fresh medium can easily be achieved by simply tilting the flask, aspirating or collecting CM, and refilling the flask with new medium. This fast and easy process allows the researcher to replace all or portions of the medium as often as necessary without the harsh mechanical stresses involved in centrifugation.

The key to transitioning traditional aggregate cultures into MAN cultures is the overall density of the hNPC initially plated. Simply stated, the greater the density of hNPC initially plated, the larger the aggregate units, the more quickly they attach, and, thus, the more quickly subsequent mature multilayer adherent networks are established. It should be noted that replating hNPC at densities greater than  $4 \times 10^6$  cells per 25-cm<sup>2</sup> flask will result in overcrowding and subsequent formation of large spheroid cellular masses, negating the entire premise for the initial dissociation. For the most part, high density passaging is only recommended for preparing small cellular clusters prior to cryopreservation, or to quickly establish mature MAN cultures for short-term study. A brief history of the early stages of MAN formation is:

- a. 0 to 24 hr: Cells equilibrate and settle to bottom of flask following an even distribution pattern.
- b. 24 to 48 hr: Cells begin to lightly attach and spread (as evidenced by small microspikes and several small projections; Fig. 2-1A,B).

c. 48 to 72 hr: Aggregated cell clusters continue to spread, elongate, and begin to proliferate and extend into adjacent neighboring clusters, becoming adherent three-dimensional clusters, creating the first evidence of an interlinked network (Fig. 2-1C).

d. 72 to 96 hr: Cell clusters continue to migrate into each other at the periphery and become anchored strongly enough to change medium. These cultures consist mainly of slightly adherent clusters and a small proportion of nonadherent floating aggregates. The cultures can be carefully removed from the incubator to change medium without disrupting the newly formed MAN (Fig. 2-1 D-F).

### ***Materials***

Human NPC (Support Protocol 1): frozen (Support Protocol 2) and freshly thawed (Support Protocol 3) or freshly dissociated as described in Support Protocol 1

25% (v/v) conditioned medium (CM; Support Protocol 4)/75% (v/v) NB-B-27 complete medium (see recipe) containing 40 ng/ml bFGF and 10 ng/ml LIF (bFGF and LIF concentrations based on total volume of medium)

NB-B-27 complete medium (see recipe)

Leukemia inhibitor factor (LIF; Millipore, cat. no. LIF1010)

Basic fibroblast growth factor (bFGF; Millipore, cat. no. GF003)

Normocin (InvivoGEN, cat. no. ant-nr-1)

Dulbecco's PBS with Ca<sup>2+</sup> and Mg<sup>2+</sup> (DPBS; Mediatech, cat. no. 21-030-CM)

Dulbecco's PBS without Ca<sup>2+</sup> or Mg<sup>2+</sup> (CMF-DPBS; Mediatech, cat. no. 21-031-CM)

Accutase (Millipore, cat. no. SCR005) *or* Cell Dissociation Buffer (Invitrogen, cat. no. 13150-

016)

Conditioned medium (CM; Support Protocol 4)

15-ml conical tubes

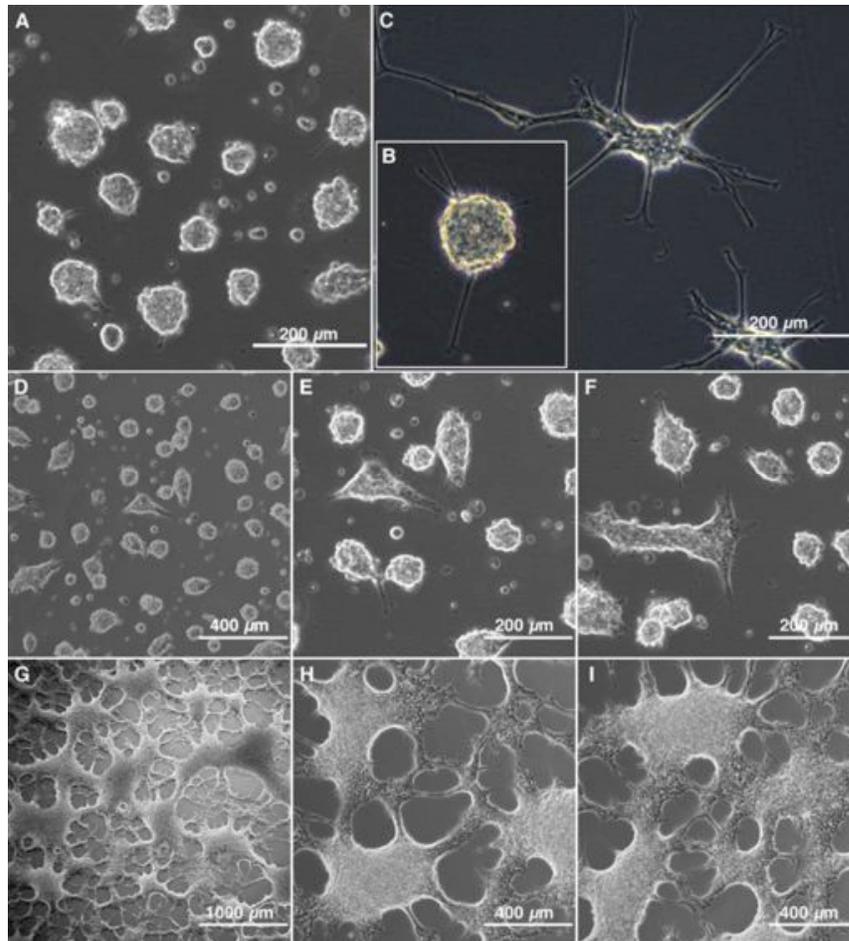
25-cm<sup>2</sup> and 75-cm<sup>2</sup> tissue culture flasks

Battery-powered pipetting aid (e.g., Drummond Pipet-Aid XP)

1000- $\mu$ l extended-length pipet tip and 1000- $\mu$ l automatic pipettor

Centrifuge

Additional reagents and equipment for counting viable cells by trypan blue exclusion



**Figure 2-1. Establishment of multilayer adherent network (MAN).** (A) 24 to 48 hr after plating, hNSC (HFB-2050) readily form evenly spaced, proliferative aggregated cell clusters. Small hNPC clusters initially attach to the culturing surface and sample the local microenvironment with meandering growth-cone like protrusions (B), and eventually flatten and spread out (C). Taking advantage of higher plating densities, the MAN culturing technique creates optimal spacing between colonies, allowing each aggregate cluster close access to neighboring signaling molecules. (D-F) After 72 hr, hNSC aggregates are lightly attached to the surface and begin to actively proliferate. Over the next 3 to 4 weeks, hNSC rapidly expand and form extensive honeycomb-shaped, mature multilayer adherent networks (G-I).

#### 2.4.1.1 Establish MAN cultures

1. To establish MAN cultures from freshly thawed cells or freshly dissociated cells, resuspend hNPC 2:1 (i.e., at  $2-3 \times 10^6$  cells/flask) in 25% (v/v) CM/75% (v/v) NB-B-27 complete medium (containing 40 ng/ml bFGF, 10 ng/ml LIF, and 2  $\mu$ l/ml Normocin) in a 15-ml conical

tube. Transfer hNPC to an uncoated 25-cm<sup>2</sup> flask (Fig. 2-2).

*Ratios such as 2:1 refer to the surface area used—i.e., if starting with one 25-cm<sup>2</sup> flask, when expanding cells, one would use a 1:2 split, meaning that one should start with one 25-cm<sup>2</sup> flask and resuspend the dissociated cells into two 25-cm<sup>2</sup> flasks—increasing the surface area from 25 to 50 or 1:2. However, if referring to establishment of a culture with frozen cells, the ratio is 2:1, i.e., the number of frozen cells that were originally in two 25-cm<sup>2</sup> flasks would need to be thawed into one 25-cm<sup>2</sup> flask. Similarly, when dealing with freshly dissociated cells, the ratio is 2:1. In this step, 2 million fresh cells or 3 million frozen cells are diluted into 8 ml media into one 25-cm<sup>2</sup> flask.*

*Plating a higher density of hNPC leads to the quicker (24- to 72-hr) formation of small (2–3 ° — 10<sup>6</sup> cells) to medium size (3–4 ° — 10<sup>6</sup> cells) clusters, respectively, initiating close cell-cell contacts critical for enhanced paracrine and autocrine support. This means that if you plate 2–3 ° — 10<sup>6</sup> cells (dissociated) into one 25-cm<sup>2</sup> flask, it will give you small clusters within 24 to 72 hr, whereas 3–4 ° — 10<sup>6</sup> will give medium-size clusters in this same period of time.*

*Interestingly, we have found that leukemia inhibitory factor (LIF) is absolutely necessary and essential for the long-term maintenance of MAN cultures. Removal of LIF from the basal growth medium results in the rapid breakdown of elongated projections into ropelike, flexible, spindly, nonadherent protrusions that eventually disappear, ultimately resulting in the loss of proliferation capacity, increased senescence, and eventual cellular crisis.*

2. Place the flask into a humidified 5% CO<sub>2</sub> incubator at 37°C and shake horizontally in both planes to evenly disperse cells throughout the flask without sloshing medium into the neck of the flask.

*It is imperative that small (4- to 16-cell) to medium size (16- to 64-cell) clusters (from thawed sample; Support Protocol 3) or single cells (from dissociation) are dispersed uniformly onto the surface to avoid clumping and uneven coating of the flask.*

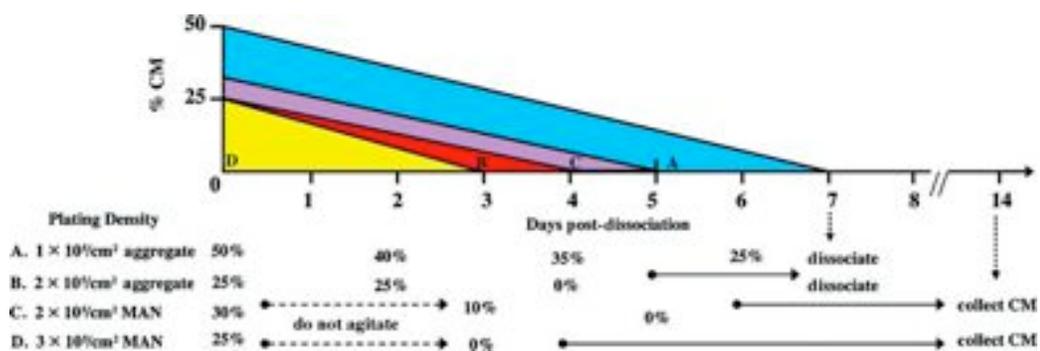
3. Once the cells have been fully distributed, allow the flask to incubate and equilibrate for 3 days without moving the flask for any reason from its original resting position.

*It is equally important that the incubator remain motionless and not be bumped or shaken in any way, or else the even hNPC coating will be disrupted and the organization of the MAN system will become disordered.*

*During the 72-hr hands-free period, the individual evenly spaced cellular clusters settle to the bottom of the flask, lightly attach, and migrate out over the surface, proliferating into each other, creating an interlinked lattice of three-dimensional adherent clusters displaying elongated processes that extend and connect each cellular island into a global multilayer adherent network (MAN). Perhaps the most important aspects of successful MAN culture setup are the initial plating conditions coupled with diligent patience and a steady hand during the initial week after hNPC thaw.*

*Throughout the process, adherent clusters can easily become detached by simply moving the flask; therefore, it is of utmost importance for the integrity of the culture system to absolutely avoid any movement of the flask or its content during the crucial aggregate-to-MAN transition process. Once hNPC clusters have detached, they will immediately merge with any other suspension aggregates they come into contact with (via integrins and secreted ECM proteins), thereby perturbing the essential spacing component of adherent growth. Even removal of the flask to view under the microscope disrupts the culture setup and should be avoided. For the same reasons, it is not prudent to supplement growth factors during this time; therefore, MAN cultures are started in 40 ng/ml bFGF to account for rapid degradation*

and resultant mitogen loss over the first 48 hr.



**Figure 2-2. Human neural precursor cell basic culture schedule.** Human NPC are grown as either lightly adherent aggregates or as multilayer adherent networks in 25-cm<sup>2</sup> flasks. Conditioned medium (CM) is gradually reduced from cultures as they progress and can be collected after MAN cultures reach ~75% confluence. Aggregate cultures are dissociated once a week or as growth parameters dictate, whereas MAN cultures can be cultured for up to 1 month before passaging.

- After 3 to 4 days of untouched growth, gently move the flask from the incubator to the sterile hood.
- Slowly tilt the flask up to 90°, then slowly rock backwards so that the flask is now upside-down, the CM is now facing downward on the top of the flask, and the cellular plane is facing upwards.

*You should be able to visibly identify exposed adherent clusters attached to the flask.*

- Aspirate all of the medium from the flask and, quickly but gently, add 8 ml fresh NB-B-27 complete medium containing 20 ng/ml bFGF, 10 ng/ml LIF, and 2  $\mu\text{l}/\text{ml}$  Normocin to the downward (noncellular) plane, being careful not to slosh medium onto the upper (cellular) plane, which would dislodge the lightly adherent cells.

*Do not allow the flask to dry out after the medium has been aspirated, as hNPC may begin to detach upon reintroduction of fresh medium to the culture.*

7. In a reverse motion, rock the flask back slowly to its original position, paying careful attention as the medium re-covers the adherent cells.

*During this process, it is absolutely imperative to reintroduce the fresh medium in a slow fluid motion to minimize waves as the medium spreads across the flask. Any major fluctuations or tapping of the flask can easily dislodge the clusters from their equally spaced positions, threatening the overall integrity of the MAN. No matter how careful you may be, there will always be a small percentage of cells that either did not attach or have detached during the feeding process. These floating cells will either reattach or can be removed from the culture at the time of the next feeding.*

8. After the medium has been changed, place the flask back into the incubator and repeat the process every 2 to 3 days as necessary to replenish growth factors (48-hr half-life) or replace metabolized medium (indicated by an orange acidic appearance).

*The literature and product datasheets support a general half-life for most of the growth factors used in this unit at 24 to 72 hr at 37°C in these medium formulations. The cells also utilize a large proportion, so we generally assume that the majority of the growth factors need to be replenished; therefore, we supplement according to the volume in the flask and adjust the concentration to the full concentration on the assumption that there is no growth factor remaining.*

*As the cultures expand, it will become necessary to alter the percentage of CM exchanged. During the first week, 75% to 100% of the medium should be exchanged to account for metabolized nutrients while maintaining adequate paracrine conditioning.*

*As cultures develop from week one onward, it will become essential to exchange 100% fresh NB-B-27 growth medium every 1 to 2 days to replenish the highly metabolized*

*nutrient stores and remove toxic metabolic by products. CM does not need to be added back in this case, as the high density-to-volume ratio leads to quick paracrine conditioning, adequate for immediate sustained survival. Furthermore, these fully developed MAN cultures can be utilized for the collection of high-quality CM (Support Protocol 4).*

*Over the next 2 to 3 weeks, MAN hNPC continue to proliferate and spread into a webbed culture, whereby adherent cellular islands will not only expand into each other but also proliferate in the vertical  $z$  dimension, creating the characteristic multilayer three dimensional appearance (Fig. 2-1 G-I).*

*As the MAN matures, it will develop into a highly mitotic (75% to 85%) confluent culture. Although clusters will continue to merge, there will always be demarcated areas on the flask surface where no hNPC grow; therefore, these cultures never attain the classic two-dimensional monolayer morphology.*

#### **2.4.1.2 Feed Multilayer Adherent Network (MAN)**

MAN cultures offer many time and growth advantages over classic aggregate or suspension sphere assays. Care should be taken to minimize sloshing of medium or excessive vibration that will detach the fragile network of cells. The basic rule for ease of use with this system is to minimize mechanical stress, especially at the edges of the flask, which can easily loosen the outer edges of the MAN, exposing the undersurface and resulting in uplifting of the entire sheet of adherent progenitor cells. Although these adherent networks of cells appear to be stably anchored to the flask, it takes relatively little force to disrupt their fragile connections. Furthermore, once detached, the cells will remain adherent in their networks and organize into large clumps, floating or partially attached to the remaining sheet of cells, which may become necrotic if not dissociated in ample time. Any cellular debris and insoluble salt residues that may develop from prolonged culture are removed by the methods described

below.

9. Slowly tilt the flask up to 90° and rock backwards so that the CM is facing downward on the top of the flask and the cellular plane is facing upwards. Carefully aspirate or collect conditioned medium

*See Support Protocol 4 for treatment of the conditioned medium.*

10. Gently rinse the flask once with 8 ml DPBS (for 25-cm<sup>2</sup> flask) or 12 ml DPBS (for 75-cm<sup>2</sup> flask) by expelling DPBS onto the downward (noncellular) plane at low speed, being careful not to slosh liquid onto the upper (cellular) plane, which would dislodge lightly adherent hNPC.

*Do not allow the flask to dry out after DPBS has been aspirated, as hNPC will begin to detach upon reintroduction of fresh media to the culture.*

11. In a reverse motion, rock the flask back slowly to its original culture position, paying careful attention as the DPBS re-covers the adherent cells.

*During this process, it is absolutely imperative to reintroduce the fresh DPBS in a slow fluid motion to minimize mechanical fluctuations as it spreads across the flask*

12. Repeat steps 9 to 11, transferring 8 to 10 ml (for 25-cm<sup>2</sup> flask) or 15 to 20 ml (for 75-cm<sup>2</sup> flask) fresh NB-B27 complete medium (containing 20 ng/ml bFGF, 10 ng/ml LIF, and 2 µl/ml Normocin) to each flask. Slowly move the flask to a humidified incubator at 37°C, 5% CO<sub>2</sub>.

#### **2.4.1.3 Dissociate Multilayer Adherent Network (MAN)**

13. When cultures are ready for passaging (see Critical Parameters and Troubleshooting), slowly tilt the flask upwards to 90°, then rock backwards so that the CM is facing downward on the top of the flask and the cellular plane is facing upwards. Aspirate or collect conditioned medium.

14. Gently rinse the flask once with 8 ml CMF-DPBS (for 25-cm<sup>2</sup> flask) or 15 ml CMFDPBS (for 75-cm<sup>2</sup> flask) by expelling CMF-DPBS onto the downward (noncellular) plane, being careful not to slosh CMF-DPBS onto the upper (cellular) plane, which would dislodge lightly adherent cells.

*Do not allow the flask to dry out after medium has been aspirated, as hNPC will begin to detach upon reintroduction of fresh liquids to the culture.*

15. In a reverse motion, rock the flask back slowly to its original position, paying careful attention as the CMF-DPBS re-covers the adherent cells. Repeat step 13 and aspirate.

*During this process, it is absolutely imperative to reintroduce the CMF-DPBS in a slow fluid motion to minimize mechanical fluctuations as it spreads across the flask.*

16. Gently add 3 to 5 ml (for 25-cm<sup>2</sup> flask) or 7 to 10 ml (for 75-cm<sup>2</sup> flask) of Accutase (prewarmed to 37°C, 10 min before use) to flask without disrupting the integrity of the cellular sheet (as described for CMF-DPBS rinse in steps 13 to 15).

17. Carefully transfer the flask into a 37°C, 5% CO<sub>2</sub> humidified incubator for 3 to 5 min (depending on density), minimizing any significant motion that will release the multilayer adherent network prematurely.

*The key to the successful dissociation of a MAN culture relies on learning to recognize the*

*following properties throughout the incubation in dissociation agent.*

*a. As the enzyme initially begins to break down cell-cell contacts, the adherent culture releases from the plastic dish from the outside in. Generally speaking, the outermost edges of the network will flap up and off of the dish, generating an organized sheet that eventually releases from the plastic dish below. If the dish is prematurely interrupted during this incubation process by moving the flask or sloshing the Accutase solution, the precise coordinated lifting of the multilayer adherent network is disturbed and subsequently leads to breakdown of the intact sheet of cells. Inadvertent disruption of the intact sheet can lead to gross clumping and compromise the integrity of cells as they dissociate.*

*b. In addition, prolonged exposure to enzymes can puncture the cell membrane and render hNPC extremely vulnerable to mechanical shearing, resulting in lysis and release of DNA into the cell suspension. The results of enzyme overexposure are visibly apparent, as evidenced by increased viscosity of the cell suspension accompanied by discernibly large floating aggregates. These aggregates have a propensity to float to the top of the cell suspension and are characterized by their sticky, slimy properties that render them problematic in culture as they accrue and amass live cells on the surface. As the aggregates continue to bind live hNPC, they become heavier and eventually fall by gravity from the top of suspension to the bottom, thus allowing for removal from the remaining population. The overall result of enzyme overexposure is decreased hNPC recovery; therefore, it is imperative to time the enzymatic process and visually inspect the flask after 3 to 3.5 min, to monitor the dissociation progress closely.*

*c. During the 3- to 5-min incubation process, the MAN layer will gradually detach completely*

*from the underlying flask, effectively shrinking into an intact rectangular sheet, resembling a miniature compacted version of the original MAN. The exact timing for completion of this process is variable, but should be minimized to account for overexposure. In general, the entire sheet should be detached and shrunk into the center of the flask for at least 1 to 3 min before the desired stage of dissociation is attained. Note that this is an extremely time-sensitive process. Lesser incubation times will result in incomplete dissociation of larger hNPC clusters, requiring additional cycles, ultimately leading to increased clumping and subsequent cell death.*

18. After 3.5 to 5 min, when the MAN displays the above characteristics, gently transfer the flask to a sterile hood, paying special care to retain the free-floating cellular sheet in its intact form for easy removal.

*The intact sheet is extremely fragile and will most likely begin to dissociate as the flask is moved. Try to retain the sheet in as many large pieces as possible. Furthermore, lower density cultures will not retain the structural integrity that their mature MAN counterparts display.*

19. Carefully tilt the flask so that the sheet of cells aggregates to the bottom corner of the flask with gravity. With a 5-ml pipet, carefully suck up the concentrated network of cells in 1 to 3 ml of the Accutase solution and transfer to a 15-ml conical tube.

*It should be possible to reclaim the cells into a small volume without extensive single cell dissociation or disruption of the cellular sheet. The remaining Accutase should appear clear and may contain a few smaller cell clusters.*

20. Gently triturate contents of the conical tube with a 5-ml pipet attached to a pipetting aid (e.g., Drummond) on medium speed (five to seven times) to break the cell suspension into

smaller floating cellular aggregates.

*Be very careful not to over-triturate, as the cell suspension is extremely fragile at this stage.*

21. Using the same 5-ml pipet, immediately triturate the remaining contents of the flask to break up remaining clusters, gently but thoroughly, paying extra attention to the removal of adherent hNPC at the edges of the flask where they tend to attach preferentially and with increased strength. Transfer the contents of the flask to the previous conical tube.

22. Continue trituration of hNPC inside the conical tube to break the cells up into smaller clusters by gently expelling the cell suspension at a 45° angle against the wall of the conical tube at medium speed (8 to 10 times).

23. If necessary, recap the conical tube and incubate in a 37°C water bath for 1 to 2 min more with constant swirling to avoid clumping of aggregates at the bottom of the tube and reduce accumulation of sticky DNA from lysed cells.

*It is very important to ensure the hNPC do not aggregate and begin clumping during the dissociation process; therefore, care should always be taken to continuously swirl or triturate the cells during steps 20 to 23.*

24. Using a 1000- $\mu$ l extended-length pipet tip with a standard automatic pipettor set to 750  $\mu$ l, slowly triturate hNPC suspension at a 45° angle against the wall of the conical tube at a consistent rate.

*Excessive or high-rate trituration against the plastic wall is not well tolerated at this stage.*

*We recommend slow to medium trituration at a position near, but not touching directly against the wall of the conical tube (five to ten times or until large clumps are no longer*

*visible and the dissociated solution has a homogenous milky and sandy appearance). Ideally passaged cultures will be fully dissociated into single cells, >95% viable, and free of floating aggregates if the time of initial Accutase exposure was within the correct window (step 17), cells are not allowed to aggregate, and trituration remains moderate and minimal. Cell clusters will readily stick to the meniscus (~750- $\mu$ l line) of the pipet tip.*

25. To recover cells that have stuck to the meniscus, reset the plunger from 750  $\mu$ l to 1000  $\mu$ l (with tip remaining intact). Rinse the 1000- $\mu$ l tip once with 1000  $\mu$ l NB-B-27 complete medium to dislodge residual clusters, and transfer the contents to a new 15-ml conical tube containing 10 ml fresh NB-B-27 complete medium containing 20 ng/ml bFGF, 10 ng/ml LIF, and 2  $\mu$ l/ml Normocin (prewarmed to 37°C) to inactivate the reaction.

26. Inactivate fully dissociated preparation from step 24 by adding it to the 10 ml medium in the conical tube from step 25.

*Variability in hNPC culture densities and morphology will dictate the specific timing and rate of dissociation for each culture. As a result, it is often the case that a small percentage of undissociated cell clusters remain and require a second round of enzymatic treatment, while the majority of cells are fully dissociated and ready to be inactivated and released from enzymatic shock.*

27. To process partially dissociated cell suspensions, place the conical tube vertically for 1 to 2 min until the visible cellular clusters have settled by gravity to the bottom. Carefully transfer the top portion of supernatant containing dissociated cells to the previously inactivated cell suspension. To the remainder of undissociated hNPC, add 1 ml fresh prewarmed Accutase, triturate twice, and repeat steps 24 to 26.

28. Transfer the appropriately dissociated cell suspension to the previously inactivated 10 ml hNPC suspension from step 26.

*In rare cases, some clusters may remain after the second round of dissociation (often seen in necrosis) and are considered behaviorally abnormal and subsequently discarded.*

*CAUTION: Overexposure to any dissociating agent will cause significant cell death and deter growth from lysed hNPC. The solution will become more viscous when this occurs. Thus, the procedure should be optimized to break up the cell clusters, while minimizing the amount of time in the dissociation agent. Generally, the larger the flask, the more dissociation agent that will be needed, which means more cell death and greater difficulty in controlling the timing of the process. We recommend 25-cm<sup>2</sup> or 75-cm<sup>2</sup> flasks for optimal conditions.*

29. Centrifuge the cell suspension for 4 min at 400 °— g, room temperature. Carefully aspirate the supernatant.

*Adherent cultures exhibit a highly branched, polarized cellular morphology, and unfortunately many of these delicate processes are cleaved by dissociating agents and mechanical stress, resulting in a greater amount of cellular debris. As a result, an additional rinse and centrifugation with 10 ml of either CMF-DPBS or Neurobasal medium (Invitrogen) is recommended to remove any problematic residual debris.*

30. Resuspend the hNPC pellet in the conical tube with 1 ml fresh NB-B-27 complete medium using an extended-length 1000- $\mu$ l pipettor and tip, gently triturating five to seven times to thoroughly liberate the cell pellet.

31. Count viable cells using a hemacytometer and trypan blue (*UNIT IC.3*) for correct

replating density.

32. After counting, add 8 ml CM (for a 1:2 dilution) to the conical tube, adjust for the desired final volume of fresh NB-B-27 complete medium to CM ratio accordingly (i.e., 8 ml fresh NB-B27 medium for 50% CM final), bring cells to desired density, and replate into new 25-cm<sup>2</sup> flasks.

*In general, more concentrated splits survive and proliferate more effectively than their diluted counterparts. As a guideline, a 25-cm<sup>2</sup> flask containing 1–3 × 10<sup>6</sup> cells is fed 25% to 50% CM, and 4 × 10<sup>6</sup> cells do not require CM as they quickly condition the medium due to high density.*

33. Add bFGF and LIF to achieve a final concentration of 20 ng/ml and 10 ng/ml, respectively. Gently swirl contents of flask horizontally to evenly disperse hNPC and place in a humidified incubator at 37°C, 5% CO<sub>2</sub>.

*Subsequent culturing methods will depend on the density of cells plated and method for further expansion.*

34. Passage MAN cultures.

*Typically, the growth parameters of hNPC MAN cultures dictate passaging once every 1 to 2 months depending on the original plating density and desired confluency. We typically split MAN cultures at a 1:2 dilution for 3–4 × 10<sup>6</sup> cells/25-cm<sup>2</sup> flask of mature 65% to 75% confluent culture, or 1:4 for 5–10 × 10<sup>6</sup> cells/25-cm<sup>2</sup> flask of very mature 80% to 90% confluent extremely high-density 2-month-old cultures, as they contain many more cells per flask than a typical aggregate culture where high density cannot be achieved at the cost of fusion, large globular aggregate formation, and ensuing necrosis.*

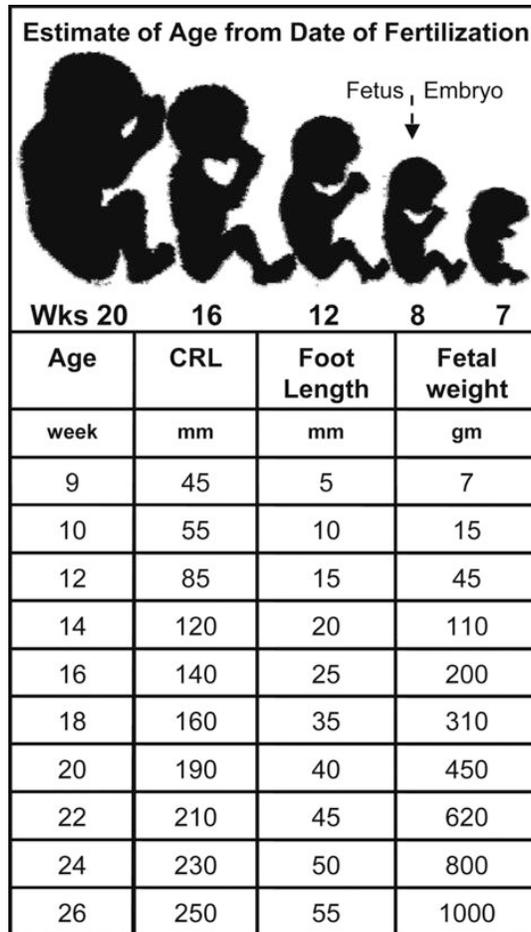
*We consider the above modifications of the enzymatic process, specifically the precisely timed controlled release of the entire MAN as an intact sheet, to be one of the key components of successful passaging and subsequent expansion of hNPC using this assay. Consistent high viability and overall health of the resultant hNPC preparations coupled with the intrinsic quantitative qualities of the assay (i.e., increased population doubling rate, apparent increase in proliferation capacity for >100 passages without senescence or decrease in rate of replication, and decreased cost in consumables and personal time) all mark the overall utility and advantages for employing the MAN assay to obtain long-term expansion of large quantities of undifferentiated hNPC.*

*MAN cultures can also be processed by traditional methods used for aggregate cultures. Simply triturate adherent cells thoroughly from the flask and proceed as described for aggregate cultures (Alternate Protocol 1). It should be noted that enzymatic dissociation times will be greatly enhanced, requiring multiple rounds of gravity-based cluster separation, enzymatic treatment, and subsequent centrifugation cycles. Unfortunately, this procedure results in significant cell death (60% to 70% viability) in even the most skilled hands, and should only be employed when cells are accidentally detached by mechanical force. In these cases, a second rinse and centrifugation step should be added prior to final plating.*

#### **2.4.2 DERIVATION OF HUMAN FETAL NEURAL STEM/PRECURSOR CELLS**

Fetal spatial features and their specific neuroanatomical coordinates are used to determine the cadaver's specific stage of CNS development and dictate the exact location for tissue dissection. Proficiency in fetal neuroanatomy is essential for efficient assessment and subsequent resection of specified CNS regions. We, along with others, have described various methods for the derivation of hNPC. Here, we detail the methodology we have successfully employed to isolate and expand fetal forebrain periventricular zone human NPC.

The investigator typically comes into possession of fetal tissue post-mortem as cadaveric material (Prior to 23 weeks of gestation, a fetus is non-viable outside of the womb). Between the eighth and ninth week of development, the embryo undergoes several distinct developmental transitions. Both the feet and hands lose their webbing and become separated into distinct digits, and the stubby tail disappears. At 9 weeks post-fertilization, by convention, the embryo is called a fetus. Estimation of fetal age can be extremely tricky without prior knowledge of the last normal menstrual period or day of fertilization (which is rarely known by clinicians); however there are a few decent methods to gauge estimated fetal age and developmental stage. Fetal entities are most accurately staged using pre-term ultrasound measurements, however, these vital records are rarely granted to the investigator, therefore, we must rely on post-mortem methods including crown to rump length (CRL), foot length, head to trunk, cheek to cheek, and fetal weight for quantitative analysis [Figure 2-3]. When utilized individually, these methods are often inaccurate, as fetal specimens are affected by stretching. A combination of strategies, however, can be used to accurately determine fetal age within several days of fertilization.



**Figure 2-3. Fetal developmental staging.** Determining the relative fetal developmental age is crucial for proper anatomical assessment prior to dissection of the brain. Specifically, Crown Rump Length (CRL) is used as an appropriate measure for staging post-mortem fetuses. All measurements are averages based on deviations from Streeter's original tables from fixed fetuses based on the *time from fertilization* and intended as general reference points to gauge developmental staging of the brain. Gestational age begins 2 weeks after fertilization age. Fetuses born prematurely after 22 weeks may survive with artificial support but have limited capacity due to underdevelopment of both the respiratory and nervous systems.

The most common and useful parameter for aging and developmental analysis has remained CRL due to its high correlation with definitive gestational changes. Starting at week 9 (from date of fertilization), the head contributes to almost half of the CRL, which more than doubles by the end of week 12. From weeks 12-16, fetal growth is rapid, albeit the head remains relatively stable, contributing less to the total CRL than during weeks 9-12.

Importantly, skull and long bone ossification begins to set in and should be considered when deriving hNPC from fetuses of this developmental period. By the end of the 16<sup>th</sup> week, the eyes are repositioned anteriorly from anterolateral and overall fetal growth slows down considerably.

As development proceeds over the next four weeks, the CRL continues to increase by about 50 mm, and the lower limbs reach their final proportions. By twenty weeks, hair and genitalia begin to develop and the mother will start to experience fetal movements, or *quickenings*. It is at this time that the fetal entity begins to fully develop and take on substantial weight. Furthermore, between weeks 21 and 25, the fetus develops blood filled capillaries, fingernails, and rapid eye movements reminiscent of newborns. A thorough and accurate assessment should be made to properly identify the age of the fetus before dissection. All procedures must be carried out in strict accordance with federal regulations. For example, in the United States, federal law mandates that the latest time a woman may legally terminate pregnancy to be roughly 24 weeks. Typically, cadavers become available to the investigator at 9-15 weeks of age, a time at which landmarks are admittedly hard to identify.

*NOTE:* Use of human fetal cadaveric CNS must follow all safety and bioethical guidelines, including but not limited to full informed consent, IRB approval, and strict adherence to all state and federally mandated laws and guidelines for the ethical use and treatment of patients or specimens derived thereof.

*NOTE:* Perform all procedures aseptically in a sterile Biosafety Level 2 hood. Sterilize all surgical tools in a hot bead sterilizer or autoclave (121°C, 2 hr), or by gas sterilization. During the procedure, place all of the tools in fresh 70% ethanol when not in use. Immediately following removal from ethanol, briefly rinse twice in fresh sterile DPBS (Mediatech, cat. no.

21-031-CM).

### ***Materials***

Fetal tissue

10% (v/v) formalin (optional)

Enzymes for tissue dissociation (optional): e.g., Accutase, trypsin-EDTA, PPD (papain-protease-DNase I)

Fetal bovine serum (FBS; optional)

NB-B-27 complete medium (see recipe)

Basic fibroblast growth factor (bFGF; Millipore, cat. no. GF003)

Leukemia inhibitory factor (LIF; Millipore, cat. no. LIF1010)

Normocin (InvivoGEN, cat. no. ant-nr-1)

Epidermal growth factor (EGF; Millipore, cat. no. 01-107)

Surgical equipment, including scalpel, sterile

15-ml conical tubes

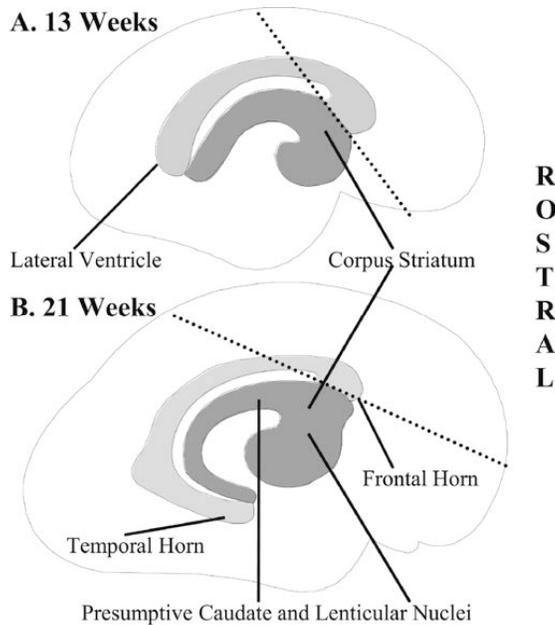
Battery-powered pipetting aid (e.g., Drummond Pipet-Aid XP)

Additional reagents and equipment for counting viable cells by trypan blue exclusion (*UNIT 1C.3*)

#### **2.4.2.1 Isolate and digest human fetal periventricular zone**

1. Stage the fetus using neuroanatomical coordinates, open the head cavity, and remove the brain.

2. Cut sagittally across the midline to separate the cerebral hemispheres then cut again coronally from frontal to occipital poles (Diagram 2-1).



**Diagram 2-1. Fetal cerebral hemispheres and ventricular zone development**

Sagittal anatomical rendering of the medial surface of the fetal brain at 13 and 21 weeks after fertilization. As the brain develops, the cerebral hemispheres expand and meet at the midline, giving rise to the corpus striatum and shaping the cerebrospinal fluid filled, C-shaped cavities known as the lateral ventricles (**A**, **B**). As the cerebral cortex begins to differentiate, the corpus striatum eventually divides into the horseshoe shaped caudate nucleus and the bulbous lentiform nucleus, which houses the putamen and globus pallidus (**B**). At 13 weeks, the surface of the cerebral hemispheres are smooth and underdeveloped, highly resembling lower order mammals, however by 21 weeks, patterned convolutions (gyri) and grooves or furrows (sulci) form complex folds increasing the total surface area with little change in cranial parameters. In addition, the lateral sulcus begins to narrow as the insula becomes buried (not pictured). We generally dissect along the *dashed line* and remove the periventricular zone adjacent to the head of the caudate nucleus.

3. Select the brain slice containing the region of interest for dissociation.

*Optional: Fix the remaining tissue in 10% (v/v) formalin for a more extensive neuropathological examination.*

4. Carefully scrape the ventricular wall and adjacent subventricular zone region from the

forebrain section with a surgical scalpel. Delicately mince the dissected tissue into small pieces with the scalpel blade.

5. Transfer the tissue pieces into a 15-ml sterile conical tube that contains 6 ml cold NB-B-27 medium, 20 ng/ml bFGF, 20 ng/ml EGF, and 4  $\mu$ l/ml Normocin.

6. Place the conical tube vertically and allow the tissue to pellet by gravity (1 to 2 min), aspirate supernatant carefully, and rinse three times, each time with 8 ml cold medium.

7. After final rinse, resuspend the tissue in 8 ml cold medium.

8. Gently triturate the fetal tissue suspension (10 to 15 times) with a 5-ml pipet attached to a pipetting aid (e.g., Drummond Pipet-Aid XP) at medium speed against the wall of the 15-ml conical tube to further dissociate the tissue into a homogenous milky solution.

*The cell suspension will contain both single cells and a few small cellular clumps. Try to avoid introducing air bubbles during the trituration process. It is important that the primary tissue not be overzealously digested into a single-cell suspension, due to the subsequent damage incurred by mechanical stress on the progenitor fraction. CNS tissue from young fetal brains is softer than that from fully developed myelinated adult brains; therefore, later-stage CNS preparations include the addition of an enzymatic agent such as Accutase, trypsin-EDTA, papain-protease-DNase I (PPD), dispase, or any commercially available reagent, according to the manufacturer's instructions, to efficiently dissociate primary cultures before their initial plating. In general, enzymatic fetal tissue dissociation averages ~5 to 10 min, while adult tissue can take upwards of 45 to 90 min to generate the desired breakdown of brain tissue.*

9. Remove large undissociated tissue bits remaining after the initial trituration by allowing

them to settle by gravity (2 to 3 min), then collect the suspension of cells in the upper supernatant. Dissociate remaining undissociated cell clumps again as in the steps above and pool together with the originally dissociated cell suspension.

10. Inactivate enzymatic preparations by diluting them 1:5 in fresh prewarmed NB-B27 medium and centrifuge for 5 min at  $400 \times g$ , room temperature. Remove supernatant and retain pellet.

#### **2.4.2.2 Establish primary hNPC cultures**

11. Following primary dissociation, bring the cell suspension to working volume in 8 ml prewarmed NB-B-27 medium with 20 ng/ml bFGF, 20 ng/ml EGF, and 4  $\mu$ l/ml Normocin at a final density of  $1 \times 10^5$  cells/cm<sup>2</sup> in one 25-cm<sup>2</sup> flask and place in a humidified 5% CO<sub>2</sub> incubator at 37°C.

*Primary cultures plated onto tissue culture treated flasks will generally produce mixed aggregate and adherent cultures. Primary cell suspensions may also be plated onto fibronectin-coated tissue culture-treated flasks for monolayer-like (two-dimensional) adherent cultures.*

12. Determine cell viability using either the propidium iodide or trypan blue exclusion assay and a hemacytometer.

*Sticky cellular debris and small undissociated neural clumps may make this process difficult initially.*

13. *Optional:* Add 0.1% to 1% (v/v) fetal bovine serum (FBS) at the time of initial derivation to enhance initial NPC expansion efficiency, promote adhesion, and decrease overall cell

death with a relatively low risk of differentiation.

*CAUTION: Using FBS may introduce unwanted variability. Serum components are removed after a short period of time and replaced with a defined, serum-free medium so as not to potentiate long-term side effects on primary hNPC cultures. In some cases, it is desired that newly derived stem cell lines be established utilizing serum-free protocols so as not to introduce animal proteins into culture.*

14. Incubate cells. At a time point 12 to 48 hr after plating, rinse any serum-containing cultures twice with 10 ml DPBS and transfer cultures to serum-free conditions in NB-B-27 medium containing 20 ng/ml EGF, 20 ng/ml bFGF, 10 ng/ml LIF, and 4  $\mu$ l/ml Normocin. Continue incubation.

15. At a time point 3 to 4 days after the primary plating, supplement cultures by carefully removing the top half of medium from each flask, termed conditioned medium (CM), and replace with fresh NB-B-27 complete medium containing 40 ng/ml EGF, 40 ng/ml bFGF, 20 ng/ml LIF, and 8  $\mu$ l/ml Normocin for the final working concentration of 20 ng/ml EGF, 20 ng/ml bFGF, 10 ng/ml LIF, and 4  $\mu$ l/ml Normocin.

*These final concentrations are based on the assumption that the growth factors have been completely deleted by this point.*

16. For more efficient recovery, remove the CM containing free-floating aggregates and small clumps of primary tissue and transfer the contents to a new flask. Triturate the cell suspension thoroughly to redissociate the remaining clumps, and supplement with fresh growth factors and antibiotics by the above procedure.

*Alternatively, centrifuge suspension aggregates and debris for 3.5 min at 400° —g, aspirate,*

*and either add the cells back to the original parent culture flask for further expansion or replate the primary cultures into 8 ml fresh pre-warmed NB-B27 complete medium containing 20 ng/ml EGF, 20 ng/ml bFGF, 10 ng/ml LIF, and 4  $\mu$ l/ml Normocin.*

17. Repeat steps 15 and 16 throughout the first few weeks of primary culture.

*Initially, hNPC will proliferate throughout the flask as a mixture of adherent and free floating aggregates and can be detached from the culturing vessel through repeated trituration. We stress the inclusion of adherent monolayer-like hNPC within primary cultures during the initial hNPC expansion stage. As cultures mature, adherent hNPC cultures may also spontaneously give rise to a few spherical balls. These aggregates detach from the initial colony and continue to expand and self-renew as free-floating suspension cultures as well.*

18. After several weeks, select the hNPC cultures that proliferate in a morphologically relevant manner and dissociate into single-cell suspensions or small clumps (3 to 8 cells/clump) with Accutase or cell dissociation buffer (CDB)/cellstripper. Dissociate when cellular aggregates are larger than 12 to 15 cells in diameter and can no longer be mechanically separated by simple trituration or when adherent cultures become greater than 75% confluent. Pool both adherent and free-floating cells and discard any remaining large clumps that do not readily dissociate.

19. Replate hNPC at a 1:1 or 1:2 ratio as either multilayer adherent aggregates or as suspension aggregates in NB-B-27 complete medium, 20 ng/ml EGF, 20 ng/ml bFGF, 10 ng/mL LIF, and 4  $\mu$ g/ml Normocin for 2 more weeks.

20. Exchange one-half of the culture medium as described in step 15 every 2 to 3 days to

replenish growth factors and antibiotics. Dissociate and replate cultures (1:1 or 1:2) once per week or as necessary. After 2 weeks, exclude LIF and EGF for mitogen selection.

#### **2.4.2.3 Mitogen-select primary hNPC cultures**

After 2 to 4 weeks of primary expansion, undifferentiated hNPC colonies will proliferate and establish a healthy culture of precursors. At this point, successful cultures are subjected to a 10-week sequential growth factor selection process utilizing parameters of growth rather than markers alone to select for the proliferative EGF/FGF responsive population of cells.

21. Expand hNPC as a mixed population of both adherent clusters and free-floating aggregates in NB-B-27 complete medium containing 20 ng/ml bFGF alone (and 2  $\mu$ l/ml Normocin) for 2 weeks with (1:1 or 1:2) dissociation once per week throughout the selection process as dictated by size exclusion and morphological parameters described above in step 18.

22. After 2 weeks, omit bFGF and supplement the medium with 20 ng/ml EGF alone (and 2  $\mu$ l/ml Normocin) for 2 weeks.

23. Maintain the bFGF/EGF 2-week rotation schedule for two to three sequential rounds (10 weeks) and complete after the final bFGF-alone cycle.

24. After the final selection process, a few primary hNSC/hNPC cultures will continue to proliferate and display appropriate morphology; dissociate these cultures and pool together into NB-B27 complete medium containing 10 ng/ml LIF, for a final hNPC complete basal maintenance medium composed of NB-B-27 growth medium containing 20 ng/ml bFGF, 10

ng/ml LIF, and 2  $\mu$ l/ml Normocin for secondary hNPC expansion.

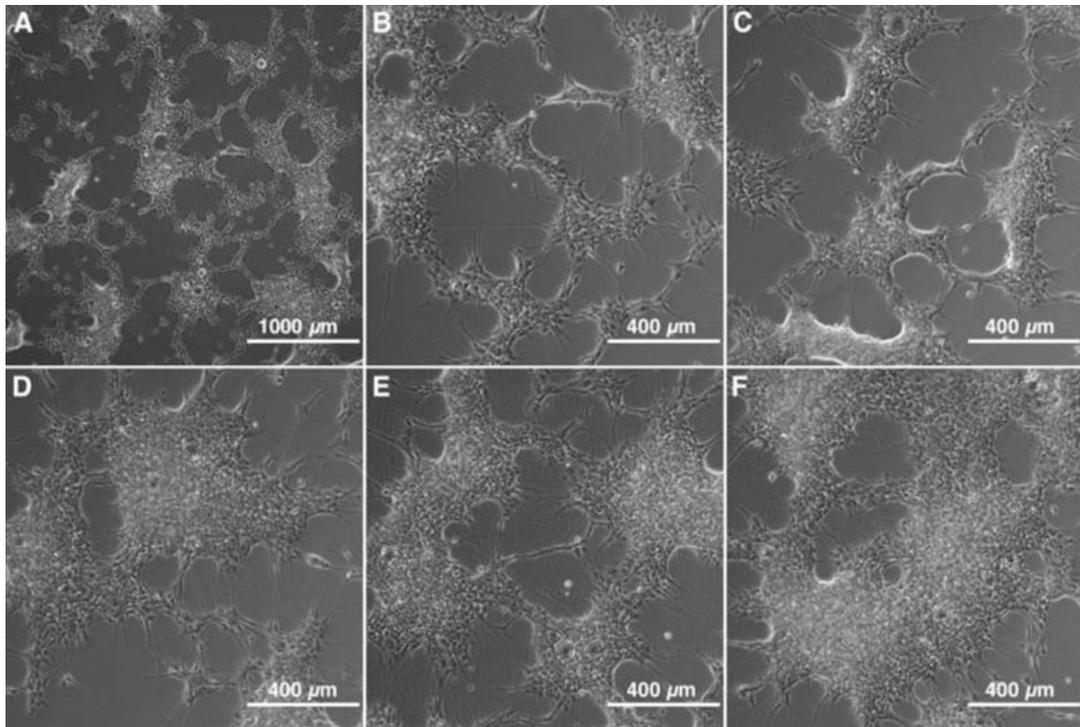
### **2.4.3 Replating Dissociated hNSC on Extracellular Matrix (ECM) as Adherent Two-Dimensional Monolayer Cultures.**

In addition to the MAN assay described in detail here, hNPC can also be replated onto a variety of extracellular matrix (ECM) components at  $1-2 \times 10^6$  cells/25-cm<sup>2</sup> flask (maximum of  $2-3 \times 10^6$  cells/25-cm<sup>2</sup> flask) to induce attachment for more traditional two-dimensional, adherent monolayer growth parameters (Fig. 2-4). As with MAN cultures, ECM attachment should not be utilized for low-density cultures where very few cell-cell contacts are present. The resulting cultures will likely become post-mitotic and differentiate prematurely. We prefer to expand primary hNPC lines without additional biological components, but we also recognize the utility and beneficial growth parameters that many ECM components confer in hNPC culture, especially when assaying and analyzing cells for migration and immunocytochemistry. That being said, not all ECM components are created equal, and each hNPC line will have its own particular characteristic adhesion properties. In our hands, hNPC tend to adhere to a variety of ECM proteins displaying a continuum for strength of adhesion—in order from weakest to strongest adhesion, fibronectin (human or mouse), laminin (human or mouse), Matrigel, collagen, and vitronectin. We recommend trying Millipore's ECM cell culture optimization assay to determine the optimal ECM protein and concentration desired for the specific growth parameters chosen. In addition, a number of commercially available cell-binding enhancement solutions (Cell Bind) or specially scaffolded substrates (Cell Web, Corning) are also available, with a variety of binding properties to circumvent the use of biological attachment substrates. Furthermore, pre-coating flasks with electrostatically charged molecules such as poly-D-lysine or poly-L-ornithine in combination with extracellular matrix proteins provide a secondary level of support, often conferring an

additional degree of adhesion. One warning is that poly-D-lysine should not be used for experiments involving electrophysiology, as it may interfere with ion-channel function.

In our hands, prolonged enhanced adhesion and exposure to matrix signaling molecules can have significant effects on hNPC phenotypic variation and related changes in cellular differentiation profile. For example, fibronectin supports a similar lightly adherent mode of growth to freshly dissociated MAN cultures on non-coated tissue culture-treated flasks, with the added benefit of slightly enhanced adhesion, quicker attachment, and higher rates of attachment. Laminin, likewise, retains many of the essential properties of the undifferentiated MAN, with the caveat that the initial adhesion is stronger, resulting in more flattened, monolayer-like, two-dimensional, multi-polar progenitor colonies. In slight contrast, Matrigel, a soluble basement membrane extract of the Engelbreth-Holm-Swarm tumor, which is composed mainly of laminin as well as collagen IV, heparin sulfate proteoglycans, and entactin, but contains trace amounts of the platelet-derived growth factor (PDGF), nerve growth factor (NGF), insulin-like growth factor-1 (IGF-1), and TGF- $\beta$ , supports exuberant growth of highly mitotic, extremely adherent, bipolar and multipolar neural precursors that will self-assemble into a highly dynamic neural niche (Watt and Hogan, 2000; Palmer, 2002; Wurmser et al., 2004; Lathia et al., 2007) composed of a heterogeneous population resembling type A, B, and C cells of the subventricular (SVZ) niche (D.R. Wakeman, unpub. observ.). Furthermore, substrates such as collagen IV and vitronectin bind hNPC, conferring an exceptional propensity for attachment, but typically at the cost of mass cellular differentiation. These findings introduce a secondary criticism of ECM components, in that ECM molecules naturally guide neuronal migration (Thomas et al., 1996; Murase and Horwitz, 2002, 2004; Labat-Robert and Robert, 2005; Flanagan et al., 2006; Hall et al., 2008) and are thought to play a critical role in differentiation of hNPC *in vivo*. As a result, culturing hNPC in the presence of these molecules *in vitro* may actually trigger primary differentiation of hNPC and

an irreversible exit from the cell cycle. It is important, therefore, to choose an ECM accordingly and with respect to the specific assay of interest, as long-term cultures will adapt to their environment and may not continue to behave as true undifferentiated hNPC. We are comfortable with prolonged undifferentiated culture and expansion on either human fibronectin or human laminin (Ray et al., 1993; Vicario-Abejon et al., 1995; Walsh et al., 2005; Flanagan et al., 2006; Ray and Gage, 2006; Hall et al., 2008) and temporary undifferentiated growth on Matrigel for 1 to 2 weeks. More adherent substrates such as vitronectin and collagen type IV are best utilized for differentiation assays. Specific brands and lots of ECM vary; therefore, proper testing is essential to determine individual growth parameters. It is worth noting that enzymatic lifting and dissociation of hNPC grown on strongly adherent ECM components generally require longer incubation times and often generate 10% to 20% cell death accordingly, due to the increased prevalence of fragile projections.



**Figure 2-4. Extracellular matrix confers two-dimensional monolayer phenotype in hNSC cultures.** Human NSC (HFB-2050) were plated onto tissue culture-treated flasks previously coated with a combination of poly-D-lysine and the extracellular matrix protein fibronectin. After 7 days in vitro, hNSC attain a similar composition and phenotype as MAN cultures, although they flatten and proliferate in a more two-dimensional manner in contrast to the three-dimensional architecture of MAN cultures (A-C). After 2 weeks, individual aggregate clusters are indistinguishable from each other, and begin to merge into a confluent layer of hNSC (D,E). In contrast to their MAN counterparts, these cultures will form a classic monolayer and lose their honeycomb appearance (F).

## 2.5 LABELING hNPC PRE-TRANSPLANTATION

In order to identify transplanted donor cells, hNPC must be pre-labeled chemically or genetically with a definitive nontransferable marker. One can also utilize human-specific antibodies, such as huNuc, after transplantation to recognize donor-derived cells. For proper validation, hNPC should be identified with at least two of these markers and preferably three to ensure that results are not simply false-positive artifacts. We generally prefer colocalization of BrdU and a reporter gene with at least one human-specific epitope to locate successful donor grafts. Upon implantation into the mammalian CNS, HNPC may undergo one to three

rounds of division before becoming post-mitotic; therefore, non-integrating labels will not become too diluted for later detection. Theoretically, thymidine analogs, iron particles, and lipophilic dyes become diluted by a factor of  $\frac{1}{2}$  for every symmetrical cell division; therefore, these markers can become diluted below standard detection levels within five to six cell divisions. Careful selection of labeling method should be based on the specific assay of interest.

### **2.5.1 Labeling hNPC with BrdU**

Human NPC that have not been genetically labeled are preincubated with the S phase, DNA-intercalating thymidine analog, bromodeoxyuridine (BrdU) for proper postmortem graft identification. Both monoclonal (Gratzner, 1982) and polyclonal antibodies have been raised to detect BrdU using immunofluorescence and multiphoton confocal microscopy for graft analysis. As a result, BrdU was reinforced as a popular prelabeling technique for grafting proliferative cellular substrates. Although BrdU only labels at best 42% to 50% (neurospheres) to 74% to 82% (MAN cultures; Fig. 2-5) of donor hNPC nuclei (in our experience), presents a variety of false-positive artifacts (Rakic, 2002a,b; Burns et al., 2006), and can be highly toxic when administered for extended periods of time (Caldwell et al., 2005), it has remained one of the most highly used pre-transplantation labeling methods to verify donor cell origin (Dolbeare, 1995, 1996; Carbajo et al., 1995; Carbajo-Perez et al., 1995).

#### ***Materials***

NB-B-27 complete medium (see recipe)

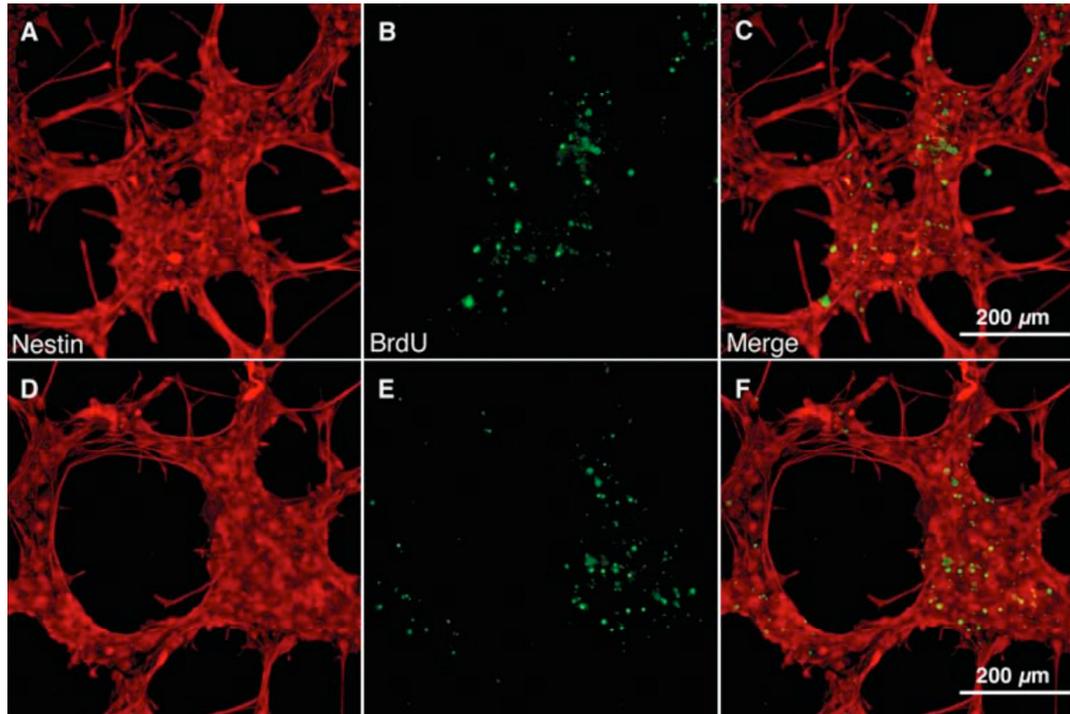
Normocin (InvivoGEN, cat. no. ant-nr-1)

Leukemia inhibitor factor (LIF; Millipore, cat. no. LIF1010)

Basic fibroblast growth factor (bFGF; Millipore, cat. no. GF003)

5-bromo-2\_-deoxyuridine (BrdU; Sigma, cat. no. 59-14-3; see recipe)

25-cm<sup>2</sup> tissue culture flasks



**Figure 2-5. hNSC MAN cultures incorporate BrdU.** Human HSC (HFB-2050) grown as multilayer adherent networks express nestin (A,D) extensively and readily incorporate BrdU (B,E), indicating that they remain in a highly proliferative, immature state throughout MAN culture. Of interest, cells within the clusters appear to proliferate preferentially in comparison to their peripheral counterparts (C,F), indicating a possible niche component within each adherent cluster.

1. To efficiently label cells, dissociate hNPC 48 to 72 hr prior to transplantation and replate as described in Basic Protocol 1 or Alternate Protocol 1, as applicable, using fresh NB-B-27 complete medium containing 2  $\mu$ l/ml Normocin, 10 ng/ml LIF, 20 ng/ml bFGF, and 10 to 20  $\mu$ M BrdU (added from 1000 $^\circ$  — stock).

*BrdU is highly toxic to low-density hNPC cultures; therefore, we recommend plating cells at no less than 2 $^\circ$  — 10<sup>6</sup> cells/25 cm<sup>2</sup>.*

CAUTION: *BrdU acts by incorporation in the place of thymidine during DNA synthesis, and thus may cause birth defects or heritable genetic effects. Be extremely careful when handling BrdU.*

2. Allow cultures to equilibrate and return to homeostasis in a 37°C, 5% CO<sub>2</sub> humidified incubator.

*Low-density cultures fail to equilibrate properly and display extremely slow division rates, impeding efficient labeling during S-phase.*

3. Every 24 to 36 hr, replenish the medium by adding fresh 1000° — BrdU to 1° — final, LIF to 10 ng/ml final, and bFGF to 20 ng/ml final. Lightly triturate.

*Gentle trituration of cells is absolutely essential to reduce merging of cell clusters and formation of premature aggregates before transplantation. Cells are extremely fragile when incubating in BrdU; therefore, slow trituration is recommended to avoid shearing.*

4. After a total of 48 to 72 hr, prepare hNPC for transplantation as described (see Basic Protocols 6 and 7).

*In our hands, 40% to 50% (neurospheres) and 70% to 80% (MAN) of the total hNPC population will be labeled after 48 to 72 hr. It should be noted that longer incubation times are notorious for introducing false positives into nondividing cells. Furthermore, hNPC do not proliferate well past 3 to 4 days in media that include BrdU (D.R.Wakeman, unpub. observ.), indicating a time threshold for toxicity.*

### 2.5.2.1 Lentiviral Infection of hNPC

Traditional methods for prelabeling NPC prior to transplantation require harsh DNA intercalating thymidine analogs such as BrdU and CldU, which have been shown to result in great underestimation of overall engraftment success and create a variety of false positives when administered for extensive periods of time. Generation of independently labeled fluorescent reporters, animals, and cell lines eliminates the need for these toxic compounds while increasing both cell viability and engraftment efficiency (Shimomura et al., 1962; Chalfie et al., 1994, 1995; Ward et al., 1998; Zhang et al., 2002; Tsien, 2003; Vintersten et al., 2004; Shaner et al., 2005, 2008; Shimomura, 2005; Giepmans et al., 2006). In addition, fluorescently labeled donor cells can be easily identified among their host counterparts, allowing for enhanced visualization of axonal and dendritic processes. These optical properties also allow us to perform classic electrophysiological assays to test the synaptic potential of differentiated cells and access the overall multi-potentiality of each subline.

Lentiviral infection has proven a reliable way to stably express genes of interest (~8 kb) into slowly dividing hNPC with little to no long-term effects on behavior or morphological phenotype (Consiglio et al., 2004; Capowski et al., 2007; also see *UNIT 2D.2*). We have used the following protocols to generate hNPC engineered to constitutively express second and third-generation lentiviruses carrying either a cytosolic CAG-eGFP (kind gift of Mark H. Tuszynski) or PGK-mCherry (kind gift of Mark Mercola) fluorescent marker protein for greater than 5 months with 75% mCherry and 85% eGFP (aggregate) and 89% mCherry and 98% eGFP (MAN) efficiency for aggregate and MAN cultures respectively.

All procedures involving live infection-competent lentivirus are performed in a Level 2 or better biosafety hood in accordance with your institution's specific safety standards. We recommend full disposable safety coat, sleeves, glasses, and double nitrile and latex gloves for adequate personal protection. Any materials (pipets, tips, flasks, conical tubes) that come into

contact with virus should be properly sanitized by soaking for at least 20 min in 10% to 20% (v/v) bleach, followed by 15 min in 70% ethanol, and properly disposed of according to safety regulations. Medium is prepared fresh prior to infection. Conditioned medium can be utilized, but metabolic components may influence the overall efficiency in cell lines (D.R. Wakeman, unpub. observ.) and subsequent gene expression (McCarthy et al., 1995).

Aggregate cultures can be infected as either single cells at high density or as small size clusters (8 to 16 cells/cluster) to medium-size clusters (16 to 32 cells/cluster) (Fig. 2-6). Utilizing cellular aggregates has the added benefit of essential cell-cell contacts, whereas single cells must be plated at high density to induce cell-cell contacts quickly following infection. In addition, larger aggregates contain internal cells that may not be exposed to the viral particles, resulting in decreased infection efficiency. Although most of these cells are migratory within each cluster and eventually become labeled with increased incubation times, we recommend infecting hNPC as either small cellular aggregates or high-density single cells ( $3-4 \times 10^4$  cells/cm<sup>2</sup>), so that all cells are accessible to viral particles for optimal infection efficiency.

### ***Materials***

Human NPC, single cells or lightly adherent aggregates (Basic Protocol 1 or Alternate Protocol 1)

NB-B-27 complete medium (see recipe)

Normocin (InvivoGEN, cat. no. ant-nr-1)

Leukemia inhibitor factor (LIF; Millipore, cat. no. LIF1010)

Basic fibroblast growth factor (bFGF; Millipore, cat. no. GF003)

10 mg/ml polybrene (Chemicon)

Lentivirus (see protocol introduction for information)

Dulbecco's PBS with Ca<sup>2+</sup> and Mg<sup>2+</sup> (DPBS; Mediatech, cat. no. 21-030-CM)

25-cm<sup>2</sup> tissue culture flasks

15-ml conical tubes

1. Dissociate hNPC as described in Basic Protocol 1 or Alternate Protocol 1, as applicable, immediately prior to infection, or utilize small clusters (dissociated 48 to 72 hr before infection, similar to cryopreservation or transplantation).

2. Replate hNPC in 6 ml fresh NB-B-27 complete medium containing 2 µl/ml Normocin, 10 ng/ml LIF, 20 ng/ml bFGF, and 6 µl of 10 mg/ml polybrene in a 25-cm<sup>2</sup> flask.

*Polybrene enhances the infection efficiency but can be omitted if desired.*

3. Carefully add lentivirus at a concentration of 100 ng p24 particles for every 1 × 10<sup>6</sup> cells.

*CAUTION: Properly sanitize any virus-exposed waste with bleach and ethanol.*

4. Incubate in a 37°C, 5% CO<sub>2</sub> humidified incubator for 12 to 48 hr.

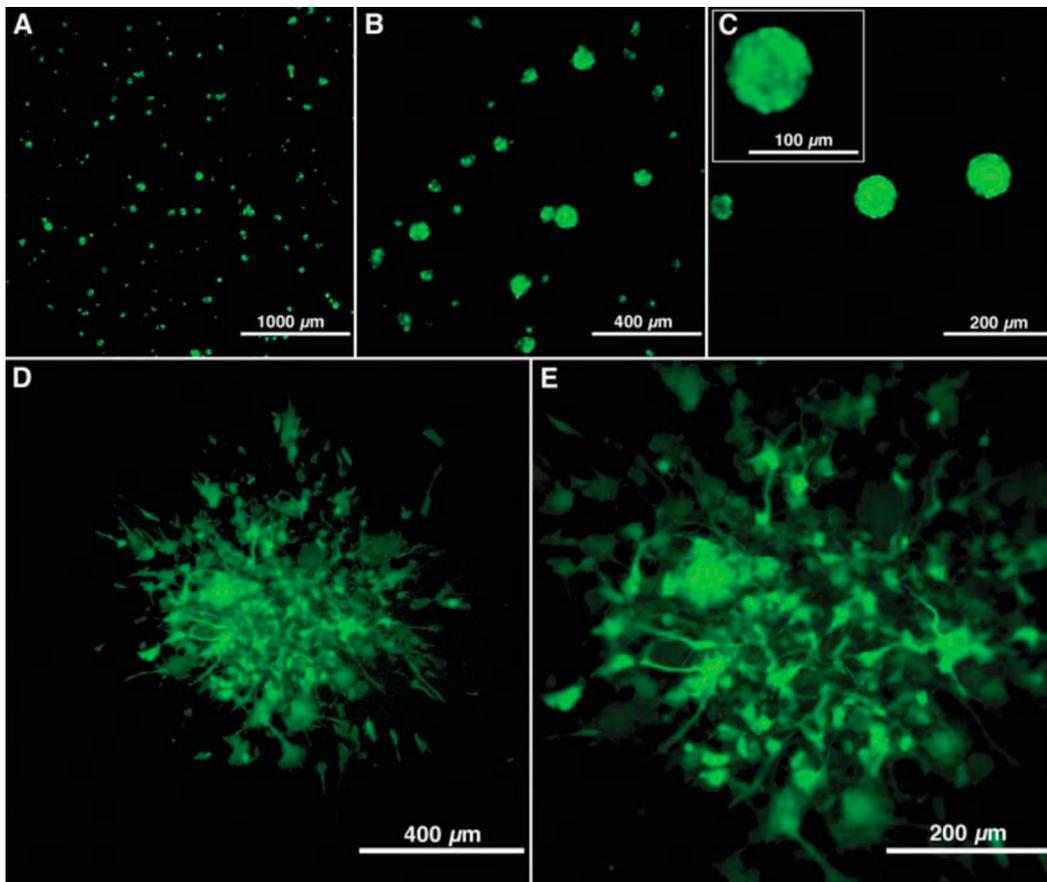
*Longer incubation times will result in higher efficiency rates and expression levels, but at the cost of multiple integration sites.*

5. To remove any remaining live virus, transfer hNPC to a 15-ml conical tube, rinse the flask once with 10 ml DPBS, combine with the cells in the tube, and allow the cells to settle by gravity. Aspirate the supernatant and repeat 10-ml DPBS rinse three times.

*Alternatively, centrifuge contents for 4 min at 400 × g in a lentivirus-approved containment centrifuge.*

6. After final rinse, add 8 ml fresh NB-B-27 complete medium containing 2  $\mu$ l/ml Normocin, 10 ng/ml LIF, and 20 ng/ml bFGF. Incubate in a at 37°C, 5% CO<sub>2</sub> humidified incubator.
7. Repeat DPBS rinse every 24 hr for the first 7 days to remove any residual viral particles.

*Excessive trituration and centrifugation can be detrimental to hNPC survival during this crucial time period. We recommend replating single cells as high-density aggregate or MAN cultures to enhance paracrine signaling required for enhanced expansion. Extensive expression of both transgenes is seen after 48 to 72 hr at an efficiency of 82% to 85% for eGFP and 75% to 79% for mCherry single cells and small clusters, respectively. Larger aggregates are much more variable and range between 45% to 85% in labeling efficiency.*



**Figure 2-6. Aggregate hNSC cultures express lentiviral eGFP.** Human NSC (HFB-2050) were dissociated into single cells at medium density ( $1^{\circ}$ — $10^6$  cells/ $25\text{ cm}^2$ ) and cultured as lightly adherent aggregates (with trituration every 12 hr) for 48 hr before transfection with a CAG-eGFP lentivirus. 72 hr after exposure (5 days in vitro), free-floating spherical aggregates readily expressed the transgene (A-C). Inset in (C), a close-up of the aggregate in the middle of the frame. When plated onto poly-D-lysine-coated tissue culture-treated flasks, hNPC-eGFP aggregates rapidly attached and flattened, confirming 80% to 90% efficiency, and sustained eGFP expression in vitro (D,E). Green = live eGFP expression.

### 2.5.2.2 Lentiviral Infection of Multilayer Adherent Network (MAN)

The MAN assay allows for temporary modification and optimization during lentivirus infections, due to the ease of changing medium; therefore, we reduce the volumes of culture medium to 5 ml fresh hNPC medium in each  $25\text{-cm}^2$  flask to concentrate virus and decrease waste. We have applied the following procedure to new and mature (60% to 80% confluent) MAN hNPC cultures ( $3\text{--}5^{\circ}$ — $10^6$  cells/flask), infecting with lentivirus containing either a

PKC or CAG promoter-driven cytosolic eGFP (Matz et al., 2002; Tao et al., 2007) or mCherry (Merzlyak et al., 2007) fluorescent protein (Figures. 2-7 and 2-8).

### ***Additional Materials***

60% to 80% confluent (2 to 3 weeks in vitro) MAN hNPC culture in 25-cm<sup>2</sup> flask

(Basic Protocol 1) or in transmembrane basket

6-well tissue culture plates

Hanging Basket Cell Culture Insert, 1.0  $\mu$ m (Millipore, cat. no. PIRP30R48)

25-cm<sup>2</sup> tissue culture flasks

1. Replace 100% of the medium in 60% to 80% confluent MAN hNPC culture (in 25-cm<sup>2</sup> flask) with 5ml fresh NB-B-27 complete medium containing 2  $\mu$ l/ml Normocin, 10 ng/ml LIF, 20 ng/ml bFGF, and 5  $\mu$ l of 10 mg/ml polybrene.

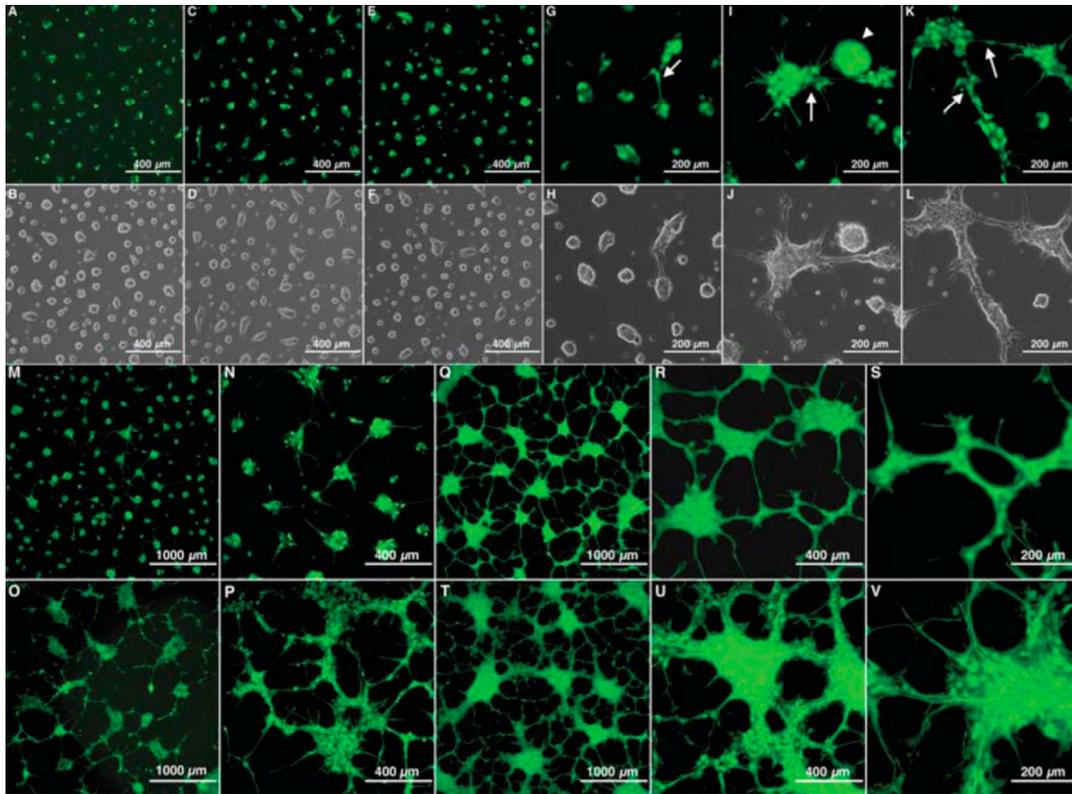
*Polybrene enhances the infection efficiency but can be omitted if desired.*

2. Carefully add the lentivirus at a concentration of 100 ng p24 particles/1  $\times$  10<sup>6</sup> cells.

*CAUTION: Properly sanitize any virus-exposed waste with bleach and ethanol. For a 60% to 80% confluent culture, there will be 3–4  $\times$  10<sup>6</sup> cells per 25-cm<sup>2</sup> flask, or 1–2  $\times$  10<sup>6</sup> cells/hanging basket cell culture insert.*

3. Incubate 12 to 48 hr in a humidified 37°C, 5% CO<sub>2</sub> incubator.

*Longer incubation times will result in higher efficiency rates and expression levels but at the cost of multiple integration sites.*



**Figure 2-7. MAN cultures efficiently express lentiviral eGFP.** Human neural stem cells (HFB-2050) were dissociated into single cells and exposed to a CAG-eGFP lentivirus (as described in text) upon replating ( $2.5 \times 10^6$  cells/25-cm<sup>2</sup> flask). hNSC were allowed to attach without agitation to induce formation of multilayer adherent networks (MAN). (A,B) 48 hr post-exposure, hNSC formed small, evenly spaced, adherent aggregate clusters and express the eGFP transgene. (C-H). By 72 hr, transgene expression markedly increased to nearly 90% of hNSC constitutively expressing eGFP throughout the cytoplasm. In addition, adherent clusters continued to proliferate and spread, making initial contacts with neighboring colonies (arrow). (I-L) Over the next week, adherent hNSC aggregates rapidly proliferated and actively migrated between adherent three-dimensional aggregates, creating the initial foundation for the multilayer adherent network. (I-J) Many peripheral anchor cells displayed long protruding feet resembling growth cones that sampled the local microenvironment and rapidly reorganized in response to local guidance cues (arrow). In addition, some clusters detached and continued to proliferate as floating suspension aggregates (arrowhead). (K-N) After 10 days, multilayer adherent clusters began to coalesce, established the classic honeycomb architecture, and actively exchanged migratory proliferative cells between colonies. Individual colonies became unrecognizable as the meandering protrusions of neighboring clusters (arrows) met and rapidly joined into one two-way hNSC highway. (O, P) At 15 days, MAN cultures continued to proliferate and expand across the culturing surface, covering nearly 40% to 60% of the flask by 20 days (Q-S). After 30 days, hNSC (HFB-2050- eGFP) assumed mature form (T-V), covering nearly 70% of the culturing surface and were ready to be dissociated. MAN cultures can be maintained for up to 6 weeks; however, overcrowding generally leads to a reduced proliferation rate, and should be avoided if possible. Green = live eGFP expression.

4. Remove any remaining virus by rinsing five times with 5 ml DPBS.

*One of the greatest benefits of utilizing the MAN assay is the ease of changing medium and rinsing live virus from the culture without extensive manipulation or harsh mechanical stress.*

*Using PBS without salts often leads to detachment of cells from the flask; therefore MAN cultures are always rinsed with DPBS containing both  $Mg^{2+}$  and  $Ca^{2+}$ .*

5. Add 8 ml fresh NB-B-27 complete medium containing 2  $\mu$ l/ml Normocin, 10 ng/ml LIF, and 20 ng/ml bFGF. Incubate in a humidified 37°C, 5% CO<sub>2</sub> incubator. 6. Repeat steps 4 and 5 every 12 hr for the first 7 days to remove any residual viral particles.

*Theoretically, all live virus will be removed with the first few rinses; however, we err on the side of safety to ensure adequate removal of all viral particles over time. In fact, we treat lentiviral-infected cultures as if they contain live virus until the cultures have been fully dissociated and passaged at least three times.*

7. After 1 to 2 weeks in culture, dissociate (1:3) and expand the newly labeled hNPC population.

*Extensive expression of both transgenes was seen after 48 to 72 hr at an efficiency of 98% eGFP and 89% mCherry. After three to five additional population doublings, fluorescently labeled hNPC populations are pooled together and sorted by FACS into three polyclonal populations based on relative fluorescence intensity, labeled low-, medium-, and high intensity expression. These cells are then expanded in 25-cm<sup>2</sup> or 75-cm<sup>2</sup> tissue-culture treated flasks (Falcon) as MAN or lightly adherent aggregate cultures and frozen for future expansion.*

*Polyclonal subpopulations can then be individually subcloned and transplanted intraventricularly into neonatal (P0) mice to assay both hNPC migration and differentiation capacity.*

8. As an example, perform MMS lentiviral infection of hNSC (HFB-2050) as follows.

*The human NPC line HFB-2050 was engineered to constitutively express either a cytosolic eGFP or mCherry fluorescent marker protein by the following methods. After 20 days of culture, hNPC MMS-multilayer adherent networks were infected with lentivirus containing either a cytosolic eGFP or mCherry protein driven by the CAG or PKC promoters respectively (Fig. 2-8).*

a. Because the MMS allows for temporary modification and optimization during lentivirus infections due to the basket's inherent mobility, reduce the volume of culture medium to 1.5 ml fresh NB-B-27 complete medium containing 2  $\mu$ l/ml Normocin, 10 ng/ml LIF, 20 ng/ml bFGF in each well of a 6-well dish, and insert a transmembrane basket with attached hNSC into each well.

*For a 60% to 80% confluent culture, there will be  $1-2 \times 10^6$  cells per hanging basket cell*

b. To each basket insert, add 10 mg/ml polybrene and lentivirus at a concentration of 100 ng p24 particles/ $1 \times 10^6$  cells. Incubate in humidified 37°C, 5% CO<sub>2</sub> incubator.

CAUTION: *Properly sanitize any virus-exposed waste with bleach and ethanol.*

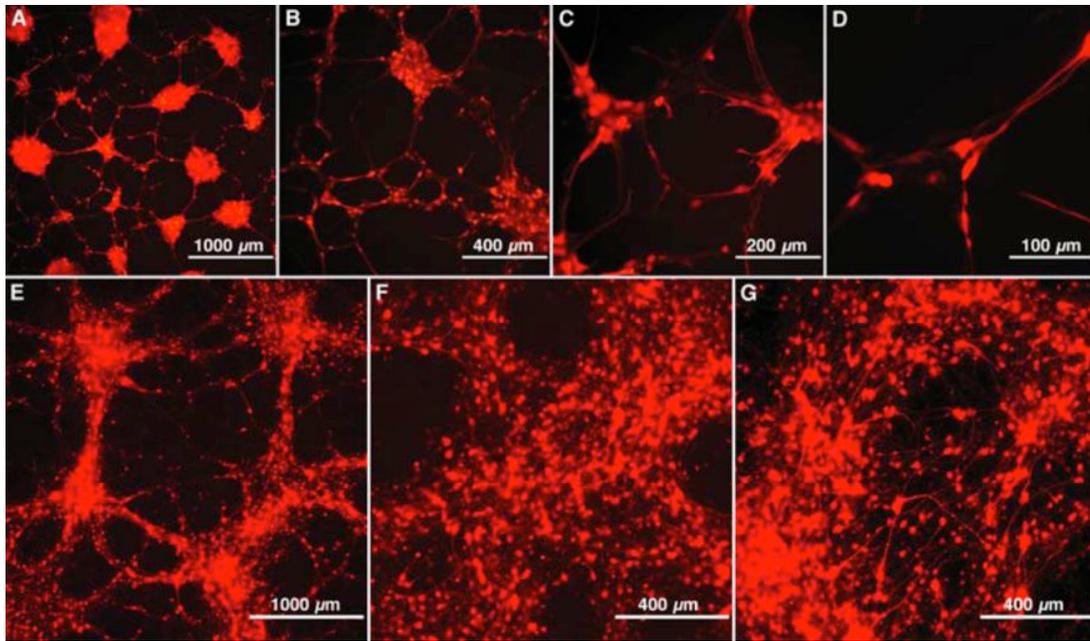
c. At a time point 48 hr later, aspirate viral waste and wash hNPC five times for 30 sec each in 5 ml DPBS. Transfer the basket insert to a new 6-well plate and return to standard culture medium volumes.

*This procedure is repeated every day for 7 days to remove any residual viral particles.*

*Extensive expression of both transgenes was seen in HFB-2050 after 72 hr at an efficiency of 98% eGFP and 89% mCherry (quantified by FACS).*

d. After 7 days, dissociate hNPC networks into single-cell suspension by adding 1.5 ml per well of Accutase and seed one well into one 25-cm<sup>2</sup> flask with 8 ml NB-B-27 medium containing 2  $\mu$ l/ml Normocin, 10 ng/ml LIF, and 20 ng/ml bFGF per flask, to induce MAN or lightly adherent aggregate cultures.

*In the case of HFB-2050, each well was dissociated into a 25-cm<sup>2</sup> flask, cultured, and expanded as a webbed MAN for 20 days before passaging again into one 75-cm<sup>2</sup>. After three total passages (90 days), all hNSCs were pooled together and sorted by FACS into three polyclonal populations based on relative fluorescence intensity, labeled low-, medium-, and high-intensity expression. These cells were then expanded in 25-cm<sup>2</sup> or 75-cm<sup>2</sup> tissue culture-treated flasks (Falcon) in modified adherent networks or as suspension aggregates, frozen for future expansion, or cloned and transplanted into (P0) neonatal mice intraventricularly (Basic Protocol 8) to assay both migration and differentiation capacity.*



**Figure 2-8. MAN and MMS mCherry lentivirus expression.** Human NSC (HFB-2050) grown as multilayer adherent aggregate (MAN) cultures maintain expression of the PGK-mCherry transgene 10 days post transfection (A-D). In addition, the MAN culturing technique can be adapted to a semi-porous membrane culturing surface and grown using the MMS technique, highly resembling MAN cultures in phenotype and composition. MMS cultures are then easily transfected with viral vectors (as described in the text) to produce fluorescent reporter cultures that are easy to manipulate (E-G).

### 2.5.3 Labeling hNPC with Super-Paramagnetic Iron Oxide (SPIO)

To follow survival and migration of transplanted donor hNPC in vivo, it has become increasingly important to develop noninvasive techniques (Manganas et al., 2007; Gilad et al., 2008; Ruiz-Cabello et al., 2008) for monitoring and imaging engrafted donor cells (Rogers et al., 2006; Slotkin et al., 2007; Schroeder, 2008; Sumner et al., 2007). Common post-mortem immunohistological staining techniques do not afford the opportunity to trace migration within the same animal over time; however, hNPC transfected with super-paramagnetic iron oxide (SPIO) particles have been used with MRI in rodents to successfully track migration of stem cells after engraftment in the normal and diseased animal (Lewin et al., 2000; Bulte et al., 2002; Frank et al., 2003; Hinds et al., 2003; Arbab et al., 2003a,b, 2004a,b; Jendelova et

al., 2004; Miyoshi et al., 2005; Magnitsky et al., 2005; Sykova and Jendelova, 2005, 2006, 2007a,b; Lepore et al., 2006a,b; Shapiro et al., 2006; Guzman et al., 2007, 2008; Politi et al., 2007; Walczak and Bulte, 2007; Neri et al., 2008; Walczak et al., 2008). Successfully labeled cells can be detected and followed weeks to months after implantation as they migrate contralaterally through predominantly white matter to sites of pathological insult. In addition, MRI findings can then be verified post-mortem utilizing immunohistochemistry by costaining for human specific markers and iron particles (Prussian blue or dextran staining) utilizing the MRI guided coordinates.

### ***Additional Materials***

Human NPC

Feridex (Bayer Healthcare Pharmaceuticals, cat. no. NDC-59338-7035-5)

Protamine sulfate injection, USP, 50 mg/5 ml (Bayer Healthcare Pharmaceuticals, cat no. NDC-63323-229-05)

25-cm<sup>2</sup> tissue culture flasks

15-ml conical tubes

Centrifuge

1. Dissociate hNPC into single cells as applicable. Replate 2–3 × 10<sup>6</sup> cells in a 25-cm<sup>2</sup> flask containing 5 ml of 25% CM/75% fresh NB-B-27 complete medium /10 ng/ml LIF/20 ng/ml bFGF/2 μl/ml Normocin. Incubate for 6 hr at in a 37°C, 5% CO<sub>2</sub> humidified incubator.

*Alternatively, skip the incubation and proceed directly to step 2. We prefer to allow freshly dissociated hNPC to recover for 6 to 12 hr after dissociation to increase viability before SPIO incubation.*

2. Prepare 2° — SPIO mixture, 45 to 60 min before labeling, by adding 10 to 20 µg/ml Feridex and 5 µg/ml protamine sulfate to 5 ml fresh NB-B-27 growth medium containing 10 ng/ml LIF and 20 ng/ml bFGF. Incubate in a 37°C water bath for 45 to 60 min with mixing.

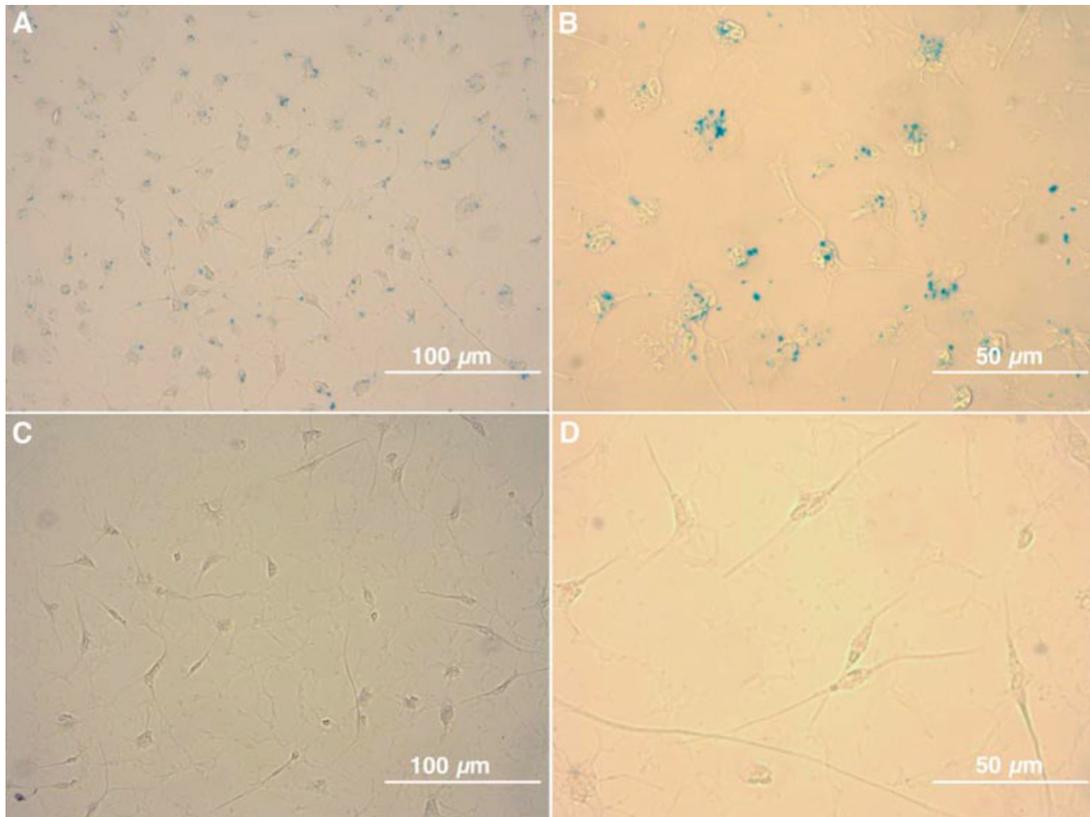
*We have had the best short-term in vitro results utilizing a final working concentration of 10 µg/ml Feridex.*

3. Add 5 ml of 2° — SPIO solution to the medium over the plated cells from step 1 and triturate gently for a final working concentration of 5 to 10 µg/ml Feridex/2.5 µg/ml protamine sulfate. Incubate for 20 to 24 hr in a 37°C, 5% CO<sub>2</sub> humidified incubator to allow iron particles to enter the cell by endosomal transport.

*Increasing the Feridex concentration will enhance the overall signal but can result in intracellular clumping and aggregation of iron particles that may be detrimental to the long-term viability, stability, and behavioral phenotype of the cell.*

4. After 20 to 24 hr, gently triturate hNPC and transfer the contents to a 15-ml conical tube. Rinse the flask once with 4 ml DPBS and add to the conical tube. Centrifuge for 3 to 4 min at 400 ° — g, room temperature.

*Increasing the incubation time will enhance the number of iron particles that enter the cell by only a small percentage and possibly at the cost of cellular differentiation or decreased post-incubation survival. In addition, the total number of particles per cell will vary within the individual culture. 24 hr is sufficient for nearly 95% labeling efficiency (Figure 2-9).*



**Figure 2-9. hNPC readily engulf SPIO particles in vitro.** Human NSC (HFB-2050) were plated onto Matrigel with SPIO particles for 20 hr and processed with Perl's Prussian blue to detect labeled cells. Most hNSC readily took in the Feridex particles (blue particles) (**A,B**), whereas unlabeled control cells (**C,D**) were completely devoid of blue staining.

5. Vacuum aspirate the supernatant and rinse by resuspending in 8ml DPBS. Centrifuge for 3 to 4 min at 400  $\times$  g, room temperature. Repeat DPBS rinse twice.

6. Resuspend the SPIO-labeled hNPC pellet in 8 ml fresh prewarmed NB-B-27 complete medium/10 ng/ml LIF/20 ng/ml bFGF/2  $\mu$ l/ml Normocin. Replate in a 25-cm<sup>2</sup> tissue culture flask and incubate in a 37°C, 5% CO<sub>2</sub> humidified incubator for 6 to 12 hr, or proceed immediately to transplantation procedures.

*Alternatively, replate SPIO-labeled hNPC onto Matrigel-coated 24-well plates for in vitro characterization and differentiation assays. Recommended plating densities are 50,000*

*cells/well for a 24-well plate, 100,000 cells/well for a 12-well plate, or 250,000 cells/well for a 6-well plate. The hNPC may also be labeled 24 hr after plating onto Matrigel by simply adding the SPIO mixture to the newly established adherent monolayer culture.*

## **2.6 PREPARING hNPC FOR TRANSPLANTATION**

Human NPC injections are performed with cells in a uniformly undifferentiated state, displaying log-phase growth, and able to incorporate BrdU during DNA synthesis. To enhance synchronization, undifferentiated hNPC are dissociated 2 days prior to transplantation and disaggregated into a single-cell suspension immediately prior to surgery. Maintaining hNPC in vitro for a longer period of time prior to injection promotes clumping due to spontaneous elaboration of extracellular matrix (ECM) or subsequent premature spontaneous cellular differentiation, both of which may deter efficient engraftment. Careful dissociation of hNPC is essential for sustained viability and long term post-transplantation survival.

Many global neurodegenerative disorders affect multiple CNS cell types; therefore, undifferentiated hNPC provide the plasticity needed for the host microenvironment to naturally direct differentiation into multiple phenotypes that may be necessary to ameliorate and restore the host cytoarchitectural milieu. On the other hand, focal CNS disorders may benefit more from lineage-directed pre-differentiation strategies, as a priori ex vivo manipulation of hNPC along specific neuronal fate pathways may be more suitable for alleviating lineage-specific CNS deficits. These cells are presumably lineage-restricted and committed, effectively eliminating nonspecific, regionally inappropriate differentiation that could lead to potentially detrimental off-target side effects. In contrast to their undifferentiated hNPC counterparts, pre-differentiated precursors and mature post-mitotic neurons do not have the same capacity for long-distance migration or the multipotency beneficial to global neurodegenerative diseases (Le Belle et al., 2004). The decision to implant a semi-

homogenous population of undifferentiated hNPC or lineage-defined progenitors is ultimately determined by the specific cellular properties that best suit the experimental paradigm (Svendsen et al., 1997a; Armstrong et al., 2000; Teng et al., 2002; Pluchino et al., 2003; Burnstein et al., 2004; Kelly et al., 2004; McBride et al., 2004; Yasuhara et al., 2006; Lee, H.J. et al., 2007; Lee, J.P. et al., 2007; Tarasenko et al., 2007). Both strategies have their merits, allowing the natural host tissue to effectively direct maturation and terminal differentiation in accordance with local signaling cues (Fricker et al., 1999).

### ***Materials***

Human NPC (hNPC): small to medium aggregate clusters (Basic Protocol 1 or Alternate Protocol 1)

Dulbecco's PBS with Ca<sup>2+</sup> and Mg<sup>2+</sup> (DPBS; Mediatech, cat. no. 21-030-CM)

0.4% (v/v) trypan blue

Hemocytometer (*UNIT 1C.3*)

Additional reagents and equipment for counting viable cells using a hemacytometer and trypan blue (*UNIT 1C.3*)

1. Triturate hNPC lightly against the bottom of the flask to detach any cellular aggregates that may have attached, and transfer the contents to a 15-ml conical tube. Centrifuge for 3 to 4 min at 400 × g, room temperature.

*Utilize small- to medium-size aggregates plated 48 hr before transplantation.*

2. Dissociate hNPC by the methods previously described (Basic Protocol 1 or Alternate Protocol 1, as applicable) and resuspend in 10 ml cold DPBS.

*Dissociation should not take longer than 1 to 3 min to fully break apart small aggregates.*

3. Centrifuge the 15-ml conical tube for 4 min at  $400 \times g$ , room temperature. Remove supernatant. Repeat steps 2 to 3.

4. With a pipettor and 100- $\mu$ l pipet tip, carefully resuspend the hNPC pellet in a small volume of cold DPBS (20 to 40  $\mu$ l), approximately equal to the volume of the cell pellet, and transfer into a 1.5-ml microcentrifuge tube.

*Resuspend hNPC in half the volume of DPBS first, then wash the 15-ml tube with the second half volume and transfer all cells into the 1.5-ml microcentrifuge tube. Be very careful not to triturate aggressively, as hNPC will easily shear when resuspended at such high density.*

5. Remove 2  $\mu$ l of cell solution and transfer to a microcentrifuge tube containing 17  $\mu$ l DPBS, add 1  $\mu$ l of 0.4% trypan blue to the tube, and triturate well.

6. Count the cells with a hemacytometer, making sure to factor in the 1:10 dilution (*UNIT 1C.3*).

*The purpose of this dilution is to use the fewest cells possible for counting, while maintaining accuracy.*

7. Adjust the amount of DPBS in the microcentrifuge tube as needed to achieve the desired cell concentration.

*We aim for ~50,000 cells/ $\mu$ l as an optimal concentration for transplantation. Cells should not be suspended at higher than 100,000 cells/ $\mu$ l for long periods of time, as excessive cell death will likely occur, resulting in a sticky DNA precipitate that will easily clog the needle.*

8. Immediately following resuspension in DPBS, place the vial of hNPC onto wet ice (4°C) and gently flick the tube every minute to deter clumping of cells that may clog the needle during stereotactic injection. Proceed without delay to Basic Protocol 7.

*Cells should be utilized as quickly as possible to reduce cell death and increase overall viability and engraftment success. For multiple animal studies, it is beneficial to prepare separate biological replicate batches as the procedures continue, preparing enough cells for use no longer than 1 hr post-dissociation.*

## **2.7 LOADING AND INJECTION OF hNPC FOR TRANSPLANTATION INTO ST. KITTS AFRICAN GREEN MONKEY**

The following methods have been refined over a number of years for stereotaxic injection of cells and tissue into the nigrostriatal system of monkeys. There are some variations in the methods which other investigators have used, but this procedure has evolved to work successfully in our hands (Redmond et al., 1986, 1988; Taylor et al., 1995; Sladek et al., 1995, 2008; Bjugstad et al., 2005, 2008; Wakeman et al., 2006; Redmond et al., 2007, 2008).

*NOTE:* All protocols involving live animals must be reviewed and approved by an Institutional Committee for Ethical Animal Care and Use (IACUC) and must conform to government regulations for the care and use of laboratory animals.

### ***Materials***

Experimental animals: St. Kitts African Green Monkey (MPTP-treated or PBS sham control)

Ketamine hydrochloride injection, USP (Ketaset, Fort Dodge Animal Health)

Atropine for i.m. or s.c. administration

Pentobarbital for i.v. administration

Lubrivet (optional; Butler Animal Health; <http://www.accessbutler.com/>)

Dulbecco's PBS with Ca<sup>2+</sup> and Mg<sup>2+</sup> (DPBS; Mediatech, cat. no. 21-030-CM)

70% ethanol

Human NPC (hNPC; Basic Protocol 6), labeled according to any of the pretransplantation labeling procedures described in this unit

NB-B-27 complete medium (see recipe) without Normocin

Phosphate-buffered saline (PBS)

100- $\mu$ l syringe (Hamilton) for stereotactic injection

22-G, 2-in. needle (Hamilton) for stereotactic injection

Single syringe microinjectors for Hamilton syringes (Stoelting)

Primate or large animal size stereotaxic head holder and bars (David Kopf Instruments)

Ear bars (David Kopf Instruments)

Animal clippers

Endotracheal tube

Sterile surgical supplies including:

Stereotactic bone drill with 0.5-mm drill burrs

Drapes

Forceps

Hemostats

Needle holders

Periosteal elevators

Scalpel with no. 10 surgical blade (Bard-Parker)

Self-retaining retractor

Towel clips

Standard neurosurgical supplies—sutures, bone wax, Gelfoam, sterile NaCl irrigation solution

3-in. sterile gauze

EKG machine with monitoring leads

Leads for temperature and O<sub>2</sub> saturation

Additional reagents and equipment for counting viable cells with a hemocytometer and trypan blue (*UNIT 1C.3*)

1. Mount the syringe into the stereotactic apparatus, calibrate according to ear bar zero, and calculate target sites from atlas or prior studies.

*For injecting monkeys, we steam-autoclave all of the equipment in the surgical field, including the stereotaxic frame with the carriers. The syringe pumps, syringes, and controllers are sterilized by ethylene oxide gas.*

2. Anesthetize monkeys initially with 7 to 15 mg/kg ketamine by i.m. injection and administer 0.02 to 0.05 mg/kg atropine by i.m. or s.c. injection. Shave and prepare the surgical area. Place an intravenous line into the saphenous or other peripheral vein and induce the animal to light anesthesia with 20 to 30 mg/kg pentobarbital for placement of an endotracheal tube to ensure adequate airway and for supplemental ventilation with room air or O<sub>2</sub>, if necessary.

3. Place the anesthetized and monitored monkey into the stereotaxic frame with the ear bars (it helps to lubricate and clean the ear canals with Lubrivet) and check the centering. Repeat if necessary.

*For accuracy of targeting, this is the most critical step of the procedure, if it is performed without MRI targeting.*

4. Place EKG monitoring leads and a lead for temperature and O<sub>2</sub> saturation.

5. Scrub and prep the already shaved head from the eyebrows to the back of the neck and from ear to ear. Remove the drape which protects the sterile cover over the ear bar on the stereotaxic frame, and then drape the head with sterile drapes and an incise drape so that nothing remains exposed except the sterile ear bar, which remains accessible for the syringe holder/carrier.

6. Recheck the zero coordinates for the drill and for the syringe carriers against a calibrated zero bar.

7. Compare the midline of the skull with the calculated coordinates for the drill. Adjust if necessary and then recalculate all targets for the drill and the carriers if the midline was adjusted. After the coordinates have been verified, slowly drill holes, lubricating with sterile DPBS. Replace the drill holder with the needle carrier and repeat steps 6 to 7.

8. Clear the Hamilton syringe by drawing and expelling 60  $\mu$ l sterile DPBS two to three times. *The DPBS rinse removes residual ethylene oxide, helps lubricate the needle, and ensures proper working condition of the syringe*

9. To fill the needle, wipe the hNPC suspension tube (optimally containing  $\sim$ 50,000 cells/ $\mu$ l) thoroughly with 70% ethanol and carefully uncap the tube inside the surgery suite. *Gently flick or triturate the hNPC microcentrifuge tube immediately prior to filling the needle.*

10. Grasp the microcentrifuge tube with hemostats and carefully move the tube up to the needle from underneath.

*The bevel of the needle should be located at the middle of the cell suspension.*

11. Slowly move the tube vertically to mix cells and begin drawing 10 to 20  $\mu$ l hNPC into the needle with the controlled-rate syringe pump.

*Cells should be drawn into the syringe immediately before the injection. Do not touch the sides or bottom of the tube, introduce air bubbles, or bend the needle. Prolonged time within the syringe will result in clumping and subsequent clogging of the needle. Always draw 2 to 3  $\mu$ l excess cell suspension into the syringe for post-injection analysis.*

12. Immediately before insertion of needle, slowly expel 1  $\mu$ l cell suspension from needle into a clean sterile microcentrifuge tube. Add 1 ml fresh NB-B-27 complete growth medium without Normocin to the microcentrifuge tube.

13. Lower the needle into target area of interest at slow speed (10 to 15 sec) to the proper depth for the given stereotactic location.

*At the appropriate vertical depth, wait 2 min to allow the brain tissue to adjust to the proper level at the needle tip.*

14. Inject the prepared cell suspension using the syringe pump at a maximum constant rate of (1  $\mu$ l/min).

15. During autoinjection, count viable hNPC in the microcentrifuge tube with a hemocytometer and trypan blue (*UNIT IC.3*) to determine viability at time of injection and replat the 1 ml of hNPC into one well of a 24-well dish. Incubate at 37°C for 48 to 72 hr and examine under the microscope to determine if any contamination was introduced during the

procedure.

16. After the cells are fully expelled from the syringe, allow the needle to remain in place for 2 min before withdrawing.

*The extra settling time will prevent significant backflow and leakage of hNPC through the needle track during retraction of the needle.*

17. Retract needle slowly, at a maximum of 1  $\mu\text{m}/\text{min}$ . If another injection is planned, flush out the syringe with sterile PBS. Refill and repeat step 8 above.

18. After the last injection, make sure that any bleeding is controlled, suture animal, and proceed to post-operative care.

## **2.8 INTRAVENTRICULAR INJECTION OF hNPC INTO NEONATAL MICE**

Neonatal mice (post-natal day 0 to 3; P0 to P3) are relatively effortless to handle and can easily be injected in the lateral ventricles with hNPC, offering several advantages over adult mice for transplantation. The undeveloped newborn skull is soft and translucent, obviating the need for drilling or cutting into the head while facilitating penetration of the needle or drawn glass micropipet. In addition, hNPC survive, migrate, and integrate within the developing neonatal CNS at a higher efficiency than when transplanted into the mature adult brain. It should be noted that engraftment, migration, synaptic maturation, and development of fully competent neuronal subtypes often take longer with human-derived NPC than with rodent-derived NPC; therefore, we highly recommend the use of immunodeficient rodent models such as the SCID genetic background to increase overall success and eliminate the need for expensive immunosuppressants (Chidgey et al., 2008). Other strains with variations

in T, B, and NK cell deficiencies such as BALB/c and Rag2 backgrounds may also be used, but may display some leakiness with age. We prefer the NIHS-beige-nude-xid (NIHBNX) or C.B-17 scid beige (CBSCBG) mice available from Taconic or Harlan. The extra cost for maintenance of these animals is far outweighed by time and resource expenditures involved in continually testing and rederiving new hNPC lines, which could be avoided by eliminating the false negatives involved in using non-immunopriveleged strains.

### ***Materials***

Neonatal mice (P0 to P3) of appropriate strain

70% ethanol

Dulbecco's PBS with Ca<sup>2+</sup> and Mg<sup>2+</sup> (DPBS; Mediatech, cat. no. 21-030-CM)

Human NPC (hNPC; Basic Protocol 6; 50,000 cells/μl) in microcentrifuge tube

Borosilicate glass (Sutter Instrument Co., cat. no. B100-75-15)

Micropipet puller (Sutter Instrument Co., Model P-87)

Aspirator tube assemblies for calibrated microcapillary pipets (Sigma Aldrich, cat. no. A5177-5EA)

Fiber optic light source for transillumination (Dolan-Jenner Industries)

Warm-water glove balloon

Additional reagents and equipment for preparing injection micropipet (Lee et al., 2008) and processing mouse brains

1. Prepare calibrated drawn borosilicate glass micropipet using borosilicate glass and a micropipet puller (Lee et al., 2008).
2. Anesthetize the neonatal mouse by placing the pup on wet ice for ~1.5 to 3 min until the

animal no longer retains locomotion or responds to gentle toe and tail pinch.

*Carefully monitor the pup to minimize time on ice, as overexposure to low temperatures can lead to death of newly born mice. Immediately proceed to transplantation*

3. Insert a calibrated drawn borosilicate glass micropipet into an aspirator tube assembly. Just prior to drawing up hNPC, rinse the micropipet by drawing up and then expelling 5  $\mu$ l of 70% ethanol ten times, followed by sterile DPBS ten times, to sterilize and lubricate the glass.

4. Gently flick or triturate the microcentrifuge tube containing the hNPC suspension immediately prior to filling the needle. Wipe the tube thoroughly with 70% ethanol and carefully uncap the tube.

5. Slowly move the tube vertically to mix cells and begin drawing 4 to 5  $\mu$ l hNPC into the micropipet by mouth suction.

*Do not touch the sides or bottom of the tube, introduce air bubbles, or break the glass needle.*

*Prolonged time within the micropipet will result in clumping, subsequently clogging the needle. Always draw 2 to 3  $\mu$ l excess cell suspension into the syringe for post-injection analysis.*

6. Loosely secure the skull by hand at the cranium and place directly over a non heat conducting (fiber-optic) light source to visualize the bregma and lateral ventricles by transillumination.

*Gentle handling of the pup throughout the procedure is important to avoid trauma.*

7. Carefully insert glass needle ~0.5 to 1 mm deep into the head at the midline between

bregma and eye and slowly inject 1 to 2  $\mu\text{l}$  hNPC suspension at 5 ° — 104 cells/ $\mu\text{l}$  into the lateral ventricle of either the left or right hemisphere.

*Correct location and accurate dispersion within the ventricles can be confirmed by addition of an inert dye to the cell suspension. A correctly placed glass needle will deliver cells without resistance by mouth-pressure delivery from a micropipet aspirator tube assembly. We recommend practicing on nonexperimental animals with trypan blue, to become acquainted with the correct pressure and distance necessary for accurate intraventricular injection of experimental animals with hNPC cell suspensions. P0 mice generally will tolerate 3  $\mu\text{l}$  total divided between both ventricles, while the later stages P2 to P3 can tolerate 4 to 5  $\mu\text{l}$  total.*

8. Slowly remove the needle and check for leakage through the needle track. Repeat step 7 in the contralateral hemisphere.

*The entire procedure starting from grasping the pup off of ice to injection should not take longer than 40 to 60 sec per animal to avoid having the anesthetic wear off.*

9. Immediately following injection, warm the lower extremities under tepid flowing water, gently pat dry with a cotton swab or piece of gauze, and place on a warm-water glove balloon on top of a heating pad to adequately increase the body temperature of each pup before returning the pup to its mother, to avoid post-operation parental rejection.

*Monitor pups after transplantation for several hours to ensure adequate post-operational recovery. It is very important to increase core body temperature quickly, for best results. The brain is not an immunoprivileged organ; therefore immunosuppression regimes are recommended to avoid rejection in applicable genetic backgrounds (e.g., wild-type strains).*

10. Process brains for differentiation potential at the following time points (see Basic Protocol 9):

a. Injection location confirmation (2 to 12 hr).

*Location and structures of interest, lateral ventricles and luminal cavity: Trypan blue tracer can easily be seen within the lateral ventricles at this time, confirming correct needle position at time of injection. In addition, the overall integrity of the brain and amount of damage incurred by the glass needle track can be assessed to validate technical procedure for optimal hNPC transplants. Cells will begin to engraft in the first 8 to 12 hr following transplantation; however, accurate assessment for cell survival can not be attained this early.*

b. Engraftment (12 to 24 hr).

*Location and structures of interest, lateral ventricular walls, including SVZ and choroids plexus: Cells will initially incorporate within the ventricular walls and preferentially accumulate in the SVZ as bulbous nodules where they eventually incorporate into the host SVZ and rostral migratory stream (RMS). Dead cells are often cleared and become trapped within the choroid plexus, the natural “kidney” of the brain that actively filters CSF. In general, there is a give and take between the moderate amount of cell death that occurs during the transplantation process and the subsequent cell divisions that take place post-transplantation.*

c. Integration and migration within SVZ-RMS niche (1 to 3 days).

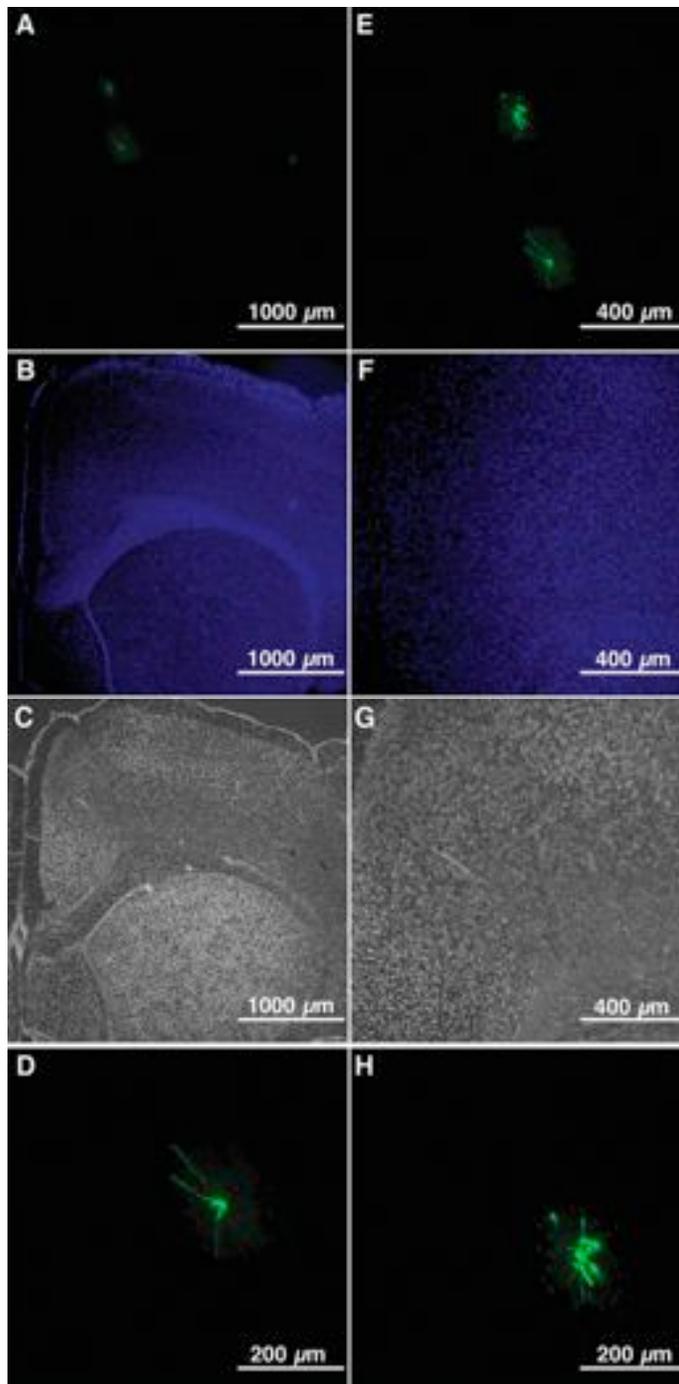
*Location and structures of interest, SVZ, RMS, olfactory bulbs (OB): Successfully engrafted hNPC incorporate within the developing ventricular walls, forming nodules reminiscent of SVZ protrusions induced by intraventricular growth factor infusion (Kuhn et al., 1997; de Chevigny et al., 2008). The bulbous cluster eventually flattens back out into a normal*

*ventricular wall, and the hNPC integrate within the host neurogenic niche. Within the endogenous SVZ niche, a small population of donor cells will continue to proliferate (Ostenfeld et al., 2000) for at most two to four cell divisions (in our experience) before exiting the cell cycle as they mature and coordinately migrate tangentially and by chain migration through the RMS. The number of cells that die during and following transplantation far outweighs the additional proliferative load; therefore, unchecked tumor-like growth does not occur.*

d. Migration, differentiation, and synaptic integration (3 to 14 days).

*Location and structures of interest, RMS and olfactory bulbs (OB), neuraxis: As hNPC preferentially migrate through the RMS to the OB, they receive signaling and guidance inputs that eventually direct them to defer from tangential or chain migration and turn radially where they continue to migrate and differentiate into synaptically integrated OB interneurons. In addition, many cells will migrate through predominantly white matter and integrate appropriately into the developing striatum and to a lesser extent contribute to the cortex as well (Fig. 2-10). Utilizing non-immunopriveleged animals that do not exhibit pathological deficit often leads to a significant decrease in overall graft survival, integration, and long-term synaptic connectivity. When testing hNPC differentiation potential, we recommend always using immunodeficient mouse models (SCID, BALB/c, or Rag2) to decrease time-consuming false negatives. Assaying donor cell survival after 2 to 3 weeks in vivo in nondiseased animals can often be misleading, as many of the cells are discarded, become quiescently integrated within the ventricular wall and SVZ, or simply undergo apoptosis in response to local microenvironment niche signaling cues. It seems that once the initial process of fetal development has concluded, only a small portion of transplanted hNPC remain quiescently undifferentiated within the neurogenic niches, while the remainder of donor cells*

*continue to migrate, differentiate, and eventually be replaced by the natural host neurogenic process. In non-immunopriveleged animals, we speculate that these donor cells undergo additional selectional pressure as the host immune system develops, and this may eventually deter the long-term survival and maturation of OB interneurons. One way or another, very few transplanted hNPC will be found after 1 to 3 months in vivo in nondiseased, non-immunopriveleged animals; therefore, careful consideration of genetic background and terminal end point is essential for transplantation success [Figure 2-10].*



**Figure 2-10. Engrafted hNSC survive, mature, and express the eGFP transgene in vivo.** Human NSC (HFB-2050) engineered to express eEGP under the CAG promoter were established and transplanted at low density into both lateral ventricles of P0 neonatal mice (as described in the text). Mature eGFP<sup>+</sup>, donor-derived neurons were found in clusters throughout the forebrain up to 10 weeks post-transplantation (**A-H**); DAPI (**B,F**), phase (**C,G**), GFP (**A,D,E,H**).

## **2.9 CHARACTERIZING hNPC**

Throughout expansion and long-term maintenance, hNPC are periodically assayed for *in vitro* expression of known “stemness” markers. Human NPC should be analyzed by standard immunoassays to demonstrate sustained undifferentiated characteristics and cellular morphology. Prior to transplantation, a subset of hNPC are set aside and assayed for the following markers (see below) in multiple combinations to confirm morphologically relevant expression profiles exist.

The fate of donor cells in the CNS can be assessed *in vivo* by dissecting the brains and processing by standard immunohistochemical methods for lineage-specific antigens such as nestin/vimentin for hNSCs; GFAP and s-100-Beta for astrocytes; and NeuN, NFM70/200, and synaptophysin for mature neurons. Antibodies to the human nuclear antigen (HuNuc) can be used to co-label hNSCs for donor confirmation. The functionality of donor cells in the brain can also be assessed by classic electrophysiological methods *ex vivo*.

### **2.9.1 Undifferentiated hNPC**

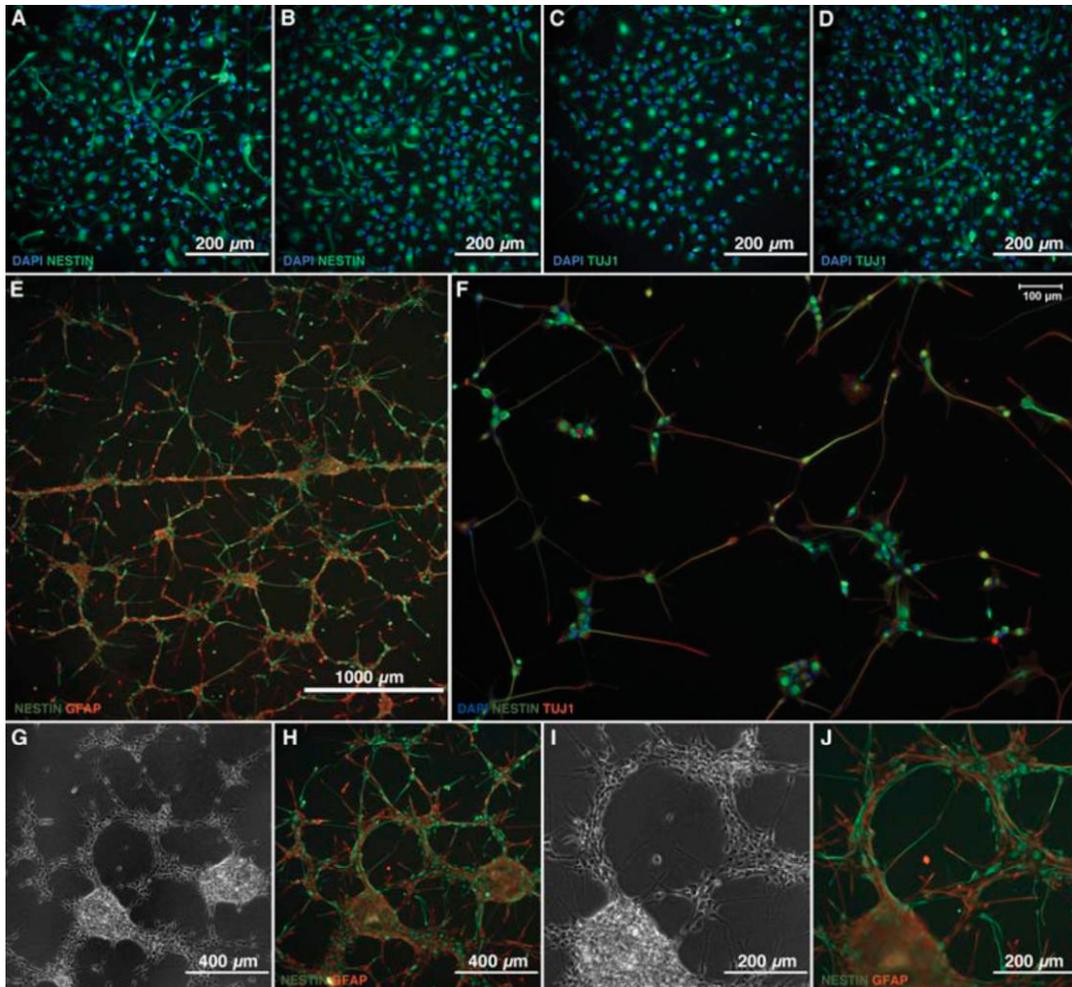
Healthy proliferative Ki-67+ (Scholzen and Gerdes, 2000; Ab available from Abcam, use at 1:200 dilution) or PCNA+ (Hall and Levison, 1990; Hall et al., 1990; Ab available from Santa Cruz Biotechnology, use at 1:100 dilution) undifferentiated hNPC express the nuclear transcription factor Sox-2 (D’Amour and Gage, 2003; Komitova and Erickson, 2004; Baer et al., 2007; Ab available from Santa Cruz Biotechnology, use at 1:100 dilution), musashi-1/2 (Chan et al., 2006; Ab available from Abcam, use at 1:100 dilution), the filamentous cytoplasmic proteins Nestin (Ab available from Chemicon, use at 1:400 dilution; Fig. 2-11A,B,E-J) and Vimentin (Ab available from Chemicon, use at 1:400 dilution), and the surface protein LexA/SSEA-1 (Ab available from Chemicon, use at 1:250 dilution) in cellularly appropriate locations (Temple, 2001; Pixley and de Vellis, 1984; Pixley et al.,

1984a,b; Hockfield and McKay, 1985; Lendahl et al., 1990; Dahlstrand et al., 1992; Zimmerman et al., 1994; Garcia et al., 2004). In addition, fetal hNPC also express glial fibrillary associated protein (GFAP; Ab available from DAKO, use at 1:400 dilution) in morphologically appropriate (nonprotoplasmic) locations, highly correlating with Nestin expression (Fig. 2-11. E,G-J), consistent with rodent data (Doetsch et al., 1999; Laywell et al., 2000; Imura et al., 2003), suggesting that GFAP also labels human fetal NPC or that astrocytes may be hNPC at specific times during early development. It is critical that these markers be utilized in combination to confirm the stem/precursor phenotype of donor cells before transplantation. In addition to the classic stem cell markers, we have also confirmed the expression of radial glia (RG) associated brain lipid binding protein (BLBP; Feng et al., 1994; Feng and Heinz, 1995; Ab available from Chemicon, use at 1:350 dilution), which has been linked to fetal NSC in vivo (Fig. 2-12; Garcia et al., 2004; Malatesta et al., 2000; Hartfuss et al., 2001; Alvarez-Buylla et al., 2001, 2002; Miyata et al., 2001; Noctor et al., 2001, 2002; Gotz et al., 2002; Gregg et al., 2002; Doetsch, 2003; Malatesta et al., 2003; Goldman, 2003; Gotz, 2003; Gregg and Weiss, 2003; Noctor et al., 2004, 2008; Merkle et al., 2004; Gotz and Bard, 2005; Merkle and Alvarez-Buylla, 2006; Merkle et al., 2007).

In addition, a small population of bipolar migratory cells express the microtubule associated protein doublecortin (DCX; Fig. 2-12E,F; Gleeson et al., 1999; Francis et al., 1999; Friocourt et al., 2003; Ab available from Santa Cruz, use at 1:400 dilution) and the membrane-bound polysialylated neural cell adhesion molecule (PSA-NCAM; Hu et al., 1996; Curtis et al., 2007; El Marouf and Rutishauser, 2008; Rutishauser, 2008; Burgess et al., 2008; Ab available from Chemicon, use at 1:250 dilution). Furthermore, some later-passage hNPC populations also express the intermediate filament protein, beta-3-tubulin (Tuj-1; Fig. 2-11 F; Caccamo et al., 1989; Geisert and Frankfurter, 1989; Lee et al., 1990; Menezes and Luskin, 1994; Menezes et al., 1995; Ab available from Covance Research, use at 1:400 dilution) and

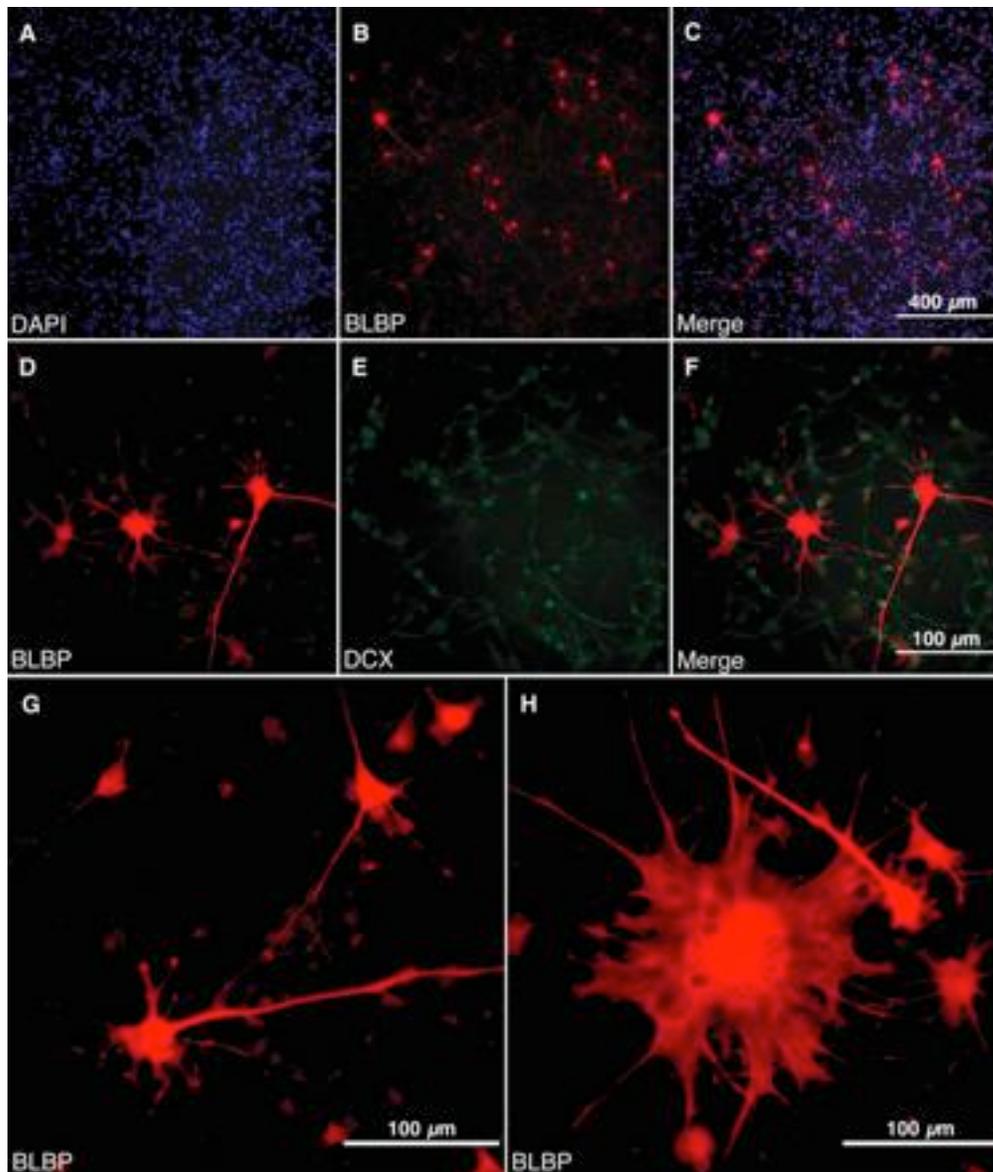
colocalizes with GFAP (Rakic, 1972; Sidman and Rakic, 1973; Levitt and Rakic, 1980; Levitt et al., 1981, 1983). Although beta-3-tubulin has been described as an early marker for immature neuron ally–restricted NPCs in rodents, these hNPC continue to self-renew and maintain multipotency in vitro and in vivo. Furthermore, they are phenotypically and behaviorally indistinguishable from early-passage predecessors. As a result, assuming the same stemness profile for human cells may not be entirely appropriate, and this should be taken into account when assessing hNPC fate. For example, expression of GFAP or beta-3-tubulin alone may not be entirely sufficient to assume loss of multipotency or regional differentiation into neurons and astrocytes, respectively. Instead, we favor a mode of characterization based on both morphology and multiple marker comparison to appropriately assess stem cell fate both before and after transplantation.

It is well known that cell lines accumulate in vitro artifacts in response to long-term artificial cell culture environments (Doetsch et al., 2002; Pollard et al., 2008). Whether the trends we see in vitro appropriately mimic the in vivo nature of hNPC remains to be determined. Certainly, it stands to reason that hNPC share many developmental markers in common with their rodent counterparts; however, it is not beyond the scope of reason to assert that perhaps some of the key players involved in human neural developmental processes may be differently regulated spatiotemporally from the corresponding processes in lower-order mammals. We have verified these findings by both standard immunocytochemistry and western blotting in three hNPC lines ranging from 10 to 22 weeks (from fertilization date), further suggesting that hNPC are not homogenous populations of identical stem cells, but rather highly dynamic heterogeneous populations (Kukekov et al., 1999; Laywell et al., 2002; Suslov et al., 2002; Steindler et al., 2003; Chen et al., 2006) of neural progenitors. Clonal analysis assays and in-depth time-lapse fluorescence video microscopy are currently being used to determine the specific lineage relationships involved.



**Figure 2-11. hNSC express classic neural stemness genes.** Human NSC (HFB-2050) express nestin (**A,B**) and Tuj-1 (**C,D**) uniformly when plated as aggregates on poly-D-lysine-coated tissue culture-treated flasks (blue = DAPI). Cells rapidly attach and begin to elongate, sending processes throughout the culture. (**A**) and (**D**) were processed 24 hr after plating, while (**B**) and (**C**) represent the more immature morphology seen after 12 hr of culture. DAPI+ nuclear morphology indicates active mitosis and sustained proliferation at both time points. In addition, nestin+ hNSC (green) also express GFAP (red) (**E,H,J**), typically in inverse proportion to each other. For example, a cell with high GFAP expression will also express nestin, but at a much lower level, whereas a highly nestin+ cell will express GFAP at a lower level. Interestingly, both cell types are intimately interwoven within each other, forming a meshwork of migratory cells (pictured here on poly- L-ornithine/fibronectin-coated tissue culture-treated flasks). Furthermore, nestin+ hNSC also express Tuj-1 (**F**) throughout most of the cytoplasm. Typically, nestin expression is highest surrounding the nuclear box, whereas Tuj-1 expression is greatest at the feet of meandering processes. Expression patterns are typically opposite of each other, and tend to colocalize in the middle of the cell's architecture, similar to nestin and GFAP coexpression. These filamentous proteins may play distinct roles at their specific positions within the stem cell, conferring or coordinating cell polarity within the *in vitro* microniche (**G,I** = phase contrast).

Whether these distinct phenotypic outliers represent bona fide stem cells or restricted progenitors that have been reprogrammed in response to mitogenic signaling cues remains to be determined. Certainly, the new wave of research dedicated to studying induced pluripotent stem cells (iPS) suggests that this process is much easier to induce than previously appreciated (Takahashi and Yamanaka, 2006; Takahashi et al., 2007 a,b; Maherali et al., 2007; Wernig et al., 2007; Okita et al., 2007; Meissner et al., 2007; Yu et al., 2007; Nakagawa et al., 2008; Park et al., 2008; Brambrink et al., 2008; Wernig et al., 2008; Stadtfeld et al., 2008; Shi et al., 2008; Kim et al., 2008; Maherali et al., 2008; Maherali and Hochedlinger, 2008a,b). Therefore, it is only through careful phenotypic characterization that we can begin to understand the nature of hNPC in vitro. Until better methods and markers are discovered to accurately assess cellular identity, we are forced to apply the borrowed phrase, “It’s hard to define, but I know it when I see it” (Morrison et al., 1997).



**Figure 2-12. hNPC exist as heterogeneous populations resembling the SVZ niche.** In addition to the classic NSC marker proteins, some cells also highly express brain lipid binding protein (BLBP) (A-H) or doublecortin (DCX) (E,F). Cells that highly express BLBP typically assume an astrocyte-like star morphology and have one or two long meandering processes with a highly arborized cell body, resembling radial glia. In contrast, most cells that do not express BLBP highly or have radial glia morphology express DCX and highly resemble the migratory transit amplifying cells found in vivo within the subventricular zone NSC niche. Images were taken from hNSC (HFB-2050) plated on Matrigel-coated tissue culture plates and cultured for 3 days in vitro to induce attachment and spreading. Of note, extracellular matrix components and growth factors found in Matrigel have a profound impact on cell morphology and may affect the differentiation profile of cultures over time. It is likely that these cultures have started an initial differentiation process and may not retain all stemness properties. Nonetheless, the heterogeneity of hNSC cultures is evident as 72 hr is probably not sufficient time to considerably differentiate hNSC.

### 2.9.2 Differentiated hNPC

In addition to the classical stemness markers, cells may be differentiated into neurons, astrocytes, or oligodendrocytes by a variety of methods and assayed for lineage-specific differentiation markers. Specific methods for in vitro differentiation are detailed elsewhere (Wakeman et al., 2009; Johe et al., 1996; Hsieh and Gage, 2004; Androutselis- Theotokis et al., 2008). Differentiated cells are fixed in 4% cold PFA, stained with the appropriate antibodies by standard protocols, and analyzed by indirect immunofluorescence for the expression of pro-neuro/gliogenic markers. To determine whether hNPC are capable of giving rise to neurons (after allowing at least 3 weeks of in vitro maturation), the cells are stained for an extensive panel of pro-neuronal markers—first (immature): doublecortin (Santa Cruz, dilute 1:500), b-III-tubulin (Chemicon, dilute 1:200), Pax6 (Covance, dilute 1:400), Ptx3 (Abcam, dilute 1:500, or R&D Systems, dilute 1:400), Lmx1a/b (Santa Cruz, dilute 1:200), Gbx1/2 (Santa Cruz, dilute 1:250), Ngn1/2/3 (Santa Cruz, dilute 1:250), and then more mature neuronal phenotypes: PSA-NCAM (Chemicon, dilute 1:100), high-molecular-weight neurofilament (Boehringer, dilute 1:150), tau (Sigma, dilute 1:400), NeuN (Chemicon, dilute 1:50), MAP-2 (Sigma, dilute 1:200), synaptophysin (Sigma, dilute 1:500), calbindin (Sigma, dilute 1:400), calretinin (Chemicon, dilute 1:500), Mash1 (BD Biosciences, dilute 1:100), Msx1 (Abcam, dilute 1:500), En1 (Iowa Developmental Hybridoma Bank, dilute 1:50), Girk2 (Alomone Laboratories, dilute 1:500), Nurr-1 (Santa Cruz, dilute 1:500), TH (PelFreeze Biologicals, dilute 1:500, or primary rabbit antiserum, Eugene Tech, dilute 1:3000), DAT (rabbit, Affinity Bioreagents, dilute 1:500), VMAT2 (PelFreeze, dilute 1:500), AADC (Chemicon, dilute 1:1000; dopaminergic), GABA (Sigma, dilute 1:5000 or rabbit polyclonal antibody for GAD65/67, Chemicon, dilute 1:1000), ChAT (Chemicon, dilute 1:400; cholinergic), 5-HT (Sigma, dilute 1:50 or rabbit polyclonal, Calbiochem, dilute 1:1200), as well as the radial glial, BLBP (Chemicon, dilute 1:1000), A2B5 (R&D Systems, dilute 1:400),

the mature astroglial, GFAP (polyclonal rabbit, Dako-Patts, dilute 1:1000, Sternberger Monoclonals, dilute 1:200, or Chemicon, dilute 1:200), S100B (Abcam, dilute 1:500), glutamate transporters Glast/GluT-1/EAAT1 (Chemicon, dilute 1:400, or Santa Cruz, dilute 1:500) and GLT1/EAAT2 (Chemicon, dilute 1:400, or Santa Cruz, dilute 1:500), and oligodendrocyte fate, MBP (Chemicon, dilute 1:200), O4/O1 (Chemicon, dilute 1:200), CNPase (Chemicon, dilute 1:200), otx2 (Santa Cruz, dilute 1:100), and RIP (Chemicon, dilute 1:400).

## 2.10 COMMENTARY

The great neuroanatomist Ram´on y Cajal wrote, “In the adult centers the nerve paths are something fixed, ended and immutable. Everything must die, nothing may be regenerated” (Ram´on y Cajal, 1928; Ram´on y Cajal and May, 1959). This observation, based on the primitive methods of the time, held up as neurodevelopmental dogma for centuries. It wasn’t until Joseph Altman and Michael Kaplan’s classic autoradiographic experiments using tritiated thymidine (Altman, 1962a,b, 1963; Altman and Chorover, 1963; Altman and Das, 1965a,b, 1966) that neurobiologists even considered rethinking the notion of adult neurogenesis (Allen, 1912; Messier et al., 1958; Messier and Leblond, 1960; Smart, 1961; Smart and LeBlond, 1961; Kaplan and Hinds, 1977), let alone embraced it as an intrinsic process active throughout adulthood until death.

In the past quarter century, it has become undeniably clear that neural stem/progenitor cells reside within the developing embryonic, neonatal, and adult songbird (Goldman and Nottebohm, 1983), rodent (Alvarez-Buylla et al., 2002; Snyder et al., 1992; Morshead et al., 1994; Weiss et al., 1996; Johanson et al., 1999), monkey (Gould et al., 1998, 1999a,b; Kornack and Rakic, 2001a,b), and human forebrain (Merkle et al., 2007; Curtis et al., 2007; Sanai et al., 2004; Howard et al., 2006; Quinones-Hinojosa et al., 2006, 2007; Sanai et al., 2007), primarily

lining the posterior to anterior subventricular zone (SVZ; Doetsch et al., 1999; Merkle et al., 2004; Menezes et al., 1995; Lois and Alvarez-Buylla, 1993; Luskin, 1993; Lois and Alvarez-Buylla, 1994; Rousselot et al., 1995; Luskin et al., 1997) of the lateral ventricular walls and within the subgranular zone of the hippocampal dentate gyrus (Ray and Gage, 2006; Ray et al., 1993; Gage et al., 1995a). These cells can be derived from various regions of the brain, with limited capacity, where they exist naturally as relatively quiescent populations of stem and progenitor cells (Palmer et al., 1995) within the complex microenvironment of a highly dynamic, tightly junctioned, neurogenic niche (Sanai et al., 2007; Alvarez-Buylla and Lim, 2004; Verdugo and Alvarez-Buylla, 2006; Lim et al., 2007).

During mammalian CNS development, hNSC undergo an initial expansion phase of symmetric divisions followed by nonsymmetric divisions and extensive migration in accordance with electrical stimulation (Deisseroth et al., 2004; Spitzer, 2006) and chemical (Ghashghaei et al., 2007) guidance cues. Differentiation is characterized by stages of neurogenesis followed by gliogenesis (Qian et al., 1997, 1998, 2000; Namihira et al., 2009), differentiating in temporal waves of first neurons, then astrocytes and oligodendrocytes, to shape and form the mature human brain (Levison and Goldman, 1993; Levison et al., 1993; Menn et al., 2006). In the adult, NPC continue to proliferate and migrate by a combination of tangential and chain migration from the SVZ through the rostral migratory stream (RMS) into the olfactory bulbs (OB; Lois et al., 1996; Doetsch and Alvarez-Buylla, 1996), generating new neurons and glia (Kuhn et al., 1996; Goldman et al., 1997; Petreanu and Alvarez-Buylla, 2002; Carleton et al., 2003; Lledo et al., 2006) as an active pool to replace or restore homeostasis to the aging or injured brain (de Chevigny et al., 2008; Lim et al., 2007; Nait-Oumesmar et al., 1999; Picard-Riera et al., 2002; Lie et al., 2004; Parent et al., 2006; Leung et al., 2007; Hellstrom et al., 2008).

In addition, cultured NSC/NPC can be directed *in vitro* and *in vivo* to give rise to all

three neuroectodermal lineages (Arsenijevic et al., 2001a; Johe et al., 1996; Gage et al., 1995b; Kirschenbaum et al., 1994; McKay, 1997; Levison and Goldman, 1997; Murray and Dubois-Dalcq, 1997; Pincus et al., 1998; Takahashi et al., 1999; Rao, 1999; Brannen and Sugaya, 2000; Dietrich et al., 2002; Riaz et al., 2004; Scheffler et al., 2005; Christophersen et al., 2006; Pistollato et al., 2007; Rao et al., 2008), as well as a variety of intermediate cellular phenotypes (Markakis et al., 2004) when presented with the appropriate signaling cues. These cells may act by a variety of mechanisms, either by providing potential new raw material for regenerating the damaged CNS or by rescuing the endogenous nervous system through secondary neuroprotection mechanisms (Pluchino et al., 2005a), thereby modulating the host microenvironment (Madhavan et al., 2005, 2006, 2008), conferring a return to baseline nonpathological stasis (Bjugstad et al., 2005). In this manner, neurological function may be restored through transplantation therapies by either directly integrating and replacing host neural circuitry, or, secondarily, by rescuing and restoring the endogenous host milieu through growth factor or neurotransmitter paracrine signaling (Brustle and McKay, 1996; Park et al., 2002a; Svendsen and Langston, 2004; Tai and Svendsen, 2004; Emsley et al., 2005; Pluchino et al., 2005a,b; Schwartz, 2006). Moreover, multipotent hNSC have been shown to display low immunogenicity (Mason et al., 1986; McLaren et al., 2001; Odeberg et al., 2005), readily express foreign transgenes (Flax et al., 1998; Ostensfeld et al., 2002a; Wu, P. et al., 2002; Park et al., 2003; Behrstock and Svendsen, 2004; Kim, 2004; Klein et al., 2005; Behrstock et al., 2006; Capowski et al., 2007; Roy et al., 2007; Suzuki et al., 2007; Ebert et al., 2008), and inherently home to sites of pathological insult (Aboody et al., 2000; Park et al., 2002b; Ourednik et al., 2002; Imitola et al., 2004; Park et al., 2006).

Traditionally, human neural stem cells (hNSC) have been operationally defined (Weissman et al., 2001; Anderson, 2001; Seaberg and van der Kooy, 2003; Parker et al., 2005; Navarro-Galve and Martinez-Serrano, 2006) by two cardinal criteria, first, the ability to *self-*

*renew indefinitely* by division into two identical (symmetric) or nonidentical (asymmetric) daughter cells, and, second, to maintain the *multi-potential* capacity to differentiate into all three neuroectodermal CNS lineages: neurons, astroglia, and oligodendrocytes in vivo (Gage et al., 1995a; Brustle et al., 1997; Lundberg et al., 1997; Winkler et al., 1998; Zigova et al., 1998; van der Kooy and Weiss, 2000; Weissman et al., 2001; Buchet et al., 2002). Asymmetric daughter cells that are not identical to the parental hNSC have a reduced stemness and are therefore referred to as intermediate or transit amplifying neural progenitor cells (Doetsch et al., 1999; Merkle and Alvarez-Buylla, 2006; Verdugo and Alvarez-Buylla, 2006; Doetsch et al., 1997; Garcia-Verdugo et al., 1998; Lim et al., 2008; for reviews see Kemperman, 2006; Slack, 2008). In vertebrates, NSC self-renewal can be viewed as a property of the entire population as opposed to just the single cellular entity, in that self-renewal is thought of more globally as the capacity to maintain the absolute number of neural stem cells in a given area at a steady level throughout time (Gritti et al., 2003). This more relaxed definition allows for fluctuations in absolute NSC population size, evenly balancing symmetric stem cell divisions with asymmetric differentiated progenitors as physiological conditions dictate.

Although there may be no true NSC capable of self-renewing indefinitely throughout adulthood in vivo, NPCs seem to maintain prolonged self-renewal (Ravin et al., 2008) and regain their multipotentiality when exposed to mitogenic growth factors in vitro (Doetsch et al., 2002; Gabay et al., 2003). In addition, transplantation of SVZ-derived NPC suggests that they retain both migratory and differentiation capabilities when homotopically reintroduced into appropriate anatomical locations, but may only differentiate and fail to migrate when heterotopically positioned (Betarbet et al., 1996; Zigova et al., 1996; Herrera et al., 1999; Yang et al., 2000; Ourednik et al., 2001; Tamaki et al., 2002; Seidenfaden et al., 2006). The results demonstrate that there may be an intrinsic spatiotemporal program that determines the developmental potential of NPC. This apparent positional restriction suggests heterogeneity of

hNPC as restricted populations of bipotent or unipotent glial and neural precursor cells in vivo; however, these findings do not exclude the possibility that precursor cells may reconfer multipotentiality in vivo (Ostenfeld et al., 2002b; Hitoshi et al., 2002; Parmar et al., 2002, 2003; Lepore et al., 2004; Kim et al., 2006; Kallur et al., 2006) upon long-term exposure to the local cellular milieu or complementary signaling of mitogenic growth factors from secondary expansion in vitro.

Recent evidence from Cre-lox and retroviral lineage mapping studies has challenged the past notion that NPC become fate restricted before adulthood (Garcia et al., 2004; Doetsch et al., 1999; Malatesta et al., 2000; Hartfuss et al., 2001; Miyata et al., 2001; Noctor et al., 2001; Gotz et al., 2002; Doetsch, 2003; Malatesta et al., 2003; Goldman, 2003; Gotz, 2003; Noctor et al., 2004; Merkle et al., 2004; Gotz and Barde, 2005; Noctor et al., 2008; Weissman et al., 2003; Martinez-Cerdeno et al., 2006). These data have led to three major changes in the tenets of developmental neurobiology (Merkle and Alvarez-Buylla, 2006; Malatesta et al., 2008). First, adult NSC are of glial origin, but are not fate restricted and can therefore give rise to all three central nervous system (CNS) cell subtypes. Second, embryonic, fetal, and adult NSC are uniformly lineage related through radial glia (RG), and, third, fetal NPC actually divide asymmetrically to increase the number of progeny they generate via symmetrically dividing intermediate NPCs (Alvarez-Buylla et al., 2001; Tramontin et al., 2003; Rakic, 2007).

The revised unified model demonstrates the lineal transition from neuroepithelium to radial glia and eventually adult astrocyte-like NPC and may help explain the relative heterogeneity that NPC display throughout embryonic and fetal development into adulthood. Furthermore, the unified theory proposes that functional CNS stem cells will display heterogeneous phenotypes throughout neural development depending on temporal and spatial cues, suggesting that the nature of in vivo NSC/NPC is highly dynamic. Moreover, the relative

heterogeneity and plasticity of the *in vivo* neurogenic niche suggests a similar component may exist within artificial *in vitro* dissociated stem cell culture preparations.

Although it has been published that passaging cells with enzymes results in “high risk of high rates of cell death, lack of adherence, or differentiation” (Nethercott et al., 2007) as well as induction of karyotypic abnormalities, utilizing the procedures described here, we have been able to maintain behaviorally normal, karyotypically stable, undifferentiated forebrain hNPC (Villa et al., 2004; Foroni et al., 2007) as highly proliferative, multilayer adherent networks for >100 passages without marked senescence or phenotypic adaptation by means of enzymatic (Accutase) single-cell dissociation. It is our opinion that overall expansion rates and possibly time to senescence (Carpenter et al., 1999; Goyns and Lavery, 2000; Wright et al., 2006) can be greatly increased by simply improving the overall condition of hNPC during and after dissociation, regardless of the technique employed. The repetitive combination of mechanical shear stress from trituration, centrifugation, and osmotic shock simply provides more opportunities to destroy the fragile neural progenitors and ultimately results in a gradual decline in hNPC numbers. Furthermore, we speculate that as the gross number of actively mitotic progenitors decreases, the subsequent loss of paracrine signaling (Taupin et al., 2000; Toda et al., 2003; Agasse et al., 2004, 2006) between hNPC eventually falls below a threshold concentration, whereby the delimited hNPC culture no longer maintains the capacity to properly condition its own basal substrate and subsequently becomes quiescently static, undergoing an irreversible halt in paracrine/autocrine regulatory signaling. The ultimate result of such events is a small population of nonproliferative hNPC in severe crisis; these cells are not suitable for study and should be distinguished from their proliferative counterparts and discarded.

We propose a model, whereby hNPC end-term senescence and proliferative potential is influenced by population density through “conditioned signaling” and can be controlled by

manipulating various combinations of these factors. Moreover, *in vitro* human manipulation can play a huge impact on the overall health and success of cultures, impacting the combined intrinsic signaling cascades that govern the phenotype of hNPC. On a global scale, the ultimate capacity for longterm self-renewal and ability to generate extremely large quantities of undifferentiated neural precursors (Svendsen and Smith, 1999) may be vastly improved with minimal adaptation to currently employed procedures. We therefore posit that the potential for somatic hNPC therapy and diagnostics would best benefit by a paradigm shift in culturing techniques from low- to high-density adherent populations, paying special attention to the importance of re-establishing essential cell-cell contacts. Investigating these properties may restructure the current theory of *in vitro* populations of somatic hNPC as limited-capacity progenitors (Hayflick, 1968; Temple and Raff, 1986; Durand et al., 1998; Svendsen et al., 1998; Quinn et al., 1999; Palmer et al., 2001; van Heyningen et al., 2001) incapable of amassing the relatively large quantities of cells (like their embryonic counterparts) necessary for regenerative therapies (Gottlieb, 2002).

## **2.11 Acknowledgements**

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## CHAPTER 3:

### BEHAVIORAL IMPROVEMENT IN A PRIMATE PARKINSON'S MODEL IS ASSOCIATED WITH MULTIPLE HOMEOSTATIC EFFECTS OF HUMAN NEURAL STEM CELLS

#### 3.1 Summary

Stem cells have been widely assumed to be capable of replacing lost or damaged cells in a number of diseases, including Parkinson's disease (PD), in which neurons of the substantia nigra (SN) die and fail to provide the neurotransmitter, dopamine (DA), to the striatum. We report that undifferentiated human neural stem cells (hNSCs) implanted into 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine- treated Parkinsonian primates survived, migrated, and had a functional impact as assessed quantitatively by behavioral improvement in this DA-deficit model, in which Parkinsonian signs directly correlate to reduced DA levels. A small number of HNSC progeny differentiated into tyrosine hydroxylase (TH) and/or dopamine transporter (DAT) immunopositive cells, suggesting that the microenvironment within and around the lesioned adult host SN still permits development of a DA phenotype by responsive progenitor cells. A much larger number of hNSC-derived cells that did not express neuronal or DA markers were found arrayed along the persisting nigrostriatal path, juxtaposed with host cells. These hNSCs, which express DA-protective factors, were therefore well positioned to influence host TH-ir cells and mediate other homeostatic adjustments, as reflected in a return to baseline endogenous neuronal number-to-size ratios, preservation of extant host nigrostriatal circuitry, and a normalizing effect on alpha-synuclein aggregation. We propose that multiple modes of reciprocal interaction between exogenous hNSCs and the pathological host milieu underlie the functional improvement observed in this model of PD.

### 3.2 Introduction

Degeneration of dopamine (DA) neurons in the substantia nigra (SN) and the consequent deficit of DA release in the striatum and other target areas appear to be responsible for the characteristic manifestations of Parkinson's disease (PD). Although substantial improvements result from the systemic administration of the DA precursor L-DOPA or DA agonists, such pharmacological replacement does not address the etiology of the disease, provide a permanent redress of the pathophysiology, or forestall progression of the degenerative process. It does, however, support the idea that DA provided by exogenous replacement cells might be therapeutic, a notion verified in rodents (1–3) and monkeys (4–6), where grafts of fetal DA neurons led to improvements in biochemical and behavioral indices of DA deficiency. However, in graft studies, the improvements in Parkinsonism have been limited and variable (see review in ref. 7). Therefore, we hypothesized that, in addition to DA replenishment, PD treatment should also restore functional equilibrium in the host SN-striatal system. A clinically relevant strategy might be to implant human neural stem cells (hNSCs) and progenitor cells constitutively capable of multiple actions, including neural differentiation and cytokine secretion, and allow them to develop within the PD affected brains of nonhuman primates to yield cells whose types, numbers, locations, and regulation are determined by the interplay of donor elements and the local host milieu. Outcomes derived from such donor–host interactions may result in a new level of bio-equilibrium among the DA-related neuro-structures (i.e., homeostasis), which could benefit behavioral recovery. hNSCs, either isolated directly from the developing normal brain (8–11) or derived from embryonic stem cells (12, 13), appear to be well suited for testing implementation of such a hypothesis. As the CNS' most primordial cell, the hNSC has attributes that appear to promote anatomical and functional preservation and/or restoration in neurodegenerative diseases. These properties include the potential for yielding appropriate ratios of cell types that constitute a normal

anatomical structure (i.e., both neurons and glia, plus other “chaperone-like” cells) (14–16). In addition, large numbers of hNSCs can be grown in culture as homogeneous, well-defined populations. For this study, we used hNSCs directly isolated from a neuroectoderm-derived structure, the telencephalic ventricular zone of normal, early second-trimester human cadavers (8). We selected two identically derived, non-immortalized hNSC lines (maintained *in vitro* as monolayers in serum-free, mitogen-supplemented medium) for their ability to engraft and migrate *in vivo* (8). One of these cell lines was known to pursue a ventral mesencephalic lineage when presented with appropriate cues *in vitro* (17) and to express a number of markers associated with a mesencephalic cell lineage [supporting information (SI) Figure 3-6]. We have reasoned that our hNSC-based comprehensive approach might better alleviate some of the limitations of previous strategies that placed partially differentiated cells, apparently without natural regulatory mechanisms, in ectopic locations such as the striatum (7).

### 3.3 Results

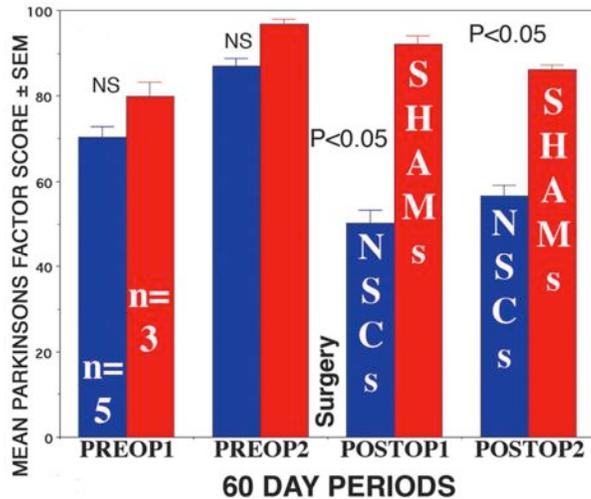
We studied 29 adult male African green monkeys of St. Kitts origin (*Chlorocebus sabaues*). Four were untreated normal control monkeys, and the remainder were injected systemically with 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP). This neurotoxin causes selective permanent bilateral destruction of mesencephalic DA neurons and their striatal projections, depletion of DA concentrations, and the full signs of Parkinsonism (4, 18, 19). hNSCs were injected stereotaxically into the right SN and bilaterally into caudate nuclei. The approach of implanting undifferentiated (as opposed to precommitted) hNSCs allowed us to investigate whether cues might be present in the host milieu that could permit, or even direct and sustain, an appropriate anatomical and physiological restoration. The animals were studied over periods of  $\leq 8$  months and categorized according to MPTP treatment,

immunosuppression, numbers of cells injected, and other treatment variables (Groups 1–5; see SI Table 3-1). To ensure reproducibility, hNSCs were obtained from two separate lines (designated as “H1” and “HFB2050”) that were initially derived by the same method: mitogen selection and expansion without immortalization (8). Numerous aliquots of early passaged cells were banked, thawed, and expanded as needed for new studies, hence minimizing cell variability from experiment to experiment over time.

To assess the possible impact of exogenous hNSCs on DA function, we studied a group of severely Parkinsonian monkeys (Group 1). Severity was determined with a well-validated and reliable behavioral scoring method consisting of time-sampled, quantitative behaviors and qualitatively rated items that reflect manifestations of Parkinsonism as well as normal behaviors in this primate species. A Parkinson’s factor score (PFS), derived from these observations, correlates inversely with postmortem striatal DA concentrations ( $r = -0.72$ ;  $n=18$ ;  $P<0.01$ ) (18, 19). Monkeys in this “most severe” category do not spontaneously or significantly improve over periods of greater than 1 year (18–20). Furthermore, the PFS in monkeys closely matches the 5-point Hoehn–Yahr scale, which is used clinically to categorize PD patients; the “most severe” category in monkeys corresponds to Stage 5 in humans.

Based on the PFS, eight monkeys that met the “Stage 5–Severely Parkinsonian” criteria were selected for study after their behavioral abnormalities were stable. Stage 5 monkeys show severe difficulty in ambulation, poverty of movement, delayed initiation of movement, lack of responses to food, difficulty eating, periods of “freezing” (remaining motionless for 5 sec), as well as head and limb tremors. The monkeys were randomly assigned to receive hNSC infusions or sham operations. Five hNSC-injected monkeys ( $10^6$  cells X3 sites per monkey) were compared to three monkeys that received sham surgical injections, with observations starting 4 months before and continuing to 4 months after surgery. The hNSCs, maintained and prepared to optimize engraftment, were injected into the SN and

caudate. These severely affected hNSC injected monkeys improved progressively and were significantly different from controls for the entire post-treatment period (Figure 3-1). These differences were highly significant functionally as well as statistically, and they included “activities of daily living” (such as ability to sit, walk, and self-feed) compared to the sham-injected monkeys, which were unable to do so. Although the hNSC-engrafted monkeys were less improved in the final 60-day period, at the end of the experiment, they remained significantly improved compared to their pre-implantation levels and compared to sham-operated monkeys, which remained severely Parkinsonian (Figure 3-1).

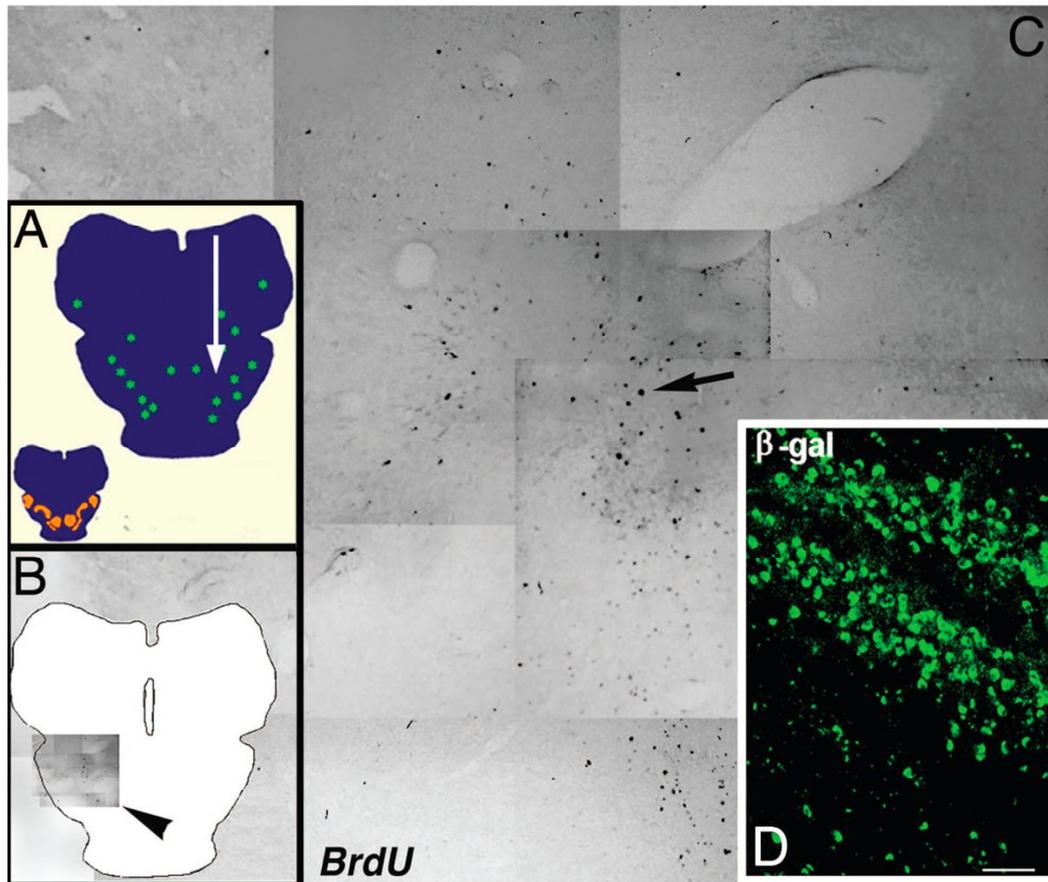


**Figure 3-1. Behavioral recovery in severely Parkinsonian monkeys after hNSC injections.** Severely Parkinsonian monkeys engrafted with hNSC (blue bars) demonstrated a significant decrease in a quantitative PFS, compared to sham operated monkeys (red bars), which remained severely Parkinsonian. Mean values  $\pm$ SEM are divided into 60-day periods (PREOP, before injections; POSTOP, after injections). After treatment, the hNSC group improved dramatically and significantly. ANOVA revealed a significant interaction among treatment group (hNSC vs. sham), treatment (before or after), and day of observation ( $F=65.87$ ,  $df=1,1096$ ,  $P<0.0001$ ). Tests of main effects showed that differences between the treatment groups were not significant before surgery ( $F=1.06$ ,  $df=1,6$ ,  $P=NS$ ), but became significantly different after ( $F=6.16$ ,  $df=1,6$ ,  $P<0.05$ ).

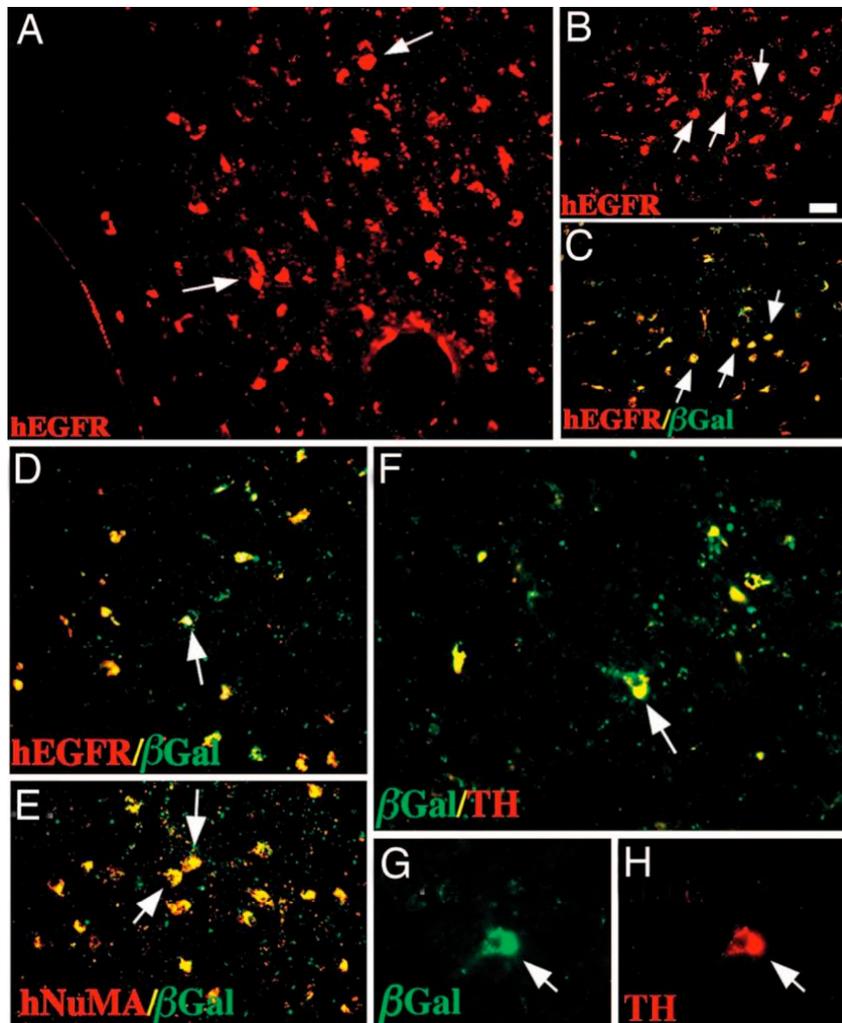
During formal (as well as extended periods of informal) observation of monkeys with “chimeric” human neural cell-bearing brain regions, there were no indications of any qualitative or quantitative behaviors that were not typical of the species, nor were any Parkinsonian dyskinesias noted. The duration and magnitude of functional recovery, compared to the controls, convinced us to terminate the behavioral experiment and begin a more extensive search for biochemical and histological correlates of improvement that might justify longer duration experiments and the investigation of immune and other factors and side effects that might help inform future long-term treatment of human PD.

To understand the basis for this functional recovery, histological sections from brains of these Group 1 monkeys and additional hNSC-injected monkeys were processed to assess the fate of donor and host cells [4 months after hNSC injections, designated as Group 2 (4-

month monkeys)]. Another group of MPTP-treated and hNSC-injected monkeys (Group 3) were studied and killed after 7 months (designated as 7-month monkeys). Four monkeys that were sham-operated but not MPTP-treated were controls (Group 4). In Group 1, although hNSCs were injected unilaterally immediately dorsal to the right SN (Figure 3-2A), we noted that donor-derived cells [identified by BrdU prelabeling (Figure 3-2C) and Beta-gal expression (Figure 3-2D)] were distributed bilaterally throughout the DA pathway, suggesting migration to the contralateral SN (Figure 3-2B and C) and/or migration from the engrafted ipsilateral caudate. Small numbers of donor-derived cells expressed tyrosine hydroxylase (TH) in the ventral mesencephalic region of Group 1 hNSC-injected monkeys (Figure 3-3). Such double-labeled TH-ir cells (identified by multiple independent markers) were not seen in non-lesioned hNSC-injected adult monkeys, although there was robust survival of hNSCs in all monkeys, whether normal or MPTP-lesioned. (In no monkeys were neoplasms, tumors, deformation, or overgrowth noted.)



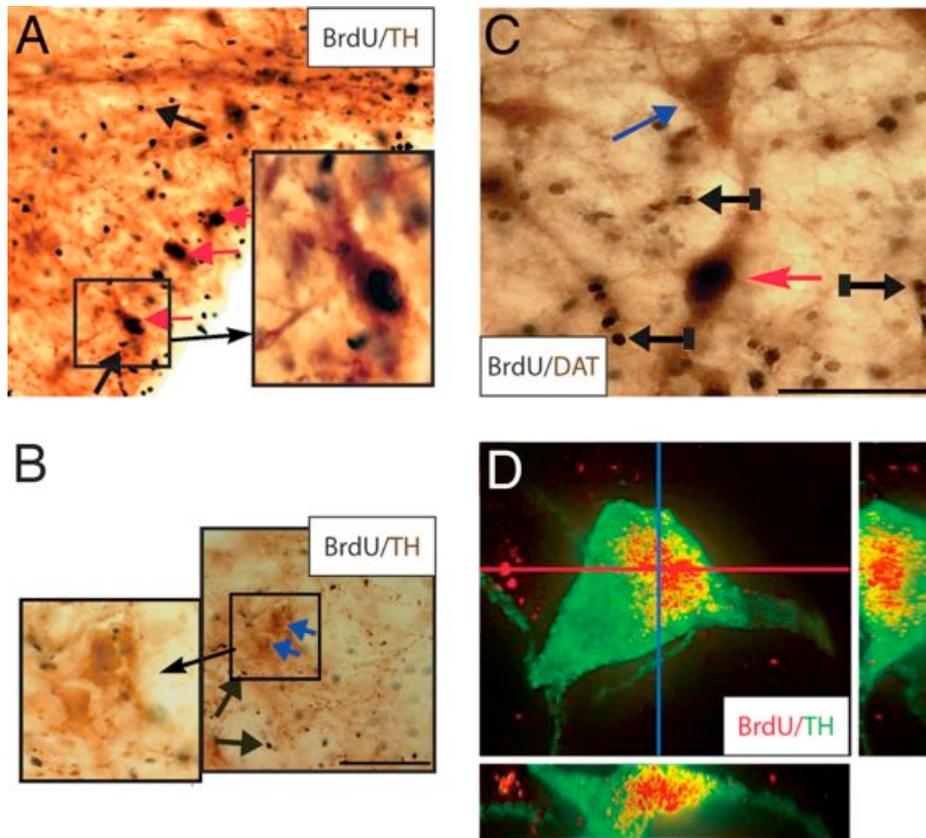
**Figure. 3-2. Survival and migration of engrafted hNSCs associated with functional improvement after transplantation into the nigrostriatal system.** (A) hNSCs were injected unilaterally, dorsal to the right SN (white arrow) in the monkeys studied behaviorally (Figure. 3-1, Group 1). Donor-derived cells were mapped (green stars) and detected throughout the area where DA nuclei are located (DA neuron distribution in this region is delineated in orange in the lower left corner, as recorded with a camera lucida). (B–C) Donor-derived cells labeled with BrdU and  $\beta$ -gal were also detected on the side contralateral to the implant, as shown by widely distributed BrdU-ir donor-derived hNSCs (black nuclei, black arrow) in the region (arrowhead in B and enlarged in C). (D)  $\beta$ -gal-ir cells (green) were present in substantial numbers in the ventral mesencephalon.



**Figure 3-3. Additional independent markers identify engraftment and survival of donor-derived hNSCs within structures relevant to Parkinsonism and differentiation of some of them into TH-ir cells.** Donor-derived cells in the ventral tegmentum are identified by antibodies to the human EGF receptor (hEGFR), nuclear mitotic apparatus (NuMA), or b-gal (arrows). EGFR-ir and NuMA-ir (*A* and *E*) co-localized with *lacZ*-expressing b-gal-ir (*C–E*) donor hNSCs. Furthermore, some donor-derived β-gal-ir cells (*G*) co-labeled with TH-ir cells (*H*) in the DA-deficient nigra (*F*) (merged). Confocal microscopic analysis of such cells (with optical dissection and *z*-stacks) is shown in Figure 3-4*D* and SI Figure 3-7.

To confirm the presence and numbers of hNSC-derived neurons expressing markers consistent with a DA phenotype, we injected hNSCs into six additional MPTP-lesioned monkeys and performed additional histological studies on them after >7 months (SI Table 3-1, Group 5). Although in the earlier studies no differences were noted between cyclosporine-

and noncyclosporine-treated animals, azathiaprine and prednisolone were added to cyclosporine in this later group of monkeys to increase immunosuppression. These animals showed extensive survival of hNSCs, yielding a variety of neural cell types, including significant numbers of TH-ir and DAT-ir expressing cells in the disabled SN (Figure 3-4 and SI Figure 3-7). Although such cells constituted  $\leq 1\%$  of donor-derived cells in the SN, they represented 4–7% of the total TH-ir cellular population in that region. Further, because of migration of the unilaterally injected hNSCs to the contralateral equally impaired SN, the percentage of TH-ir cells that were donor-derived was not significantly different between the two sides [implanted,  $6.74 \pm 1.75\%$  vs. unimplanted,  $5.99 \pm 1.74\%$ ;  $F(1,8) = 0.15$ ,  $P = \text{NS}$ ]. Accordingly, the actual concentrations of DA measured biochemically in punches from these regions were also statistically not different [ $t(3) = 0.087$ ,  $P$  value not significant]. The number of BrdU-ir cells that were also TH-ir was not significantly different from those that were also DAT-ir [TH-ir,  $6.37 \pm 1.23\%$  vs. DAT-ir,  $4.69 \pm 1.03\%$ ;  $F(1, 8) = 0.70$ ,  $P$  value not significant]. (Because not all hNSCs become pre-labeled *ex vivo* with BrdU, a larger number of TH-ir and/or DAT-ir cells in the SN may have been derived from donor hNSCs.) There was no significant difference between the total number of counted BrdU-labeled cells between the implanted and unimplanted sides ( $5,931 \pm 312$  vs.  $4,672 \pm 988$ , respectively) [ $t(3) = 1.86$ ,  $P$  value not significant].



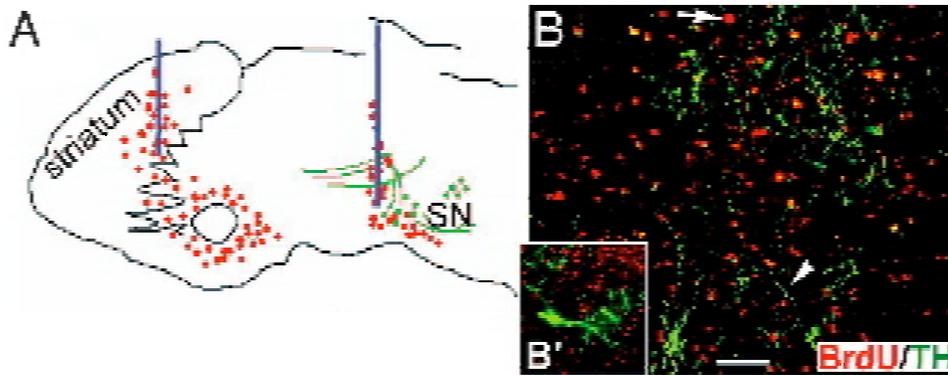
**Figure 3-4. Some hNSCs transplanted into the SN of MPTP-lesioned monkeys showed key markers of DA neurons.** (A) A black BrdU-ir nucleus indicates a donor-derived cell, with a small proportion also containing brown cytoplasmic TH-ir ( $6.75 \pm 1.28\%$  of all TH-ir cells, red arrows). (Inset) Magnification of blocked cells. (B) Most BrdU-ir cells are not TH-ir (black arrows). Compared with host BrdU-negative TH-ir cells in B (blue arrows), cells in A are most likely donor-derived. (Inset) Magnification of blocked host BrdU-negative TH-ir cells. (C) Some donor-derived BrdU-ir cells in this region were also immunoreactive for the DAT (red arrow with tail);  $3.91 \pm 1.04\%$  of DAT-ir cells were also labeled with BrdU. A DAT-ir neuron with a nucleus void of BrdU (blue arrow), presumably an endogenous host cell, is seen above the hNSC-derived neuron (red arrow), as well as many DAT-negative BrdU-ir (black nucleus) hNSCs (black arrows with tails) juxtaposed to DAT-ir fibers. (D) Double-label immunofluorescence of an hNSC cell expressing TH viewed by confocal microscopy with z-stacks; a BrdU-ir nucleus (red) is surrounded by a TH-ir cytoplasm (green). Red and blue lines indicate corresponding points in the orthogonal planes, confirming localization of the label within the pictured cell after the summation of serial optical sections. See also SI Figure 3-7. (Scale bars: B, 100  $\mu\text{m}$ ; C, 50  $\mu\text{m}$ .)

Besides bilateral distribution of hNSCs in the SN after a unilateral injection, the hNSCs and their progeny appeared to migrate from the depleted striatum toward the SN along the nigrostriatal pathway (Figure 3-5 A and B). Most donor-derived cells were found between

SN and the striatum and ventral to the SN. Donor hNSC-derived TH-ir cells were closely associated with host-derived cell bodies and TH-ir fibers in the SN and nigrostriatal pathway (Figure 3-5B). Indeed the close physical association suggested that stimulus–response intercellular relationships might be in process between donor-derived non-neuronal cells and host DA neurons and their fibers (Figure 3-5B). This nonrandom distribution pattern was reminiscent of one of the proposed routes followed by progenitors during embryonic emergence of the nigrostriatal functional unit and might suggest that this pathway can still be used by progenitors, with possible behavioral consequences in the adult primate brain with PD pathology. No cells in the striatum of any of the monkeys were double-labeled for TH-ir and B-gal, nuclear mitotic apparatus (NuMA), or other human-specific markers studied. (Because of the extensive migration of hNSCs and their progeny, it was not feasible to count the proportion of implanted cells that survived. A more detailed study of cell migration in these monkeys is in progress.)

We also noted significant increases in the size of host TH-ir neurons in the SN by 7 months after hNSC injections (Figure 3-5C and SI Figure 3-8 A and B), associated with the presence of donor hNSC that were not TH-ir or DAT-ir. Thus, hNSCs appeared to exert homeostatic effects on host circuitry, increasing the size of abnormally small endogenous TH-ir neurons of the SN toward normal values. MPTP-induced changes in the size and number of TH-ir host cells in the striatum were also normalized after hNSC injections (SI Figure 3-9). Although the molecules mediating the impact of hNSCs on host DA systems are unknown, some BrdU-ir cells expressed a marker associated with an astrocytic lineage (Figure 3-5G) and expressed glial cell line-derived neurotrophic factor (GDNF) (Figure 3-5H), a growth factor known to augment and/or protect DA systems (21–24). Also, increased aggregation of  $\alpha$ -synuclein has been reported after MPTP treatment in rodents (25) and primates (26). We found by immunohistochemical analysis of the nigrostriatal system in eight animals that  $\alpha$ -

synuclein aggregation was present in approximately 80% of cells in monkeys that were MPTP-exposed only, but aggregation was found in less than 20% of cells after hNSC implantation had followed MPTP exposure (Figure 3-5I and SI Figure 3-10). No aggregates were seen in non-MPTP-lesioned monkeys regardless of whether they were transplanted with hNSCs. In summary, hNSC implantation appeared to return a number of abnormalities after MPTP lesioning to the parameters seen in normal animals.



**Figure 3-5. hNSC engraftment is associated with multiple influences on the host DA nigrostriatal system that might contribute to the observed functional improvement.**

Migration of hNSCs, normalization of pathological numbers and sizes of host TH-ir cells, and effects on alpha-synuclein aggregation are shown, with evidence of secretion of a growth factor known to preserve fibers in the host nigrostriatal system. (A) Four and 7 months after hNSCs were placed in the SN and caudate, the majority of donor-derived BrdU-ir cells had migrated to the nigrostriatal pathway as illustrated in a composite of serially sampled sections from an entire brain. Red dots represent an approximate density and locations where the majority of BrdU-ir cells were found. Green dots and lines indicate host TH-ir cells and their fibers. Blue lines indicate the locations where hNSCs had been implanted. (B) Many non-neuronal hNSC-derived cells (BrdU-ir in red, marked by white arrow) were found in the SN and closely associated with host TH-ir cell bodies and their neurites (green, marked by white arrowhead). (Inset) Robust, healthy host DA neuronal soma with extensive processes (see also SI Figure 3-8 A and B). (C) In the SN, MPTP reduced the size of host TH-ir cells, which were then significantly increased 7 months after hNSC injections, compared to sham-operated MPTP-lesioned monkeys [ANOVA post hoc group differences; \*, smaller than corresponding control group only; \*\*, smaller than all other treatment groups ( $P < 0.05$ )]. (D) Endogenous TH-ir cells are also found in small numbers in the primate striatum. The arrow points to the most prominent type of striatal TH-ir neuron, which is small and bipolar (see SI Figure 3-9 A–D). (E and F) Their size-to-number ratios become disordered after MPTP lesioning. After MPTP lesioning, striatal TH-ir neurons increase in number (E) and decrease in size (F), a compensatory but abnormal change. They do not restore DA function and, in fact, are at their peak in animals that show the greatest signs of DA deficiency. In monkeys receiving hNSC implants, the aberrant size-to-number relationships of striatal DA neurons return to near normal control parameters (see SI Figure 3-9). (G and H) Some hNSCs (BrdU-ir cells, black nuclei) along the nigrostriatal pathway were also immunoperoxidase-positive (brown cytoplasm) for glial fibrillary acidic protein (G) and GDNF (H), suggesting that they had differentiated into astrocytes spontaneously and constitutively produced this trophic factor as a potential mechanism for hNSCs' effects on host neurons. (I) hNSCs transplanted into MPTP-lesioned monkeys appeared to diminish the alpha-synuclein-ir aggregation pattern (arrows) in the host striatum, approximating a more normal profile (as seen in non-lesioned monkeys with and without hNSCs). (Scale bars: B, 100  $\mu$ m; D, 20  $\mu$ m; H and I, 50  $\mu$ m.)

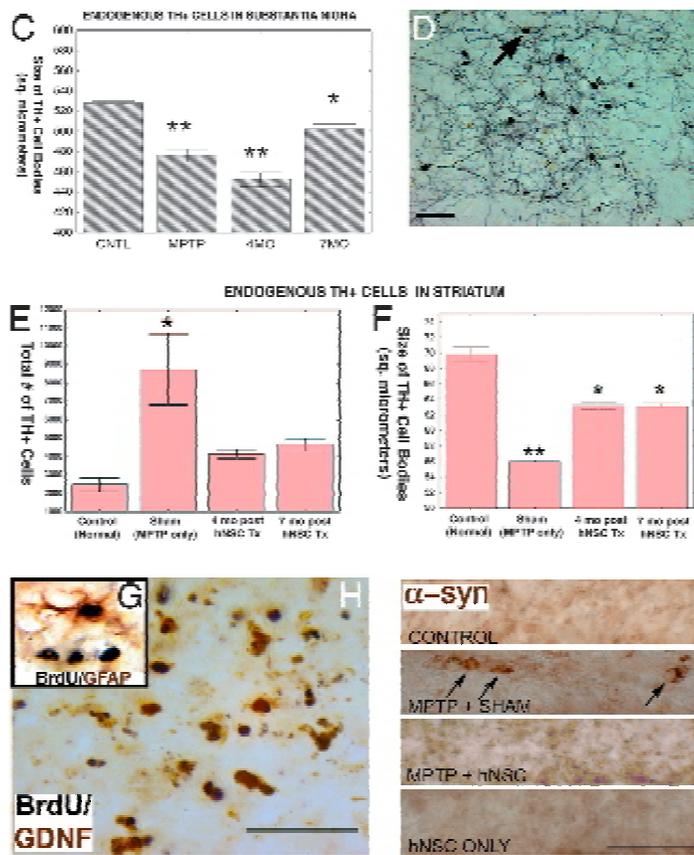


Figure 3-5. Continued

### 3.4 Discussion

Our studies demonstrate that the MPTP-lesioned adult monkey brain retains intrinsic microenvironmental signals that may direct differentiation of an uncommitted human stem cell toward a DA phenotype and suggest that hNSCs have the capacity to respond to a DA deficiency (27) even without pre-induction by factors or transgenes. However, the predominant functional action of hNSCs in the presence of damage to DA systems was most likely one of promoting homeostatic adjustment of host nigral DA neurons and their nigrostriatal projections. Some of the hNSCs, particularly those juxtaposed to host cells and fibers along the nigrostriatal trajectory, pursued an astrocytic lineage, which included expression of the neurotrophic factor GDNF. This observation is consistent with previous findings that epigenetic signals promoting the differentiation of Nurr1-expressing precursors may emanate from neighboring astrocytes (28). In fact, the role of the astrocyte in directing neurogenesis (15), mediating rescue (14), and potentiating the function of other neural cells is becoming increasingly appreciated (29). In particular, GDNF (one of possibly several natural products of the astrocytic progeny of these hNSCs) shows developmental, trophic, and protective support of DA neurons and promotes effective processing and release of DA (21–23). The normalization of  $\alpha$ -synuclein aggregation in the presence of injected hNSCs in this study illustrates another potentially beneficial effect of hNSCs, but is probably independent of GDNF (30).

Most prior studies have focused on the concept that the host environment, as it changes during the course of development and aging, or after injury or cell degeneration, influences the transplanted stem cell, as exemplified here by the homeostatic emergence of some donor-derived TH-ir and DAT-ir cells. Based on past reports that small reversals of DA depletion can underlie large functional improvements, even a small elevation in DA might be

behaviorally relevant, whether from stem cell differentiation into DA neurons or from preservation and even augmentation of host DA pathways via hNSC-derived trophic/neuroprotective effects. This study also reveals that improvement in function might result from significant reversal of abnormalities in sizes and distributions of endogenous TH-ir cells in the nigrostriatal system as well as an interesting normalization of alpha-synuclein aggregation. These effects are consistent with other studies that have shown NSCs yielding multiple interacting cell types, not only key effector neurons but also undifferentiated progenitor cells that mediate neuroprotection and neuroplasticity (31) and glia that nurture, detoxify, myelinate, or direct the differentiation of neurons (32, 33). NSCs have also been suggested to restore equipoise to a disequibrated milieu by fueling cell turnover (8, 34) and regulating gene expression and signaling pathways (35, 36). We believe that our data suggest, therefore, that the Parkinsonian primate CNS may benefit from such homeostatic effects, including (i) replacement of degenerating DA neurons by differentiated human stem cells, and (ii) the trophic, protective, and guidance effects of non-neuronal stem cell-derived progeny. These latter actions may manifest themselves by promoting recovery through the variety of effects described here, as well as by others that remain to be elucidated by additional experiments and controls. Although long-term studies of these effects and potential side effects, such as dyskinesia (although not observed in the present study) and possible immunorejection of exogenous stem cells, are needed before attempting clinical application, this report provides evidence that permissive signals are present in the milieu, and that stem cells respond with multiple homeostatic actions to restore functionality to an adult primate brain that presents with severe Parkinsonian pathology.

### **3.5 Materials and Methods**

#### **3.5.1 Source and Maintenance of hNSCs**

Cells were obtained from stable, self-maintaining populations of hNSCs dissected from the ventricular germinal zone of a 13-week-old human fetal cadaver (8, 37) and maintained in neurobasal (Gibco/ Invitrogen, Grand Island, NY) medium supplemented with N2 or B-27 (Gibco) plus bFGF (20 ng/ml) (Chemicon International, Temecula, CA), heparin (8 mcg/ml), and LIF (5 ng/ml) (Chemicon International). Both adherent cells and floating clusters were chemically dissociated with Accutase (Chemicon International), triturated, and passaged every 3 to 10 days. hNSC lines (8, 37) were propagated with mitogens alone. More details are provided in *SI Materials and Methods*.

#### **3.5.2 MPTP Lesioning of Monkeys and Treatment Groups.**

Five groups of monkeys were studied with or without injections with MPTP-HCl (RBI/Sigma–Aldrich, Natick, MA). Seventeen monkeys received cumulative doses of 2.25 mg/kg over a 5-day period to induce degeneration of the nigrostriatal pathway, and six monkeys received 1.5 mg/kg aimed to produce DA depletion but without functional impairments. Seven monkeys were sham-injected, and 20 received hNSCs injections. Four monkeys, which were not treated with MPTP, were studied as controls (see SI Table 3-1 for individual details of cell numbers, immunosuppression, cell types, and numbers). The animal experiments were approved by the relevant institutional animal care and use committees of the collaborating institutions.

#### **3.5.3 Behavioral Scoring and Statistical Analysis.**

Blinded observers scored the MPTP-treated monkeys by using a published and

validated quantitative time-sampling method (4, 18) two periods per day, 5 days per week, a regime that has been shown empirically to sample Parkinsonian behaviors efficiently and accurately. Statistical analysis of behavioral changes used a multifactor ANOVA of the daily PFS of each monkey. All 1,304 individual observations were analyzed in 60-day blocks from 120 days before to 120 days after hNSC implantation, when monkeys were killed; >95% concordance was recorded among five blinded observers for all behaviors tested.

#### **3.5.4 Preparation and Transplantation of hNSCs.**

hNSCs were injected into the right SN and the right and left caudate nuclei by using stereotaxic procedures. Donor-derived cells were identified by multiple independent techniques. Dissociated hNSCs were preincubated *ex vivo* with BrdU for 48 to 72 h *in vitro* before transplant. Some hNSCs were subcloned to stably express the *lacZ* transgene and produce B-gal. Control experiments confirmed that donor cell-specific markers were never transferred to host cells after cell destruction; donor cells never produced recombinant replication-competent helper virus. hNSCs were dissociated and passaged 24 to 48 h before transplantation to help synchronize and make as uniform as possible their state of differentiation and stage in the cell cycle (8, 37). The number of hNSCs injected was 1,000,000 cells in most animals, although it ranged from 1 to  $8.75 \times 10^6$  in Group 5. Additional monkeys were MPTP-treated only and received injections of vehicle or needle passage alone (“sham-transplanted”) (see SI Table 3-1).

#### **3.5.5 Histological Analyses.**

Donor-derived and host cells were distinguished in postmortem fixed tissue by immunocytochemistry with antibodies against multiple independent markers, including BrdU, B-gal, and human-specific epitopes (8, 37, 38). Unbiased stereology was used for counting

labeled cells. The number and size of TH-ir immunoreactive (ir) cells in the caudate and SN were quantified, and BrdU-ir, TH-ir and DAT-ir cells were calculated as a percentage of total cells of each type. Immunocytochemistry for alpha-synuclein was performed with standard methods and primary antibody (Cell Signaling Technology, Danvers, MA) at 1:500 dilution and second antibody (Jackson ImmunoResearch, West Grove, PA) at 1:500 dilution and second antibody (Jackson ImmunoResearch, West Grove, PA) at 1:200.

### 3.6 Acknowledgements

Chapter 3, is a reprint in part of material as it appears in *Proc Natl Acad Sci U S A*. 2007 Jul 17;104(29):12175-80 with co-authors Redmond DE Jr, Bjugstad KB, Teng YD, Ourednik V, Ourednik J, Parsons XH, Gonzalez R, Blanchard BC, Kim SU, Gu Z, Lipton SA, Markakis EA, Roth RH, Elsworth JD, Sladek JR Jr, Sidman RL, Snyder EY. The dissertation author was a co-author of this paper. DRW performed transplantations, immunohistochemistry/ cytochemistry, analyzed data, co-prepared manuscript, co-assembled figures, and edited all text. DER was principal investigator, performed transplantations and oversaw all animal operations, co-prepared entire manuscript, and managed the project. KBB performed stereotactic counting. BCB, EAM and ZG performed IHC. JRS prepared histological figures. YDT, VO, JO, RG, XHP cultured cells. RHR, JDE, RLS, EYS, SAL, SUK supervised. Together with his coauthors, we thank the staff of St. Kitts Biomedical Research Foundation for their contributions to the *in vivo* primate studies, Csaba Leranthy and Robert Makuch for histological and statistical advice, and Marcel Daadi for advice and studies differentiating hNSCs into DA neurons. This work was supported by National Institute of Neurological Disorders and Stroke Grants RO1NS40822, PO1NS44281 (to D.E.R.), and R21NS053935; Veterans Affairs Biomedical Laboratory Research and Development Grant 121F (to Y.D.T.); the National Institutes of Health/National Institute of General Medical

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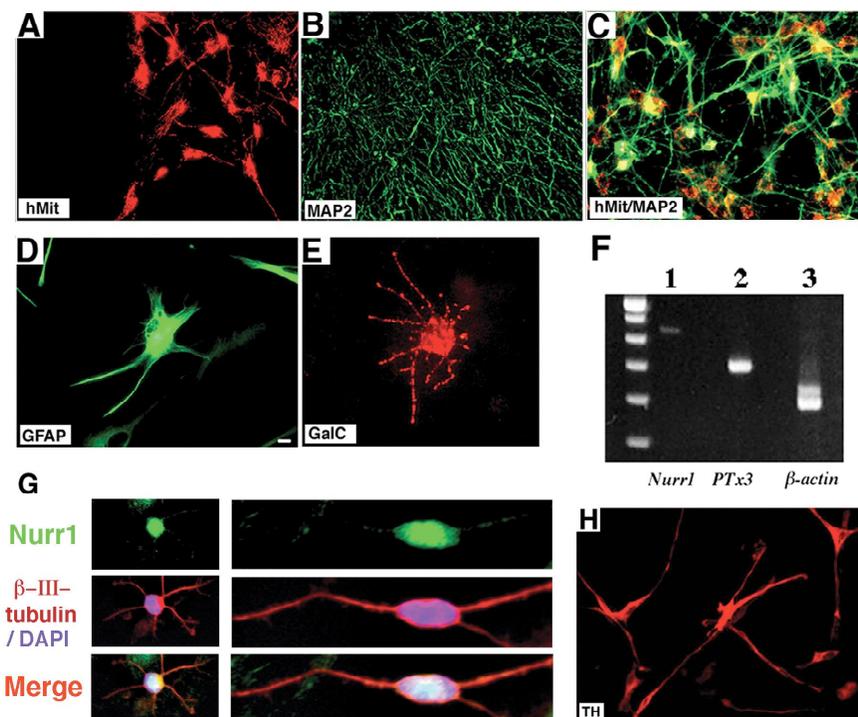
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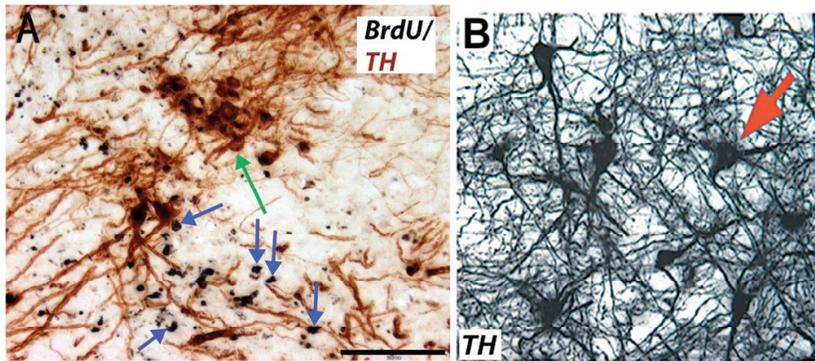
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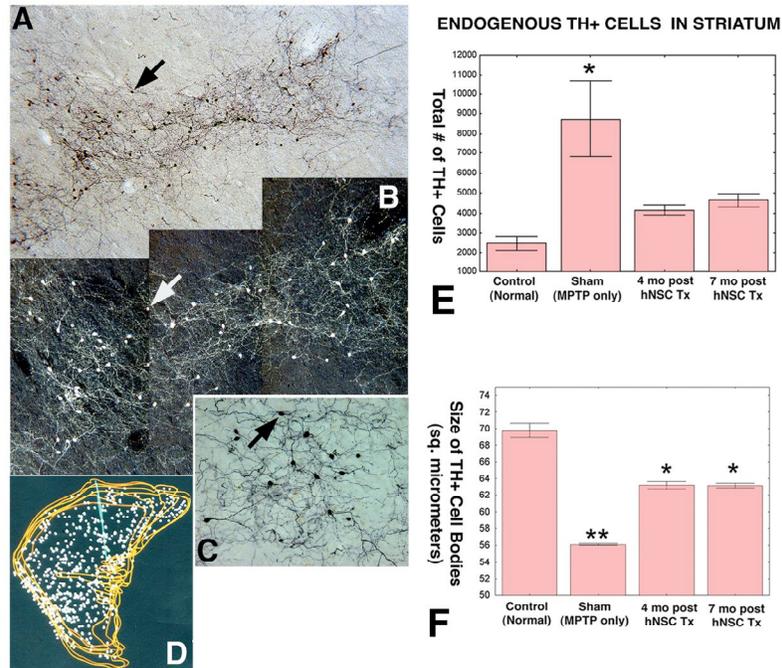
## 3.8 Supplemental Figures and Methods:



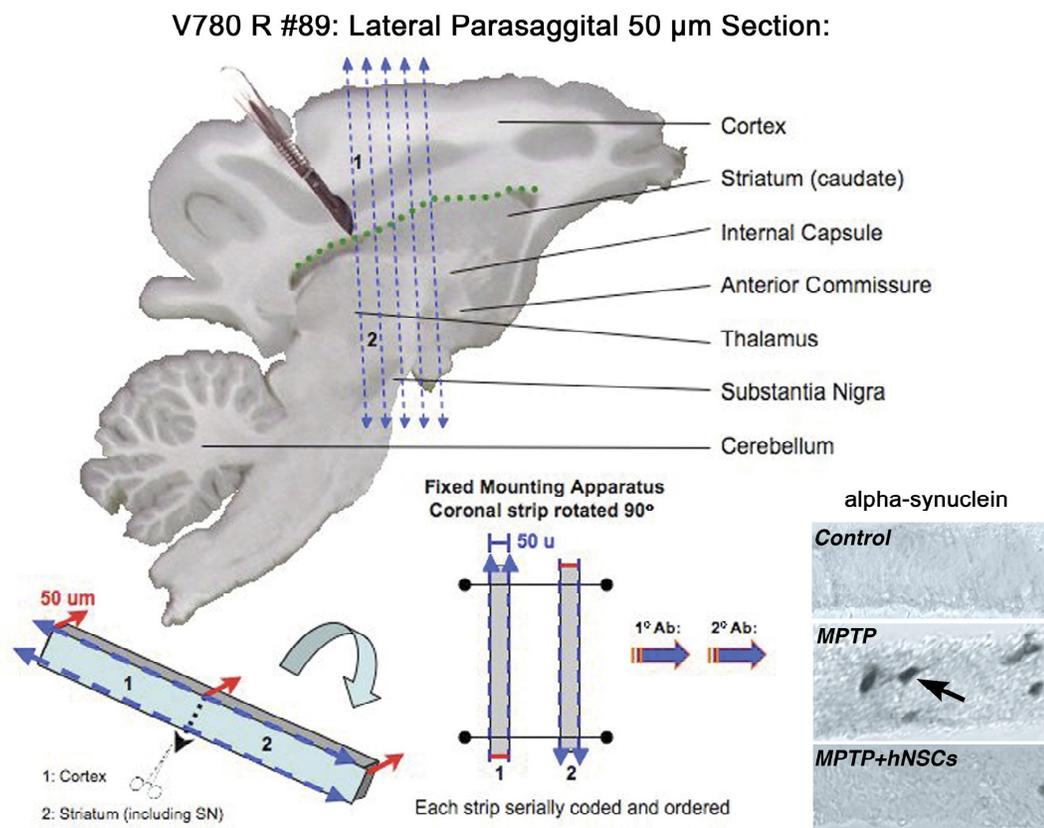
**Figure. 3-S6.** Human neural stem cells used in these studies show a number of markers associated with potential development into a DA phenotype when presented with appropriate cues in vitro. (A) Human NSCs were identified in vitro by human-specific markers as illustrated by their immunoreactivity to a monoclonal antibody against human mitochondria (hMit). (B-E) The hNSCs differentiated into mature neurons [MAP-2 (B), dual-labeled with hMit in (C)], astrocytes [GFAP (D)], and oligodendrocytes [GalC (E)]. (F) When maintained under appropriate culture conditions (27), the hNSCs expressed midbrain-specific molecular markers, Nurr1 and Ptx3 (the RT-PCR amplification product shown). (G) Prolonged culture produced neurons that co-labeled for  $\beta$ -III-tubulin (red), Nurr1 (green nuclear/perinuclear-ir product), or TH (H).



**Figure 3-S7.** hNSC engraftment normalizes some MPTP-induced changes in host DA neurons in the SN of the recipient Parkinsonian monkey. (A) Many undifferentiated hNSC (BrdU+, black nuclei, blue arrows) were located within and just ventral to the SN 7 months after transplantation. Furthermore, they were often closely juxtaposed to large BrdU-negative, endogenous TH+ (brown cytoplasm) cell bodies and their (brown) fibers (green arrows) (see also Figure 5 A-C). Typically, MPTP decreases the size of surviving nigral DA neurons, as well as their numbers. In MPTP-lesioned monkeys transplanted with hNSCs, however, as shown in B, host SN neurons were significantly increased in size 7 months after grafting. Note large exuberantly TH+ host nigral DA neurons (arrow). These sizes are quantified in Figure 5.C.



**Figure 3-S8.** hNSC engraftment also normalizes MPTP-induced changes in the Parkinsonian primate host striatum (i.e., aberrant size-to-number ratios). (A-D) Endogenous TH<sup>+</sup> cells are also found in the primate striatum. A representative overview of TH<sup>+</sup> cells in the caudate post-transplantation (in a representative monkey whose behavior was analyzed in Figure 3-1) is presented in brightfield (A) and in darkfield (B). Note the exuberant network of extensively branching TH<sup>+</sup> processes. At higher magnification (C), arrows indicate a typical TH<sup>+</sup> caudate neuron, characteristic of >95% of neurons in the caudate: small (60 mm), TH-immunoreactive, bipolar with smooth or varicose processes. It is this cell, meeting these criteria (similarly indicated by arrows in A and B), that was analyzed to yield the quantitative data (E and F) (see also Figure 3-5 D-F). The computer reconstruction (D) demonstrates the dense yet uniform and extensive distribution of endogenous dopaminergic neurons throughout the caudate of this monkey (as culled from multiple serial parasagittal 50mm histological sections) (needle track is indicated as a blue line). (E and F) The size-to-number ratios of these endogenous striatal TH<sup>+</sup> cells become disordered following MPTP-lesioning: they increase in number (E) and decrease in size (F), a compensatory but abnormal change with unclear functional consequences. They do not restore DA function and, in fact, are at their peak in animals that show the greatest signs of DA deficiency. In monkeys receiving hNSC implants, the size and number of striatal DA neurons return to normal control parameters (E) and (F) (repeated from Figure 3-5 E and F). Details of specific identifiable TH cell subtypes and their normalization after hNSC implantation are reported elsewhere (34).



**Figure 3-S9.** Processing of monkey brain to provide tissue for  $\alpha$ -synuclein immunohistochemistry. Three serial sections each  $\sim 200$ - $300$  mm wide were cut from lateral to medial surfaces from an existing 50-mm parasagittal tissue slice in an orientation indicated by the blue dotted lines. Individual strips were then divided at the green dotted line into region 1, consisting principally of cerebral cortex, and region 2, including principally the striatum and extending ventral to the nigra (below the green dotted line). These regions were numbered for record keeping and sectioned for  $\alpha$ -synuclein. Multiple samples from two or three monkeys in each of the following four experimental conditions were analyzed by two observers for extent, distribution, and pattern of  $\alpha$ -synuclein immunoreactivity in both cortex and nigrostriatum (see SI Table 1). Group 1: Not MPTP lesioned, not transplanted with hNSCs ( $n = 14$  samples); Group 2: Not MPTP lesioned, transplanted with hNSCs ( $n = 7$  samples); Group 3: MPTP lesioned, sham-treated but not transplanted with hNSCs ( $n = 37$  samples); and Group 4: MPTP lesioned, transplanted with hNSCs ( $n = 27$  samples). Representative photomicrographs of  $\alpha$ -synuclein-immunostained sections are shown in Figure 3-5 and here (from Groups 1, 3, and 4). The arrow indicates a cell immunopositive for  $\alpha$ -synuclein.

## **SI Materials and Methods**

### **Parkinsonian and Healthy Behavior Score and Scoring Methods.**

Blinded observers who are trained and maintain a coefficient-of-concordance (Kendall's)  $>0.95$  in their ratings of all behaviors scored the monkeys using a time-sampling method for the presence or absence of 20 normal and abnormal behaviors, in addition to ratings of a number of Parkinsonian signs (1-3). Monkeys were observed and scored two periods per day, 5 days per week, a regime that has been shown empirically to sample Parkinsonian behaviors efficiently and accurately. Monkeys were selected into treatment groups based on quantitative scores 1 month after MPTP exposure. A Parkinsonian factor score (PFS) was derived from a prior large study involving a principal component factor analysis on 15,000 individual behavioral observations from 77 MPTP or control monkeys, and the score was averaged per week of observations. For the purposes of predicting outcome and assigning monkeys to treatment groups, the monkeys were classified into five severity categories based on their Parkinsonian scores. These were quintiles from the initial study. Monkeys with the highest severity category, which, like Parkinson's patients, have previously been shown not to recover spontaneously, were assigned to hNSC treatment or sham surgery for the studies of function (2). Validity of the score has been tested by a number of methods. The score is highly responsive to pharmacological changes in DA function and correlates highly (inversely) with striatal DA concentrations postmortem (4). These categories at 1 month predict the extent of postmortem DA damage. Categorizing monkeys in this fashion at 1 month allows the selection of treatment groups that have similar extent of DA depletion and are known to remain significantly parkinsonian over long periods. Although the categories were derived empirically and mathematically, the five categories correlate well with the five levels of the Hoehn and Yahr scale, which is used clinically to categorize PD patients.

Eight monkeys that matched criteria for "severe Parkinsonism" (corresponding to Hoehn and Yahr category 5) were randomly assigned to receive hNSC injections or sham surgery. A total of 1,304 observations of the monkeys were made from a period from 120 before to 120 days after hNSC implantation. Five monkeys implanted bilaterally into the caudate nucleus and unilaterally dorsal to the SN with »1 million hNSCs per site were compared with three monkeys that received sham surgery using a multifactor analysis of variance. All monkeys were observed using the above-described standard behavioral scoring system (2-10).

### **Induction of Markers of Mesencephalic Precursors by hNSCs in Vitro.**

This procedure is derived from that reported by Daadi and Weiss (27). hNSCs are induced to differentiate by first omitting growth factors from the medium. During the first week of differentiation, 7 days in vitro (DIV), ~5% of the total population express the neuronal marker b-tubulin class III. After 15 days in vitro, ~60% of the differentiated progeny express b-tubulin and MAP2. Of the remainder, 38% are astrocytes as identified by their immunoreactivity to GFAP and <2% are oligodendrocyte that express the cell surface antigen galactocerebroside (SI Figure 3-6). To induce the dopaminergic phenotype, hNSCs are single cell dissociated at  $0.5 \times 10^6$  cell per ml and plated at a density of  $2.5 \times 10^5$  cell per cm<sup>2</sup> on PLO-coated glass coverslips in 24-well culture dishes containing the baseline differentiation medium. After a 2-h culture period, the baseline differentiation media is replaced by the TH-inducing media containing 20 ng/ml of bFGF and 75% (vol/vol) of glial conditioned media (CM).

CM is prepared from cultures of postnatal striatal astrocytes and from the B49 rat glial cell line. B49 cells are cultured in DMEM/10% FBS until confluency. Confluent astrocyte or

B49 glial cell cultures are rinsed once with PBS and twice with serum-free DMEM/F12 (1:1) with hormone mix and replaced in the incubator with 20 ml of the same medium. The CM is collected after 24, 48, or 72 h and centrifuged at 1,000 'g and 2,000 'g to remove cellular debris. The CM is carefully removed, filtered, aliquoted, and stored at -80°C until used. The CM likely fulfills a role similar to that reported by Wagner et al. (28).

The cultures are incubated and investigated for the TH induction within a period of 7 DIV. In each culture, the neuronal and dopaminergic lineage species are determined using immunocytochemistry for b-III-tubulin and TH, respectively. These cultures demonstrate a stable and steady increase of the TH-IR cells in cultures. After 24 h in culture, 1% of the total cells express TH, which represents ~30% of the total number of b-III-tubulin-IR neurons. The number of TH-ir cells steadily increases to 3% of the total DAPI positive live cells after 7 DIV. This proportion of TH-ir cells represents ~76% of the total b-III-tubulin-ir neuronal population (SI Figure 3-6). In vivo, the specification and maintenance of the midbrain DA fate during development is controlled by the nuclear receptor transcription factor Nurr1 and the homeobox transcription factors engrailed and Ptx-3. Semi-quantitative RT-PCR analysis confirms that DA-induced cultures (i.e., FGF2+CM-treated hNSCs after 7 DIV) markedly up-regulate expression of Nurr1 and Ptx-3, supporting the acquisition by these forebrain-derived hNSCs of a molecular profile consistent with mesencephalic dopamine precursors (SI Figure 3-6). The expression of Nurr1 and Ptx-3 without gene transduction in these cultures, a technique that has been previously used (12), suggests that they would be able to develop into dopaminergic neurons in vivo if exposed to the appropriate conditions.

### **Immunosuppressive Drug Administration.**

Approximately half of the monkeys in each experimental and control group were

immunosuppressed with cyclosporine (0.6 mg/kg) in the behavioral and initial histological studies. In the final triple "immunosuppression" study, all monkeys also received prednisolone (0.3 mg/kg) and azathioprine (0.5 mg/kg) twice daily starting 3 days before implantation and continuing until they were killed, a dose that was tolerated by the monkeys.

### **Quantitative Morphology.**

Unbiased stereology was used for counting all labeled cells. The number and size of TH cells in the caudate and SN were quantified, and BrdU+, TH+, and DAT+ cells were calculated as a percentage of total cells of each type. For cell size, a digital camera was used to take 4 to 10 pictures of TH-positive cells found at three different levels of the caudate and corresponding SN (Olympus MicroSuite software; Olympus, Tokyo, Japan). Each picture contained 1 to 15 different cells. Scion Image for Windows (Scion Corp., based on NIH Image by Wayne Rashan, NIH, Bethesda, MD) was used to measure the diameter and area of each cell pictured. ANOVA based on weighted means with posthoc Fisher's LSD was used to determine significant differences in cell soma size. Total cell counts were computed from the mean number of cells in five slides taken from a 2-mm lateral expanse and multiplied by the number of sections in 2 mm. For total cell counts, a standard ANOVA, with post hoc tests, was used to determine significance between groups of animals at  $P < 0.05$  (two-tailed). In the Group 4 monkeys, this procedure was adjusted to compensate for a postmortem punch to obtain tissue for analysis of monoamines. Paired t tests of the adjusted cell counts ( $P < 0.05$ , two tailed) compared the injected versus the uninjected sides. Double-labeled TH/BrdU, TH (endogenous), DAT/BrdU, DAT (endogenous), and BrdU alone (NSC-derived) were statistically analyzed (five tests). None was significant.

### **Statistical Analyses of Behavioral Effects.**

For the behavioral studies, sample size was estimated from the numbers required to show significant effects in previous studies of fetal tissue transplantation in MPTP-treated monkeys. All of the monkeys at a Stage 5 severity level were studied behaviorally because, as confirmed by our prior studies (2, 10), this level of severity does not spontaneously recover even after observation periods of >1 year. Monkeys were randomly assigned to receive hNSC's or sham surgery. Because we anticipated the possibility that actual cell engraftment could have adverse consequences (either perioperatively or afterward), the group sizes were skewed to have more hNSC animals than sham-operated animals. Data analyses were done using methods we have tested repeatedly to construct a PFS and Healthy Behavior score (similar to the clinical rating subscore, "activities of daily living." Significance was determined by ANOVA using the multifactor repeated measures model described by Winer (33) and carried out using the Statistical Analysis System. The two-tailed significance level was set at  $P < 0.05$ . As described in Figure 3-1, ANOVA revealed a significant interaction between treatment group (hNSC vs. sham) and treatment (before or after) and day of observation ( $F = 65.87$ ,  $df = 1,1096$ ,  $P < 0.0001$ ). Tests of main effects showed that differences between the treatment groups were not significant before surgery ( $F = 1.06$ ,  $df = 1,6$ ,  $p = NS$ ), but became significantly different afterward ( $F = 6.16$ ,  $df = 1,6$ ,  $P < 0.05$ ). In summary, hNSC transplantation had a statistically significant and sustained impact on improving behavior (i.e., diminishing the PFS while improving the Healthy Behavior Score). These differences were functionally significant and included the ability to perform critical activities of daily living, compared to the controls, which could not.

We also performed linear regressions on the PFS data to determine whether there were significant trends toward improvement before hNSC injection in either the active hNSC or the

sham-injected group and whether any slopes were significantly different between the groups (Statistical Analysis System). This analysis examined the possibility that the Stage 5/severely Parkinsonian monkeys were already spontaneously improving before hNSC transplantation. There were no differences in the highly significant increases in Parkinsonism in both groups during the first 60 days of observation, but both groups were stable (had no significant slopes) and were not different from each other in the 2 months before surgery. After surgery, the hNSC group showed a highly significantly slope of improvement ( $P < 0.001$ ). The sham-injected animals also showed a statistically significant improvement ( $P < 0.01$ ), but this slope was tiny and was different from the hNSC animals. In the final 60-day period, the shams had no significant slope and remained highly symptomatic, whereas the hNSC's appeared to worsen slightly, consistent with the interpretation that differentiated TH+ and possibly other hNSC cells were being rejected over this time period. They remained dramatically better, however, than before the hNSC injections. The significance of linear regression slopes were determined using Statistical Analysis System, and predicted slopes and 95% confidence intervals were calculated and plotted to compare the two treatment groups.

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## CHAPTER 4:

### **HUMAN NEURAL PROGENITOR CELL GRAFTS PROJECT NEURITIC PROCESSES ALONG HOST CIRCUITRY CONCOMITANTLY WITH AAV5-GDNF CHEMOTAXIS IN THE MPTP-LESIONED PRIMATE BRAIN**

#### **4.1 Summary**

Human fetal and embryonic stem cell (hESC) derived neural precursor cells (hNPC) have been suggested as a renewable alternative substrate to fetal dopaminergic (DA) neurons/progenitors for transplantation in Parkinson's disease; however, little is known about the long-term potential of these grafts to differentiate and integrate into anatomically correct loci in the adult non-human primate brain. We transplanted undifferentiated hfNPC that were derived from human fetal forebrain subventricular zone homotopically into the ventral midbrain of MPTP-lesioned monkeys. In the same animals, we concomitantly delivered striatal, glial-derived neurotrophic factor (GDNF) by an AAV serotype5 vector. The fate of grafted cells was assessed after 11-months in vivo. Donor soma remained predominantly within the ventral mesencephalon in the area of injection and extended numerous, morphologically-relevant, monoamine fiber types including both smooth and beaded varicose profiles. While these neurofilament positive neurites appeared to project in multiple directions, they also coursed in trajectories, often circuitous, to adjacent disease relevant targets, such as the substantia nigra. Donor processes paralleled tyrosine hydroxylase positive fibers of the host nigrostriatal tract, but did not differentiate substantially into fully mature, striatally-integrated, A9 dopaminergic neurons. This work demonstrates that human fetal NPC are capable of generating neuronal phenotypes long-term, retain the capacity to direct axonal projections with trajectories comparable to the intrinsic nigrostriatal pathway, and respond to

specific local endogenous signaling cues in the adult dopamine-depleted primate brain; suggesting that the adult primate brain retains specific axonal guidance cues and maintains a permissive environment for xenotransplantation of hNPC and possible pathway construction.

## 4.2 Introduction

Human stem/precursor cells derived from multiple cellular origins<sup>1</sup> have been touted for their potential as regenerative substrate in a broad range of human disorders. Experimental evidence supports the functional benefit of cell based therapies for diseases of the blood, bone, pancreas, and central nervous system (CNS)<sup>2-5</sup>, specifically neurodegenerative disorders such as Parkinson's disease (PD)<sup>6-15</sup>. Degeneration of midbrain A9 DA neurons in the substantia nigra pars compacta (SNpc) and the resulting deficit in dopaminergic (DA) innervation in the striatum and other areas appear to be responsible for the majority of the characteristic motor and cognitive dysfunction of PD<sup>16-18</sup>.

Significant motor improvements result from systemic administration of the dopamine precursor, L-Dopa, or dopamine agonists<sup>19</sup>, supporting the rationale that dopamine release provided by transplanted replacement grafts might provide therapeutic effects. Supporting evidence from numerous studies in rodents<sup>3,20-23</sup> and non-human primates<sup>24-31</sup> have shown that grafts of fetal DA neurons can lead to biochemical and functional behavioral improvements<sup>31-36</sup>. However, in clinical studies, the improvements in parkinsonism have been rather modest and variable<sup>37-49</sup>.

GDNF is essential for nervous system development and survival of ventral midbrain dopaminergic (DA) neurons in-vivo<sup>50-52</sup> and has been shown to act as a potent chemo-attractant in stem cell migration and neuritic maturation in the brain<sup>53-56</sup>. Exogenous delivery of GDNF by numerous modalities including, protein pump, cell secretion, and viral vectors<sup>52,57-71</sup> have yielded promising functional improvement in several PD models<sup>72-85</sup> at the

pre-clinical level, but present a more complicated picture in the clinic thus far<sup>86</sup>.

To further aid in these attempts, we and others have shown that exogenous overexpression of AAV-driven GDNF in the host striatum can increase the survival of ectopically placed fetal VM grafts by several fold as well as elicit highly directional axonal outgrowth<sup>87,88</sup>. In addition, we demonstrated by fluorogold tract tracing, that VM grafts placed in the substantia nigra (SN) harbor the capacity to project TH-ir neuritic processes across substantial distances and innervate the host striatum when exposed to exogenous GDNF<sup>89</sup>. Furthermore, dopamine neuroblasts implanted into the SN of 6-Hydroxydopamine (6-OHDA) lesioned mice are capable of regenerating a new anatomically aligned, functionally relevant nigrostriatal pathway<sup>90</sup>. This target directed growth was further enhanced by striatal GDNF overexpression and supports the rationale for cellular transplantation and gene co-therapy as a dual modality therapeutic approach for nigrostriatal (NS) reconstruction in animal models of PD.

Until recently<sup>89,90</sup>, attempts to establish new anatomically appropriate nigrostriatal axonal projections from fetal DA neurons placed within the VM had been met with relatively moderate success<sup>91-95</sup>. As a result, exogenous growth factors, such as GDNF, were utilized to help overcome what was suspected to be a relatively non-permissive growth environment<sup>84,96,97</sup>. In addition, studies utilizing immature neuroblasts or early post-mitotic neurons transplanted into other adult central nervous system locations (homotopic and heterotopic) *clearly demonstrate* the capacity to extend long distance, target specific axonal projections<sup>21,23,26,98-119</sup>. While these results are extremely promising, it is still relatively unknown whether undifferentiated hNPC also retain the capacity to differentiate<sup>120</sup> and respond to relevant signaling cues in the adult dopamine depleted primate brain.

While clinical trials have demonstrated *promising* results<sup>37-42,45,46,48,49,121-133</sup>, acquiring sufficient high-quality donor tissue remains a definitive obstacle for fetal VM cell

transplantation. To circumvent this issue, undifferentiated hNPC can be readily expanded in-vitro to provide sufficient cellular substrate for multiple patients from a single donor source. These cells retain an inherent developmental plasticity and respond to local excitatory and inhibitory axon guidance molecules<sup>134</sup>, differentiating in relation to the changing microenvironment in-vivo.

Previously, we demonstrated that human fetal forebrain neural stem/progenitor cells (hfNPC) engraft, migrate, and promote functional improvements in behavioral deficits when transplanted into the MPTP-lesioned, African green monkey<sup>11,135,136</sup>. Graft analysis revealed few donor-derived DA cells (BrdU+/TH+), suggesting that the reduction in parkinsonian behaviors and subsequent normalization of endogenous TH-ir cell numbers/area were most likely a result of indirect, secondary support mechanisms. Specifically, we hypothesized the homeostatic effect to be a result of alternative mechanisms: release of neuroprotective growth factors, astrocytic differentiation, and recruitment of inflammatory cytokine signaling molecules.

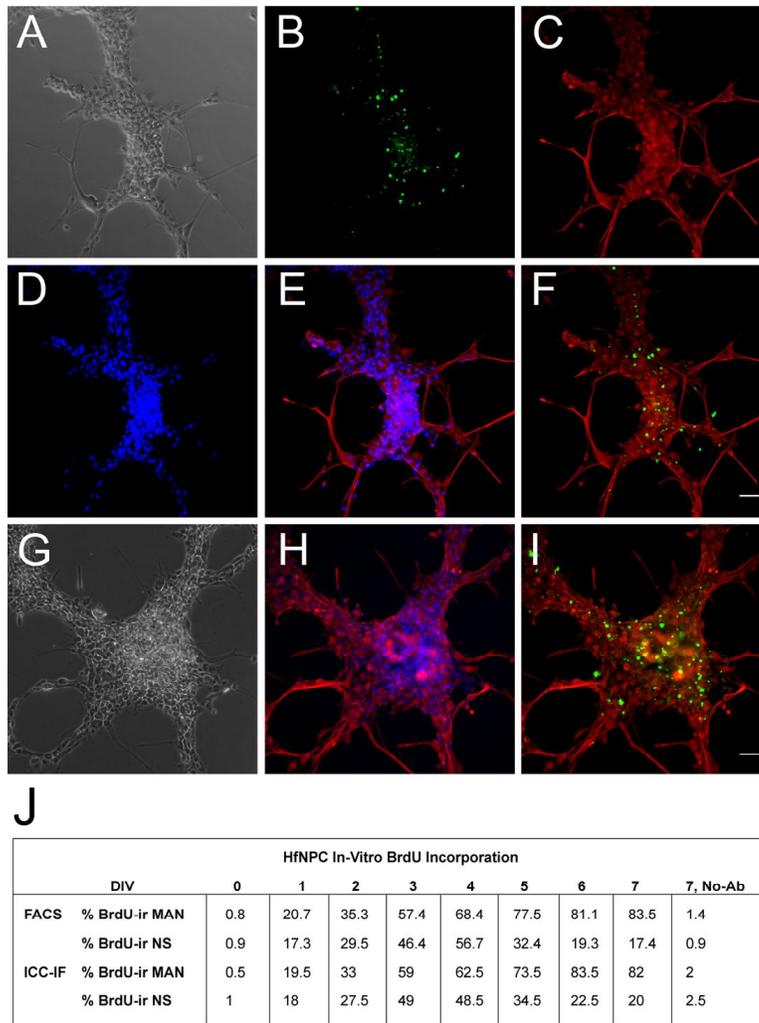
Given strong evidence for a permissive nigrostriatal growth environment to newly generated neuritic processes in rodents<sup>90,101,102</sup> and primates<sup>89</sup>, it is feasible that hNPC might also respond to similar trophic stimuli in-vivo. Immature fetal VM DA neuroblasts and other post-mitotic VM cell types have been shown to respond to environmental stimuli<sup>137,138</sup> and develop into mature DA neurons in-vitro<sup>139,140</sup> and in vivo<sup>105,106,141</sup>, however, it is still relatively unclear whether undifferentiated SVZ-derived hfNPC retain the intrinsic capacity to significantly differentiate into mature disease relevant A9-DA neurons and innervate appropriate target regions in the adult degenerative brain. Therefore, we sought to determine whether overexpression of AAV5-GDNF, delivered into the primate striatum could enhance graft integration and DA differentiation of donor hfNSC injected into the SN, as well as elicit directional neuritic outgrowth from the SN to the caudate nucleus. After eleven months,

healthy donor-derived cells with multiple morphologically relevant neuronal phenotypes successfully engrafted and extended axonal projections parallel to endogenous fiber tracts, indicating that hfNPC retain the capacity to respond to endogenous signaling molecules and differentiate in accordance with a permissive growth environment in the adult dopamine-depleted primate brain.

### **4.3 Results**

#### **4.3.1 Pre-Labeling hfNSC for Transplantation**

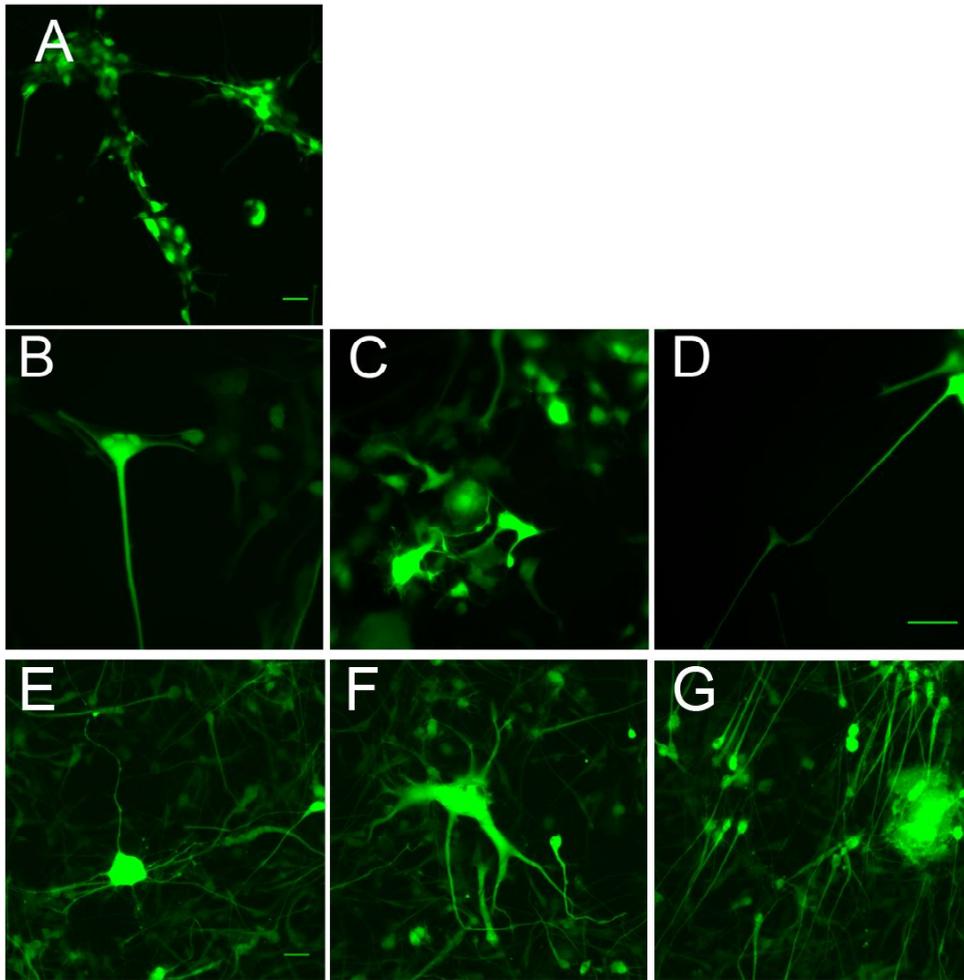
To our knowledge, no single marker exists to adequately distinguish human from non-human primate cells in the CNS; therefore, donor hNPC require pre-labeling in-vitro to ensure accurate graft identification post-mortem. While harsh DNA-intercalating thymidine analogs like BrdU have traditionally been used to label proliferative donor hNSC, we found that exposure beyond 4-DIV was toxic and resulted in only  $\approx 48-57\%$  labeling efficiency when grown as suspension aggregates. In contrast, when hNPC were grown as multilayer adherent networks (MAN)<sup>142</sup>, cell density appeared to overcome cellular toxicity up to 7-DIV, allowing for a larger percentage ( $\approx 83\%$ ) of cells to pass through the cell-cycle and acquire a BrdU label [Figure 4-1].



**Figure 4-1. HfNPC readily incorporate BrdU and are efficiently transduced with eGFP when cultured as multilayer adherent networks (MAN).**

HFB-2050 hfNPC were cultured utilizing the MAN technique or as aggregate “neurospheres” in the presence of BrdU for up to seven days in-vitro (DIV) and analyzed by immunocytochemistry (ICC) with both FACS and immunofluorescence (IF) to determine labeling efficiency. MAN culturing conditions improved labeling efficiency and allowed for incubation without toxicity beyond 4DIV. HFB-2050 hfNPC grown as MAN cultures were labeled for Nestin (red) and BrdU (green) at 4-DIV (A-F) and 7-DIV (G-I). Proliferative BrdU-ir cells were typically located within the center of the three-dimensional cellular clusters, opposed to the highly arborized webbed end-feet protruding from and connecting each larger cluster. [J], The efficiency of BrdU incorporation was determined for hfNPC grown as either neurospheres (NS) or multilayer adherent networks (MAN) for up to 7 days in-vitro (DIV) and analyzed by fluorescence microscopy (ICC-IF) or fluorescent activated cell sorting (FACS). Cells grown for 7-DIV in the presence of BrdU but not stained with fluorescent antibody are shown as a control. Data for 7-DIV without BrdU but stained with antibody were similar (not shown).

In previous studies, nigrostriatal tract tracing and donor-derived neuritic profiles could not be assessed due to histological limitations with BrdU and coronal sectioning. Generation of a stably-integrated fluorescent hNPC line eliminates the need for these toxic compounds<sup>143</sup>, eliminates false-positive artifacts<sup>144</sup>, while increasing both cell viability and graft detection efficiency. Therefore, in addition to the nuclear BrdU tracer, donor hfNPC were also pre-labeled with a stable, constitutively active, cytosolic, eGFP lentivirus [HFB-2050-eGFP] to capture donor cell morphology and accurately trace neuritic branching and cellular differentiation profiles post-mortem [Figure 4-2]. After two months of expansion, stable polyclonal populations were analyzed by flow cytometry (average, 90.5% eGFP+ n=5) and fluorescence microscopy (average, 87.5% eGFP+, n=3) to ensure efficient stable transgene expression [Figure 4-2 A-D]. Upon addition of 10% fetal bovine serum for seven months, HFB-2050-eGFP became extremely adherent and elongated into a morphologically differentiated profile with cells resembling astrocytes and neurons similar to non-transduced counterparts [Figure 4-2 E-G].

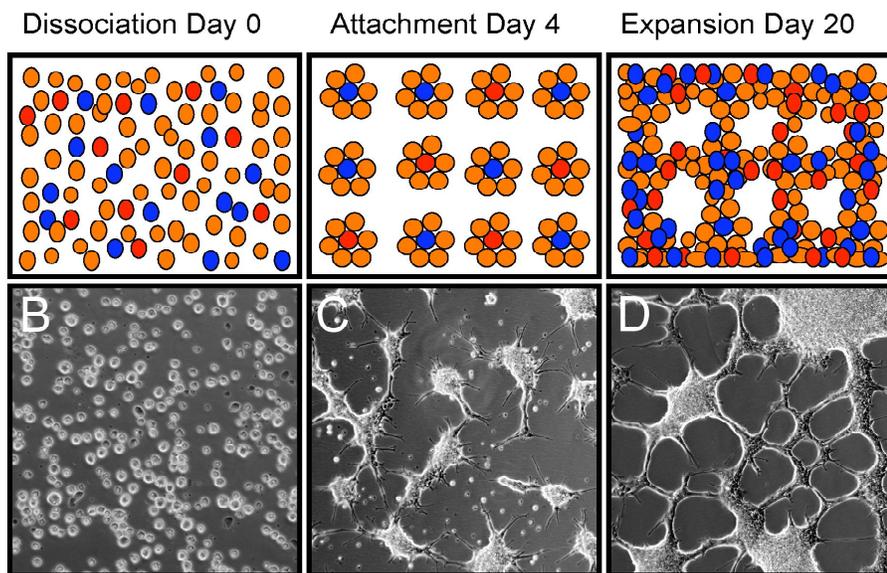
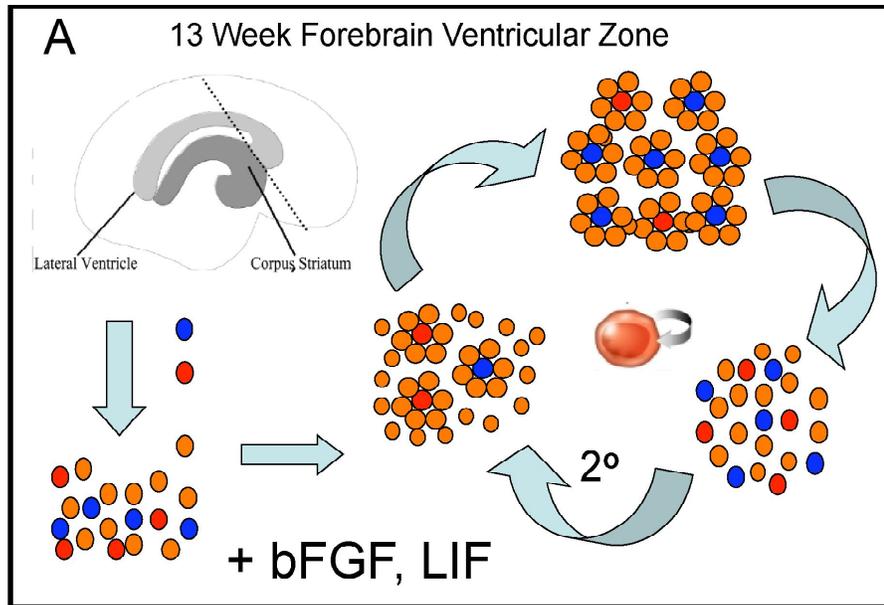


**Figure 4-2. Undifferentiated hfNPC were stably transduced with a cytosolic-eGFP lentivirus and expanded for several months in-vitro.** While often found in multi-cellular clusters [A], examination of individual migratory cells at low density typically revealed a bifurcated leading edge growth cone adorned with meandering micro-cilia [B-D]. Cells appear to constantly survey and sample the local microenvironment, protruding and retracting, often at great distances, until they come into contact with an established “colony,” whereby they ascend and migrate toward the leading edge, eventually joining into aggregate units. In addition, HFB-2050-GFP assumed multiple differentiated cell morphologies after addition of 10%FBS for seven months (E-G) confirming that lentiviral transduction did not affect multipotency.

#### 4.3.2 Multilayer Adherent Network hfNPC Culturing System

Adherent culturing conditions<sup>142,145-152</sup> have several advantageous characteristics compared to their suspension “neurosphere” counterparts and have gained attention for their utility in high-throughput applications like chemical screening<sup>153</sup>. In order to produce

sufficient cellular substrate for multiple transplants, we developed a novel culturing method for expanding hfNPC as multilayer adherent networks (MAN) [Diagram 4-1], with enhanced growth parameters compared to aggregated suspension spheres<sup>142</sup>. Utilizing these conditions, primary hfNPC, HFB-2050, were maintained for over 120 population doublings without significant chromosomal aberration or apparent senescence as previously reported utilizing suspension aggregate methods<sup>154,155</sup>. MAN cultures demonstrate an increased population doubling time as well as a greater percentage of actively dividing cells [Figure 4-1 J], while retaining a gene expression profile signature (mRNA and miRNA) similar to hNPC derived from several other central nervous system tissue sources. When compared to human embryonic stem cells (hESCs) and neural stem cells derived using alternative in-vitro manipulation techniques, HFB-2050 hfNPC mRNA and miRNA gene expression profiles clustered most closely within other human fetal CNS NPC preparations<sup>156,157</sup>. In fact, they were extremely similar to a primary human fetal telencephalon NPC line derived utilizing identical techniques, termed HFT-13<sup>158</sup>, that were recently shown to be fully reprogrammable to induced pluripotent stem (iPS) cells utilizing only the transcription factor Oct-4<sup>159</sup>.



**Diagram 4-1. Derivation and multilayer-adherent-network (MAN) expansion of hfNPC, HFB-2050.** Human fetal forebrain periventricular-zone from a 13-wk cadaver was roughly dissected and dissociated in the presence of mitogens. After growth-factor selection for proliferative stem-like cells, primary cultures were further cultured in bFGF and LIF [A]. Expansion of hfNPC was enhanced utilizing the MAN culturing method [B-D]. Dissociated cells are re-plated at a greater density [B] compared to “neurosphere” suspension culture techniques and allowed to grow for 3-4 days without disruption inducing attachment of small aggregate clusters [C] that eventually proliferate and expand into densely organized adherent 3D-networks [D].

### 4.3.3 HfNPC HFB-2050 Gene Expression

QRT-PCR confirmed a neural stem/progenitor cell lineage identity for HFB-2050 when compared to undifferentiated WA-H9 hESCs. HFB-2050 had a marked increase in expression of the NSC associated SoxB transcription factors, *sox1* and *sox2*<sup>160,161</sup>, as well as similar levels of *Myc* and significantly less *Klf4* compared to H9 hESC [Table 4-1]. These genes are essential in maintaining pluripotency in hESC as well, acting in a regulatory network with *Oct-4*, *Nanog* and several other factors<sup>162-174</sup>. In agreement, HFB-2050 displayed significantly lower mRNA levels for hESC regulatory genes *Oct-3/4* (*Pou5F1*), *Nanog*, and *Lin28*, as well as similar expression of the stem cell associated intermediate filament *Nestin*, excluding the possibility that these cells were partially undifferentiated pluripotent progenitors [Table 4-1]. Furthermore, compared to hESC, HFB-2050 hfNPC expressed moderately elevated levels of *Pax6*, an essential regulator of NSC self-renewal, neurogenesis, and fate specification in vivo<sup>175-178</sup>, as well as low levels of *Ptx3*, an important factor in DA maturation in the mouse brain. *Pax6* and *SoxB* family members have been shown to regulate the transition from neuroepithelial precursors to radial glia in-vivo and in hESC derived cultures in-vitro<sup>179-181</sup>, while *Ptx3* appears to play a role in downstream signaling events during DA differentiation. Moreover, undifferentiated HFB-2050 hfNPC retained little to no gene expression for multiple pro-neurogenic genes, specifically those associated with signaling and maturation of the DA lineage including *Wnt1*, *Lmx1a*, *Lmx1b*, *EN1*, *FoxA2*, *Nurr1* (*NR4A2*), and *TH*<sup>182-191</sup> [Table 4-1]. These combined expression data strongly support HFB-2050 lineage as a bona-fide fetal neuroectodermal hNPC population.

**Table 4-1. Quantitative PCR profile of HFB-2050 hfNPC and hESC-WAH9 for known regulators of pluripotency and neural differentiation.** Gene expression levels are consistent with a primitive neuroectodermal fate (SoxB family members, Nestin, pax6) and argue against a pluripotent ESC-like (Oct-4, Nanog) or neuronally restricted (Lmx1, EN1, TH, etc.) progenitor. NS = Not Significant; - denotes a significant decrease in expression; + denotes a significant increase in expression; NE = Not Expressed

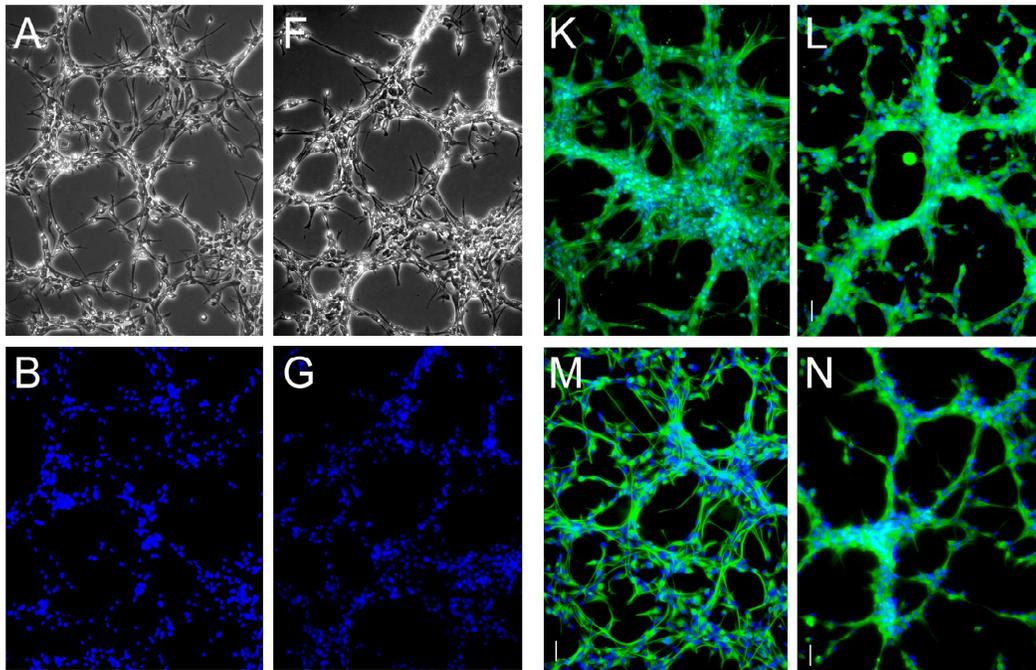
GENE	WA-H9 $\Delta$ CT	hfNPC $\Delta$ CT	$\Delta$
NANOG	4	11.5	-
POU5F1	0.4	8.2	-
LIN28	1.1	14.4	-
KLF4	9.4	13.4	-
MYC	7.6	6.3	NS
SOX1	10.6	5.8	+
SOX2	4.4	2.3	NS
NES	6.5	5.8	NS
PAX6	11.9	9.8	NS
WNT1	12	17.9	-
FOXA2	11.1	NE	-
EN1	15.1	19	-
LMX1A	13.9	20.4	-
LMX1B	13.6	19.3	-
PTX3	17	13.1	+
NR4A2	12.4	13.1	NS
TH	15.4	14.4	NS

$\Delta$ CT= (CTgene of interest) –(CTGAPDH);  
A lower  $\Delta$ CT denotes higher gene expression

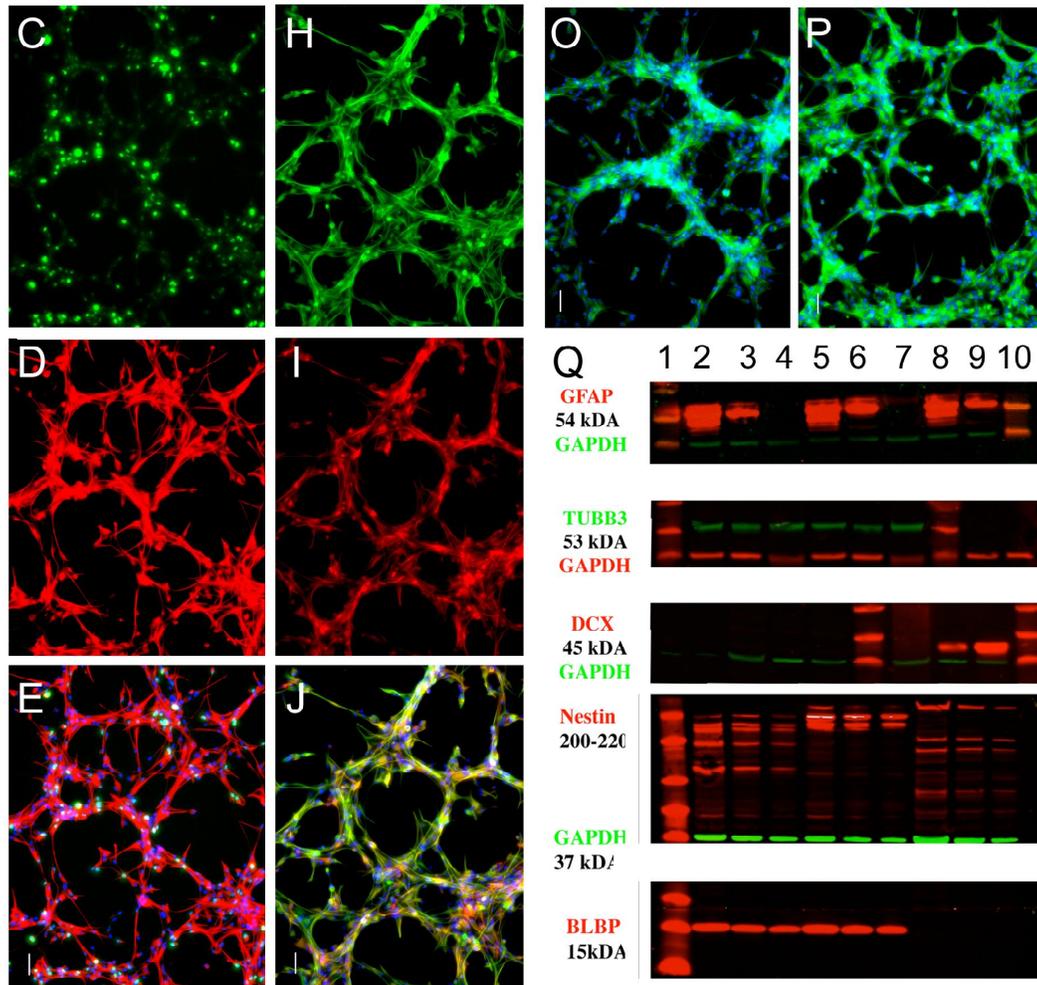
#### 4.3.4 HfNSC HFB-2050 Express Neurogenic Proteins:

We next examined expression of NSC-niche related proteins in-vitro by ICC-IF to characterize their morphological distribution in hfNPC. Highly proliferative (Ki-67-ir) MAN cultures of HFB-2050 hfNPC [Figure 4-3 C] displayed robust cytoplasmic co-expression of the NSC-lineage related filamentous proteins Nestin [Figure 4-3 D, H] and GFAP [Figure 4-3 I] as well as transcription factors sox2 [Figure 4-3 K] and sox-3 [Figure 4-3 L] and membrane

bound Tie-2 [Figure 4-3 N]. Surprisingly, hfNPC also markedly expressed the “immature” neuronal markers, cytoplasmic filament beta-III-tubulin/Tuj1<sup>192,193</sup> [Figure 4-3 M] and extracellular polysialic-neural cell adhesion molecule (PSA-NCAM) [Figure 4-3 O] attributed to migratory type-A neural progenitors<sup>194-202</sup>. In addition, HFB-2050 hfNPC express cytoplasmic brain lipid binding protein (BLBP) [Figure 4-3 P], a marker for radial glia<sup>203,204</sup>, the NE derived NSC of the fetal brain<sup>205-213</sup> suggesting HFB-2050 express developmentally appropriate forebrain hfNPC lineage markers<sup>194,204,214-229</sup>



**Figure 4-3. Human fetal NPC express proteins characteristic of stem/precursor cell in-vitro.** Proliferative, Ki-67-ir HFB-2050 cultures [A-E] expanded and maintained as multilayer adherent networks (MAN) expressed the neural stem cell related filaments Nestin, [A-J] and GFAP [F-J], as well as the NSC transcriptional regulators sox2 [K] and sox3 [L] and membrane associated Tie-2 [N]. Of interest, other neurogenesis related proteins such as Tuj-1 [M] and PSA-NCAM [O] (markers for immature neuroblasts in-vivo), along with BLBP [P] (marker for radial glia) were also highly expressed in-vitro. (A,F = phase contrast, B, G = DAPI, G = Ki-67, H, D = Nestin, I = GFAP, E, J = Merged images. K-P are merged with DAPI. Western blotting confirmed expression of neurogenic proteins in undifferentiated hfNPC [Q]. When compared to “neurosphere” (NS) suspension cultures, MAN cultures express increased levels of high molecular weight Nestin, multiple isoforms of GFAP, and similar levels of BLBP and Beta-3-tubulin/Tuj1, while NS cultures had a marked increase in DCX protein levels. N2A neuroblastoma cells were used as a +/- controls where applicable. Lanes labeled 1-10, Running order as follows. GFAP, ladder = 1, 10; 2,5,8 = MAN, 3,6,9 = NS; 4,7 = N2A neuroblastoma negative control. TUBB3/TuJ1/Beta-3-Tubulin: ladder = 1,8; MAN = 2,5,9; NS = 3,6,10; N2A = 3,6 as a positive control. Lanes 9, 10 probed with antibody from Chemicon vs. Covance in lanes 1-7 demonstrating specificity to only the Covance antibody. DCX: ladder = 6,10; Lanes 1,2 = BSA – loading control, lanes 3-5 stained without primary DCX antibody as a control for background from secondary antibody alone. Ladder = 6, 10; MAN = 4, 8; NS = 5,9 N2A = 3,7 as a negative control. Nestin and BLBP: Ladder = lane 1; NS = lanes 2-4; MAN = 5-7; N2A = 8-10. Nestin and BLBP lanes represent serial 1/10 dilutions (100X, 10X, 1X)



**Figure 4-3. Continued**

Biochemical analysis of hfNPC grown as either MAN or suspension “neurospheres” confirmed the expression of NSC niche associated proteins. Western blotting revealed elevated expression of high molecular weight, human-specific Nestin as well as multiple isoforms of GFAP<sup>224</sup>, when grown under MAN conditions [Figure 4-3 Q]. In addition, BLBP and Tuj1/beta-3-tubulin were expressed at similar levels, while the migratory neuroblast associated protein doublecortin (DCX)<sup>230-232</sup> was elevated under suspension “neurosphere” conditions [Figure 4-3 Q]. The combined results suggest that hfNPC exist as proliferative,

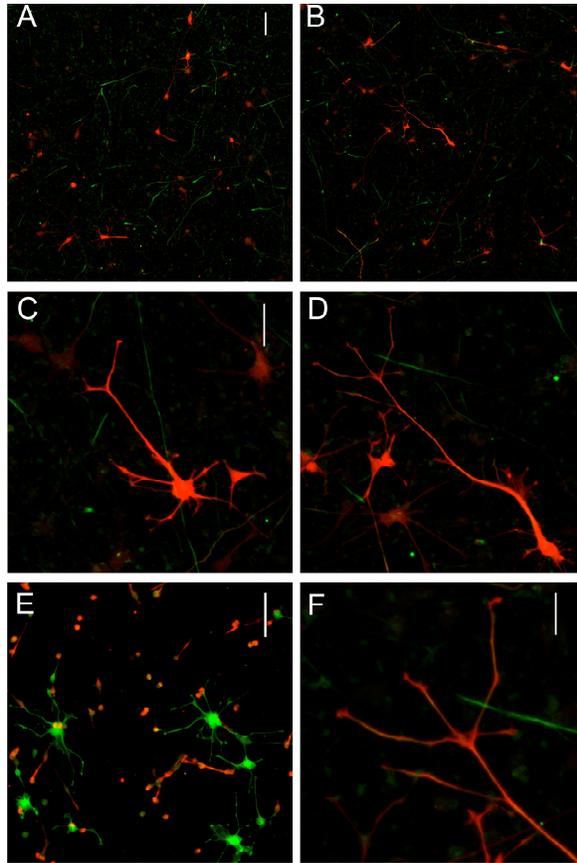
highly plastic stem/precursor cells, “primed” for neurogenesis. In addition, biochemical analysis indicates that in-vitro culturing conditions dramatically alter NSC lineage related protein expression, and neurosphere growth parameters may actually artificially select for slightly differentiated neuroblast like neural precursor cells (NPC). These observations may help explain previous reports of early senescence and loss of neurogenic potential in hfNPC (<60 passages) utilizing suspension aggregate culturing methods<sup>154,155,233</sup>.

#### 4.3.5 HFB-2050 hfNSC Recapitulate the Subventricular Zone Niche

To further investigate the developmental plasticity of hfNPC, HFB-2050-WT cells were cultured in an extracellular matrix (ECM) protein rich environment. Matrigel contains laminin, collagen-IV, heparin sulfate proteoglycans, and entactin, as well as trace amounts of platelet-derived growth factor (PDGF), nerve growth factor (NGF), insulin-like growth factor-1 (IGF-1), and TGF- $\beta$ ; proteins with functional roles in maintenance of the NSC niche, as well as differentiation and migration during normal CNS development<sup>145,234-244</sup>. We therefore sought to emulate the early neurogenic microenvironment, utilizing Matrigel to model early neurogenesis and induce differentiation in-vitro.

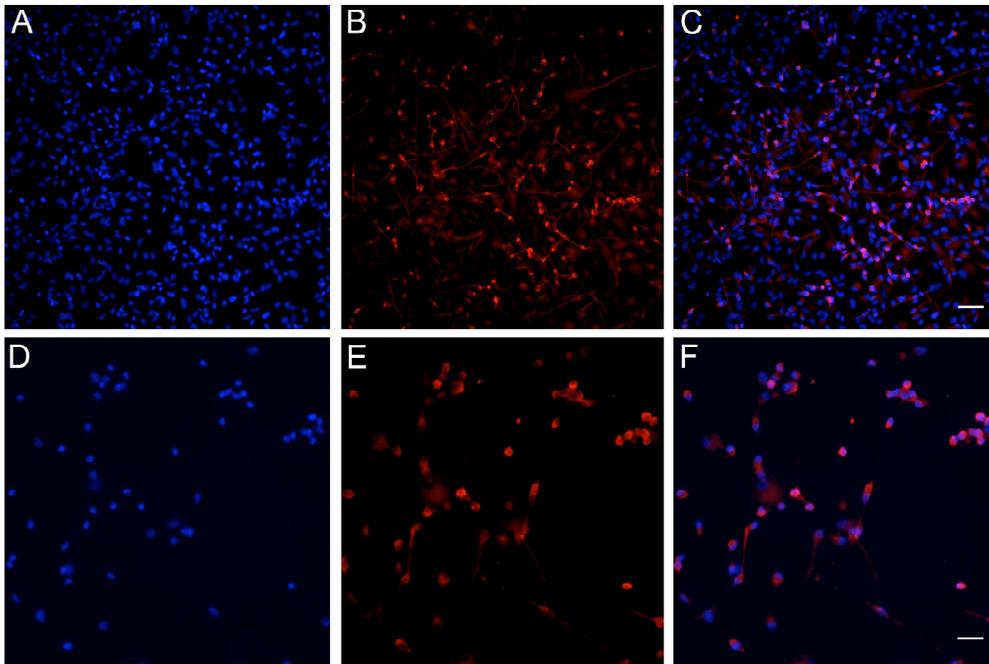
HfNPC were cultured on Matrigel to induce intermediate cell phenotypes and analyzed by ICC-IF for lineage specific NSC niche proteins. Immunostaining revealed a phenotypically *heterogeneous* population of precursor cells comprised of at least two specific replicating sub-populations of cells; supporting radial glial like (BLBP<sup>high</sup>/DCX-) and migratory (BLBP<sup>low</sup>/DCX+) precursor cells [Figure 4-4]. BLBP<sup>high</sup>/DCX- cells phenotypically resemble type-B astrocyte like NSCs of the adult subventricular zone (SVZ) niche<sup>194,207,208,214-217,219</sup>. Many BLBP<sup>low</sup>/DCX+ cells displayed distinct  $\beta$ -3-tubulin+(Tuj-1) leading and lagging processes [Figure 4-5], highly reminiscent of the in-vivo neuroblast found in the murine rostral migratory stream. These cells appear to migrate along the BLBP<sup>high</sup>/DCX- radial glial

scaffolding forming a multifaceted, in-vitro *microniche* resembling the SVZ neurogenic niche, in-vivo<sup>207,216,245</sup>. Overall, these data suggest that HFB-2050 populations retain characteristic properties of multipotent forebrain hfNSC, in-vitro<sup>246</sup>. Specifically, when introduced to ECM instructive cues, they readily differentiate into early precursor phenotypes and maintain critical components reminiscent of the ventricular zone neurogenic niche.



**Figure 4-4. Modeling early subventricular-zone (SVZ) neurogenesis in-vitro.**

HFB-2050 hfNPC were cultured on Matrigel to induce developmental programs involved in early neurogenesis. Stem/precursor cell phenotype were analyzed by immunocytochemistry revealing a relatively heterogeneous population comprised of two morphologically distinct phenotypes. Nearly all cells expressed some degree of Nestin, Vimentin, and GFAP (not shown); however, there were marked differences in structural characteristics for cells that highly expressed either BLBP (green) or DCX (red) independently [E]. Morphologically,  $BLBP^{high}/DCX^{-}$  cells were characterized by a large stellate cell body and nuclei [A-D, red = BLBP, green = NFM], as well as extensive outgrowth of multipolar arborizations [C, D], often with one or two elongated extensions protruding up to several mm in length, resembling radial glia or astrocyte-like type-B cells in-vivo [D, F]. The branched processes appear to function as neural scaffolding or “highways” upon which the  $BLBP^{low}/DCX^{+}$  sub-population migrates [E]. In contrast,  $BLBP^{low}/DCX^{+}$  cells [E] are distinguished by their highly migratory behavior, smaller cell body and nuclei, bipolar morphology, and leading edge protrusion, resembling both type-C transit amplifying cells and type-A neuroblasts. After 3DIV [E], cells are still beginning to develop in accordance with ECM cues (red= DCX, green = BLBP). By 7DIV [A-D, F], cultures mature significantly revealing extensive arborization and outgrowth from BLBP-ir (red) radial-glia like cells as well as the emergence of neuronal-related NFM-ir (green) fibers.



**Figure 4-5. Extracellular matrix induction of immature-neuron associated Beta-3-tubulin/Tuj-1.** Undifferentiated cells cultured on Matrigel extracellular matrix proteins expressed Tuj1 in immature bipolar, migratory cells after 3-DIV [D-F]. Development of more mature Tuj1-ir neuronal morphologies were seen after 7-DIV [A-C] indicating HFB-2050 hfNPC retain developmentally appropriate programs relevant to early neurogenesis in-vitro.

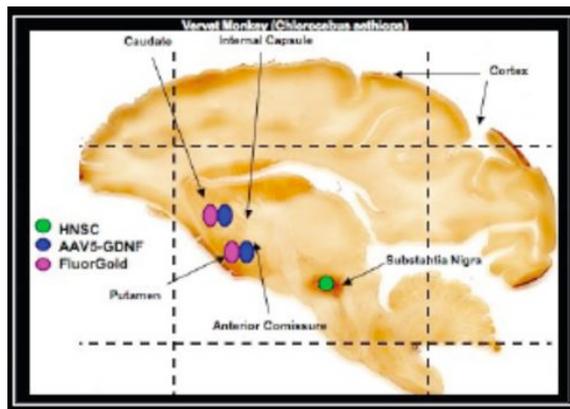
#### 4.3.6 HfNSC HFB-2050 Are Multipotent In-Vitro and In-Vivo

Previously, we demonstrated that undifferentiated HFB-2050 hfNPC retain the capacity to express proteins associated with the VM lineage in-vitro, expressing Nurr1, Ptx3, and TH<sup>11</sup>, when cultured in the presence of glial conditioned media and growth factors<sup>247-249</sup>. In addition, we have shown that these cells can be selectively directed into oligodendrocytes in-vitro<sup>250</sup>. Furthermore, upon transplantation into the lateral ventricles of P0 neonatal mice, HFB-2050-eGFP hfNPC participate in normal CNS development, including migration from germinal zones through the RMS to olfactory bulbs, into the corpus callosum, and along vascular networks throughout the brain<sup>142</sup>. These data suggest HFB-2050 hfNPC retain a responsiveness to regional and temporal developmental cues to become multiple cell types in these regions as well as integrating long-term and non-disruptively within the host SVZ niche.

Collectively, our data support HFB-2050 hfNPC as long-term self-renewing, undifferentiated, multipotent (in-vitro and in-vivo) progenitors suitable for in-vivo transplantation.

#### 4.3.7 Nigral HfNPC Engraft Long-Term Concomitantly with Striatal AAV5-GDNF:

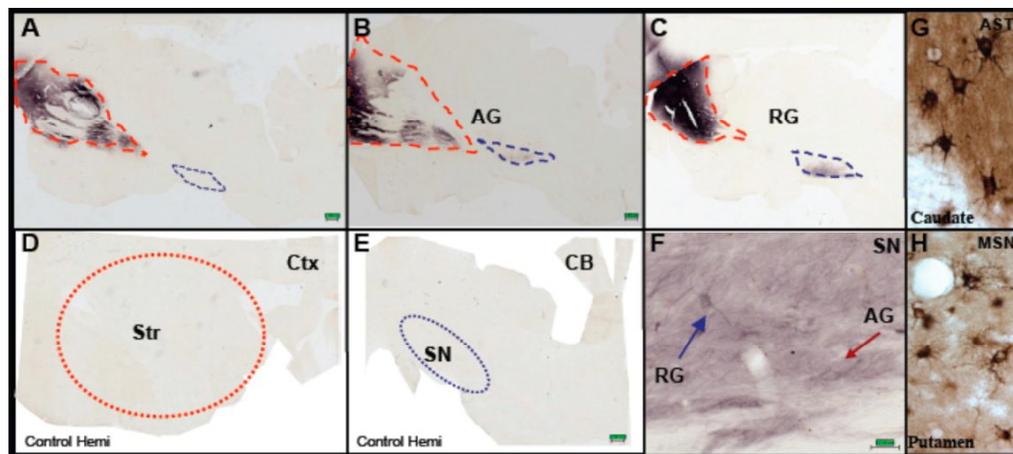
To determine whether undifferentiated hfNPC grafts retain the capacity to differentiate and respond to GDNF chemotaxis in-vivo, ten MPTP-treated adult male St. Kitts green monkeys received injections of AAV-5 GDNF unilaterally into both the rostral caudate and post-commissural putamen, followed by a hfNPC graft unilaterally into the ipsilateral hemisphere, immediately rostral to the SN [Diagram 4-2; Supplementary Table 4-1]. Animals were sacrificed at 1.5 months or 11 months and analyzed for GDNF expression and hfNPC-derived graft distribution. Parasagittal sections were utilized in order to capture the complex neuronal circuitry of the entire NS unit.



**Diagram 4-2. Schematic representation of injection targets in veret brain.** HfNSC (green oval) were homotopically transplanted into the rostral SN, and AAV5-GDNF (blue oval) was administered into both the caudate and putamen. After 11-months, Fluorogold (Purple oval) was injected into the identical striatal coordinates previously used for AAV5-GDNF. The dotted lines demarcates where the tissue was cut for histological analysis.

#### 4.3.7 AAV5-GDNF Distribution

Extensive long-term overexpression of AAV-5-GDNF at 1.5 months [Supplementary Figure 1] and 11 months [Figure 4-6] was confirmed unilaterally throughout the injected striatum, emanating and diffusing outward from the injection sites throughout both the caudate and putamen. GDNF-ir cells were distributed from the most dorsal-rostral regions of the far-lateral extent of the striatum extending ventrally throughout the caudate and into the most caudal regions of the putamen [Figure 4-5 A,B,C]. Transduced host striatal cells highly expressing GDNF had the characteristic morphology and molecular profile of predominantly medium spiny neurons and to a lesser extent, stellate astrocytes [Figure 4-6 G,H]. Immunostaining for GDNF, NeuN and GFAP revealed the majority of transduced cells had a NeuN-ir neuronal phenotype and morphology consistent with medial spiny neurons throughout most of the striatum and a small percentage of astrocytes in the most dorsal-rostral regions of the far-lateral extent of the caudate (data not shown).



**Figure 4-6. Exogenous AAV5-GDNF expression after 11-months.** Extensive diffusion and expression of GDNF (black staining) from AAV-serotype-5 was verified throughout the striatum (striatum = red dotted lines) [A, B], emanating from the injection sites [C] into adjacent tissue. Exogenous GDNF was not found in contralateral hemispheres [D, E]; however GDNF-ir cells bodies and fibers were also seen within the ipsilateral SN (SN = blue dashed lines), suggesting retrograde (RG, blue arrow) and anterograde transport (AG, red arrow) [F], respectively, from striatal injection sites. [G, H] Most GDNF-transduced cells (brown staining) had the characteristic morphology of striatal medium spiny neurons throughout the striatum as well as astrocytes in some regions of the caudate.

GDNF-ir was also abundant within the ipsilateral SN of the injected hemisphere [Figure 4-6 F] providing evidence for transport of exogenous GDNF from the striatum to the SN. Specifically nigral neuronal soma with DAergic morphology and processes located within the nigra were densely immunoreactive, suggesting either retrograde transport from nigrostriatal terminals or anterograde transport downstream along the striatonigral pathway to relevant targets including the SN. In addition, exogenous AAV-GDNF was never seen in the contralateral hemisphere within the striatum or SN [Figure 4-6 D,E], indicating localized delivery. Furthermore, no adverse behaviors or changes in feeding or body mass previously reported using AAV2 serotypes<sup>58</sup> were observed at any point throughout the experiments.

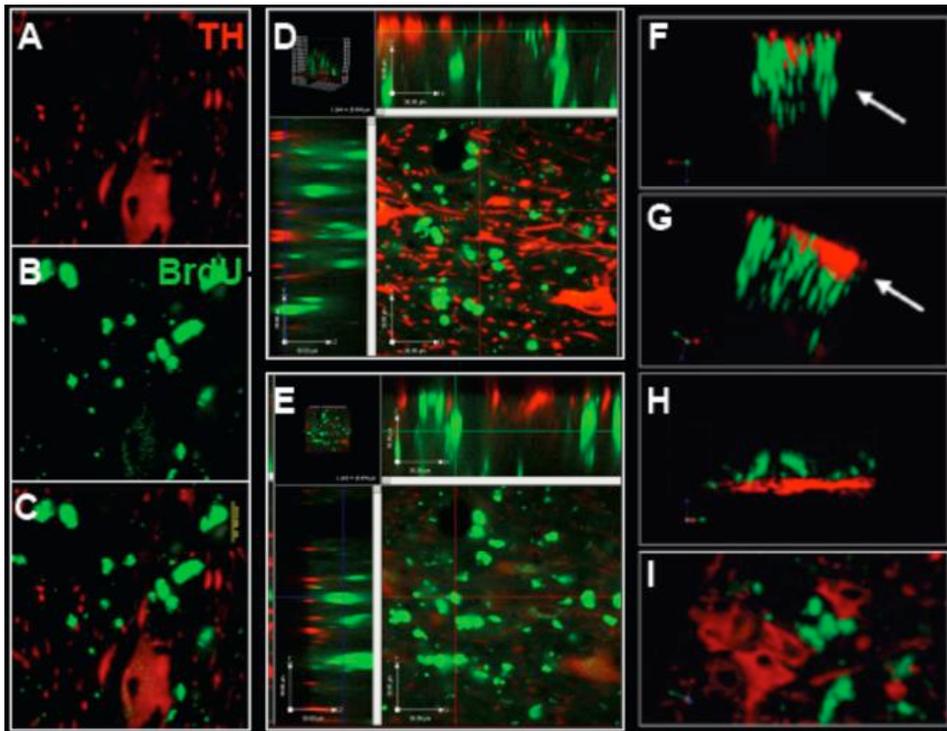
#### **4.3.9 Endogenous TH-IR within the Nigrostriatum:**

Analysis of TH-ir cell bodies located within the SN and their axonal projections to the striatum revealed a characteristic reduction in the number of endogenous TH-ir cell bodies (asymptomatic monkey) compared to control non-MPTP injected animals (data not shown). Interestingly, by 11 months, host TH-ir cells appeared to expand and fill the subsequently degenerated interstitial spaces with extensively arborized neuritic processes, indicative of endogenous regeneration. In addition, a moderate level of intact host TH-ir nigrostriatal fibers<sup>251</sup> were spared or rescued from degeneration. Host outgrowth was likely a result of predominantly GDNF neurotrophic support within the SN, as well, to a lesser extent, hfNPC graft derived factors in concert with local signaling cues.

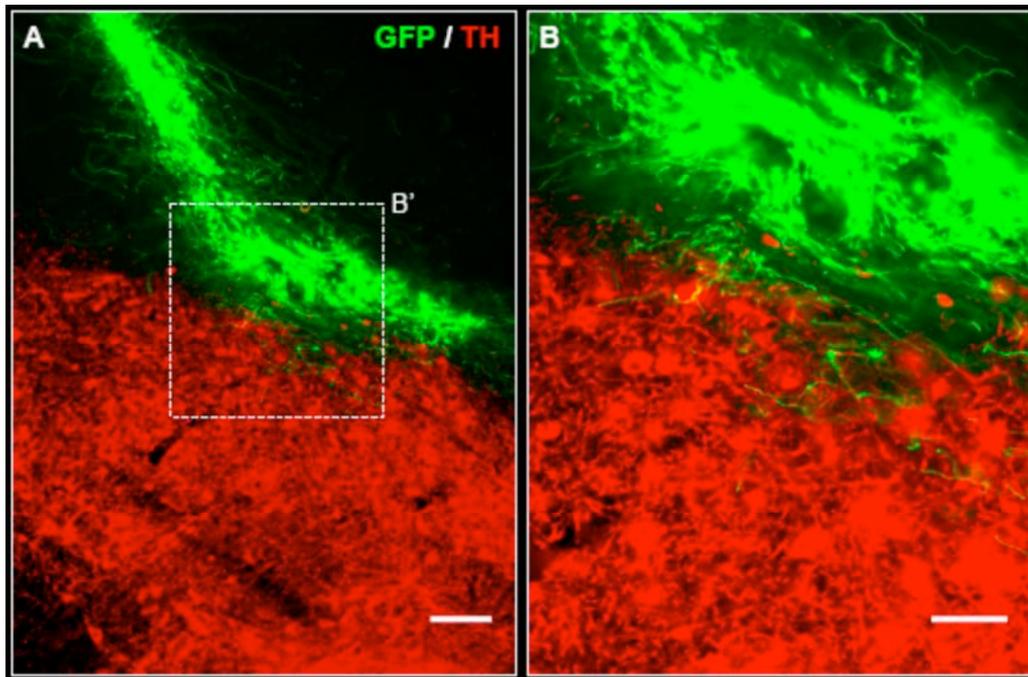
#### **4.3.10 HfNPC Graft Distribution and Morphology:**

Robust donor-derived grafts were typically located immediately dorsal and adjacent to the most dorsal-rostral region of the host SN, with some variability toward more dorsal-caudal extents of the SN in one animal. After 1.5 months, donor cells generally remained within the

target site, and did not migrate to the hemisphere contralateral from the injection site in any animals [Figure 4-7 A-C]. Grafted BrdU-ir cells were typically clustered within and immediately adjacent to TH-ir dense regions of the SN, located in an adjacent plane to endogenous TH-ir cell bodies [Figure 4-7 D-I, Supplementary Video 1] and directly juxtaposed to some endogenous TH-ir neurite tracts. After 1.5 months, donor cells retained a relatively undifferentiated phenotype, typically with little to no arborization.



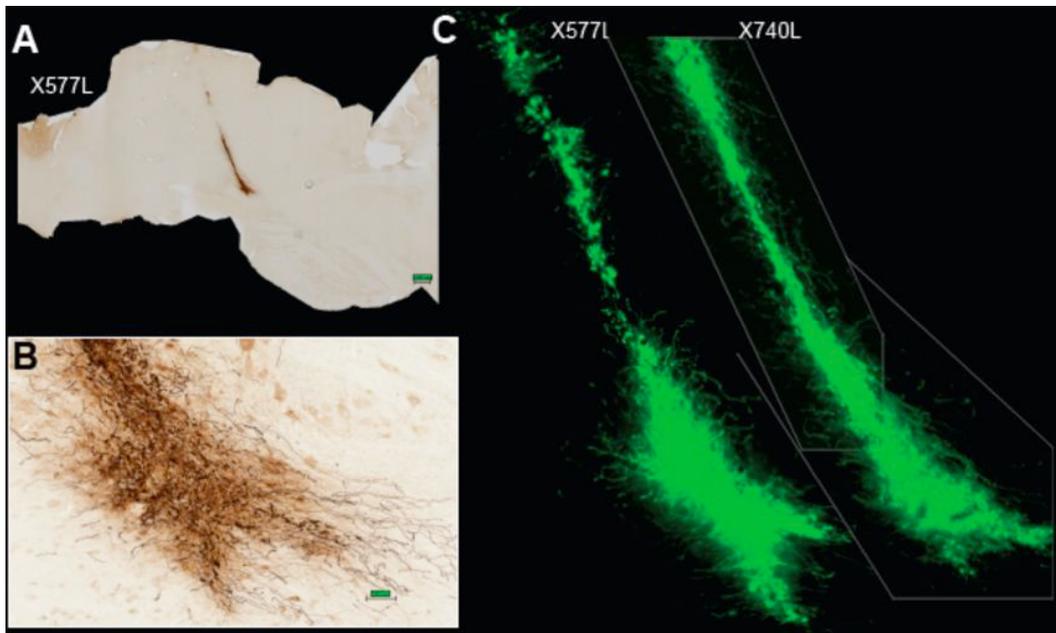
**Figure 4-7. Robust engraftment of hfNPC adjacent to host neurons after 1.5 months.** BrdU-ir donor cells (green) [A-I] were located at the more dorsal and rostral regions of the SN and typically were found to engraft near host TH-ir cells (red). No TH-ir/BrdU-ir cells were found at 1.5 months [A-C]; however several BrdU-ir nuclei had morphological features consistent with mitotic-figures (top left in panel B) but were not Ki-67 or PCNA-ir. [D-I], Confocal z-stack reconstruction. Different z-planes 30um apart at the identical (x, y) plane clearly indicated that BrdU-ir cells had integrated immediately adjacent to, but typically not within, the endogenous layer of TH-ir (red) DAergic cells [D, E]. 3D-reconstruction of z-plane images further demonstrate that donor cells occupy space near but not generally in the same plane as endogenous TH-ir cell bodies [F-I].



**Figure 4-8. Long-term engraftment of GFP-ir donor cells.** Fluorescence immunohistochemistry confirmed significant GFP-ir graft survival (green) at the dorsal extents of the SN, as indicated by TH-ir dopamine cells (red) [A, B (inset of dashed box in A)]. Another GFP-ir graft was located in an equivalent location, while two other grafts were located at the most dorso-rostral extent of the SN. In all four cases, donor-derived fibers projected within the host SN.

Analysis of hfNPC grafts at 11-months post-transplantation confirmed long-term survival of dense GFP-ir grafts at the dorsal region of the SN [Figure 4-8 A, B], bearing extensive neuritic processes in 4 out of 5 animals [Figure 4-9 A-C]. Hemispheres were analyzed and scored arbitrarily from (0-5) for overall engraftment success [Supplementary Table 4-2] and GFP-ir serial sections were analyzed volumetrically to determine overall graft density [Supplementary Table 4-2]. The one animal that did not show markers for graft survival and the two animals transplanted with dead cells (control) displayed no aberrant host pathology in the SN or apparent long-term injury from the transplantation procedure. In addition, GFP-ir grafts retracted into the needle tract in three out of four animals [Figure 4-9

A, C; Figure 4-10 A], where many dorsally oriented GFP-ir cells displayed a highly branched stellate morphology, but lacked GFAP-ir. Interestingly, the largest grafts, such as the ones in [Figure 4-9 A, C, Figure 4-10 A], bear a striking morphological resemblance to grafts reported previously utilizing human dopaminergic mesencephalic neuroblast implanted into the rostral mesencephalon or internal capsule<sup>106</sup>.

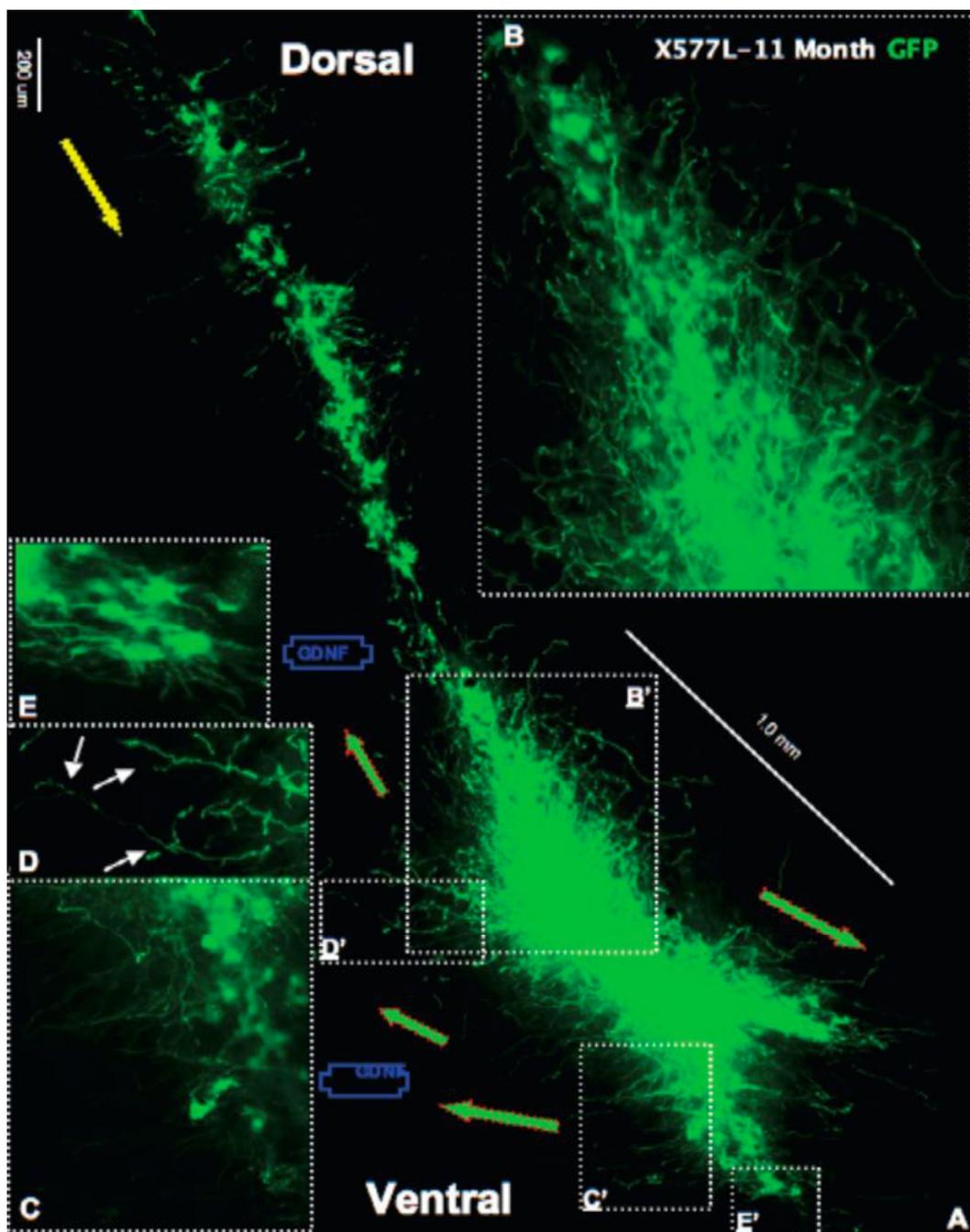


**Figure 4-9. Long-term engraftment of donor cells and extensive neuritic profile 11-months post-transplantation.** After 11-months, robust GFP-ir donor-derived grafts [A-C] were found to project extensive fibrific output into peripheral tissue [B], including SN. Cells were also found filling the injection cavity created from needle insertion-extraction in 3 of 4 animal successfully transplanted with GFP-ir [A, C]. The most robust grafts analyzed at 11-months are remarkably comparable to grafts reported previously utilizing human DA-neuroblasts implanted into the NS, however lack extensively long (> 3mm) neuritic projections growing toward striatal target regions. Migration of cell bodies from the graft into peripheral regions was never seen [C], including to the contralateral hemisphere.

#### 4.3.11 Characterization of Neuritic Outgrowth

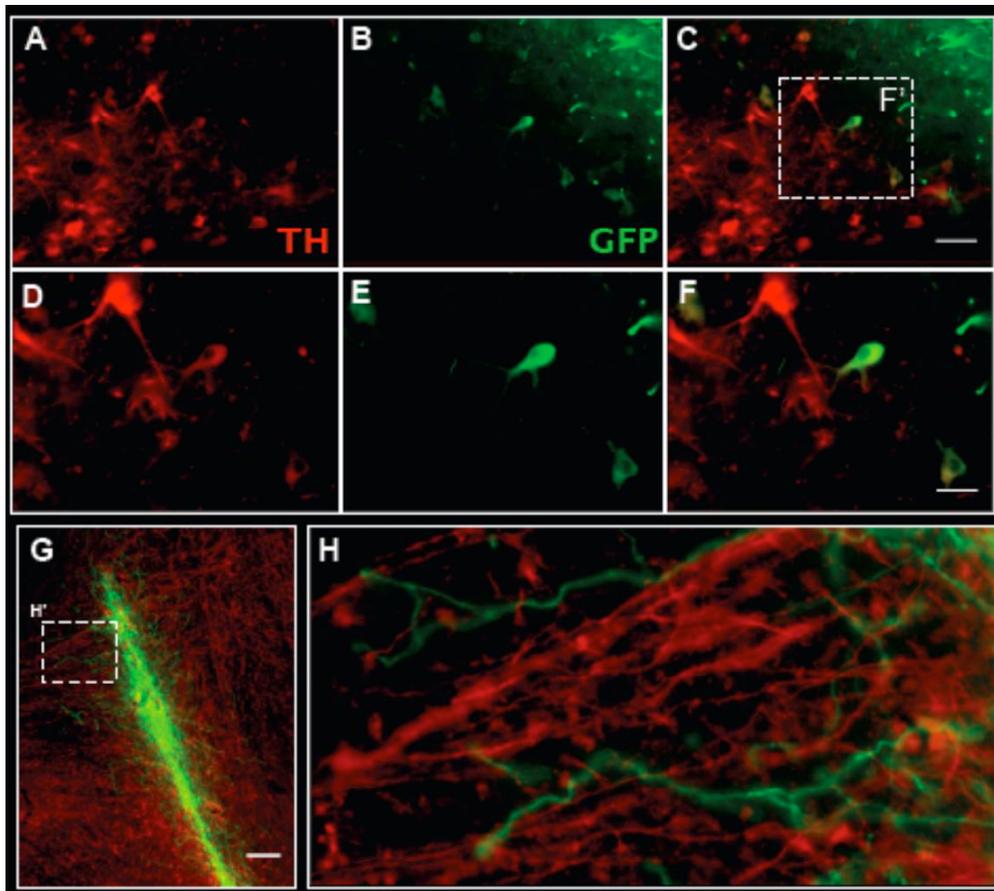
Morphologically, donor GFP-ir cells generally bore a soma located within the core of the graft site [Figure 4-10 B, E] and one or more processes extending up to greater than 1 mm out into host tissue from the periphery of the graft zone [Figure 4-10 C, D]. Numerous neurites

were found coursing in multiple trajectories, with no definitive directional preference of growth; however, GFP-ir cells located at the dorsal-rostral region of the graft core [Figure 4-10 D] displayed a more rostral pattern of growth up through the zona inserta ascending past the subthalamic nuclei toward the internal capsule and fields of Forel, while more caudal cellular protrusions generally remained within the graft or projected ventrally into the SN and further toward the cerebral peduncle and medial lemniscus of the lower brainstem. Interestingly, many GFP-ir fibers coursed immediately adjacent and/or through endogenous myelinated white matter tracts.



**Figure 4-10. HfNPC develop an elaborate neuronal profile after 11-months in-vivo.** A typical GFP-ir section (same section from Fig. 4-3C) from within the core of a hfNPC-derived graft after 11-months demonstrating a variety of morphological characteristics within the graft, including concentration of nuclei within the immediate graft zone [B, E], as well as extensive neuritic branching resembling developing axons [C, D] into peripheral tissue.

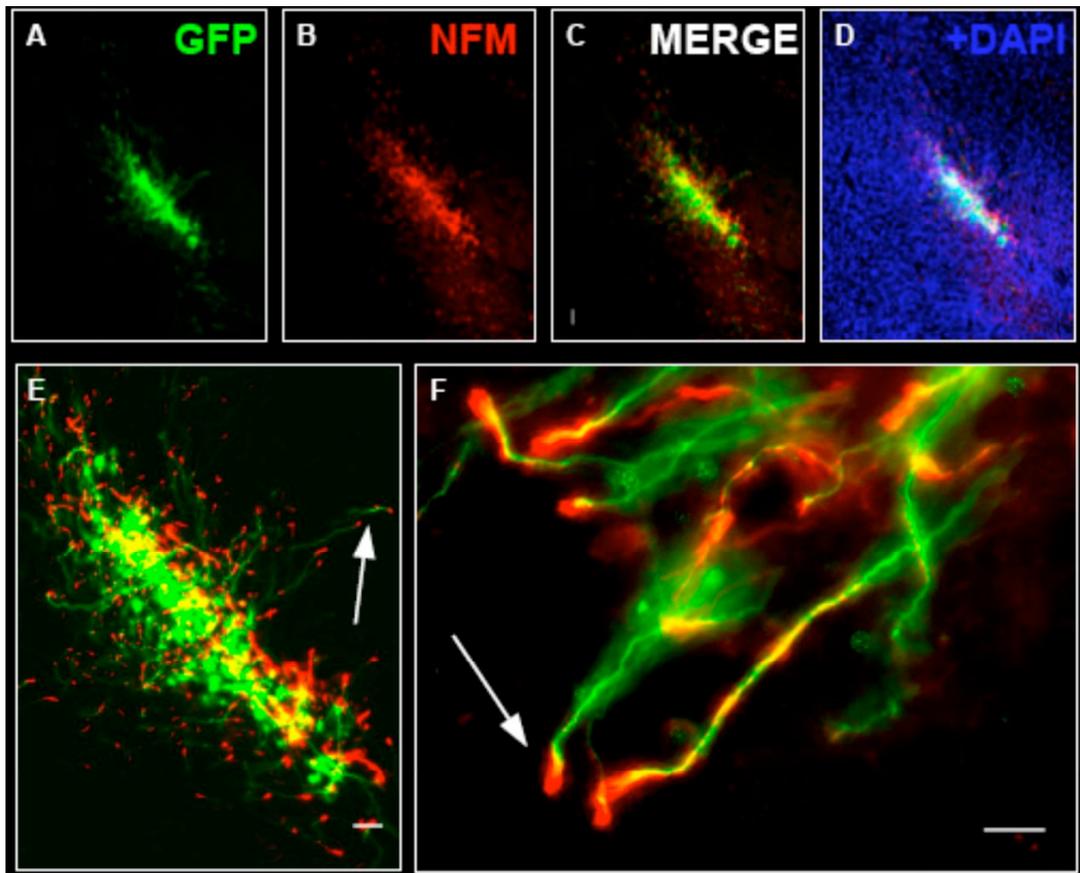
Analysis of GFP-ir neurites revealed patterned growth along a trajectory congruent with host fiber tracts [Figure 4-11]. Long protruding GFP-ir neurites coursed the SN and the needle tract, extending rostrally with a striatal trajectory in parallel with host-TH-ir fibers of the medial forebrain bundle (MFB) and NS pathway [Figure 4-11 G, H]. GFP-ir processes located at the dorsal-rostral region of the graft, as well as those located where the needle tract overlapped with the NS tract, often terminated in bifurcate forked end-feet, morphologically resembling growth-cones [Figure 4-11 H]. Throughout the graft, intricate rostro-caudal branching of thin GFP-ir fibers could be seen at the ends of thick axons, with highly ramified terminals located at both short and long distances from the transplantation zone.



**Figure 4-11. HfNPC graft development is influenced by host-environment.** Donor derived GFP-ir cells (green) co-express TH (red) [A-F] and project growth cone-like processes parallel to spared endogenous TH-ir nigrostriatal circuitry G, H]. Some GFP-ir soma were also TH-ir but typically lacked significant arborization in areas where grafted cells overlapped with spared endogenous TH-ir dopamine cells in the SN [C, F]. Regions where grafted cells overlapped with host fiber tracts displayed a clear preference of growth congruently alongside these pre-existing networks [G, H], arguing for a permissive, instructive environment with the adult primate brain.

In addition, some GFP-ir/pan-NF-ir fibers projected rostrally into host tissue then coursed back caudally towards the direction of the SN, resembling a circuitous loop [Figure 4-12 A-F]. It is possible that these projections are responding to local signaling molecules and coursing back towards a nigral target trajectory, in the same manner as young striatonigral neurons would correctly home to their target and innervate the developing brain in-vivo. Directional outgrowth, therefore, appeared to also be significantly altered by a permissive

local microenvironment and relevant endogenous excitatory and inhibitory guidance molecules in addition to direct chemotropic guidance from AAV5-GDNF.

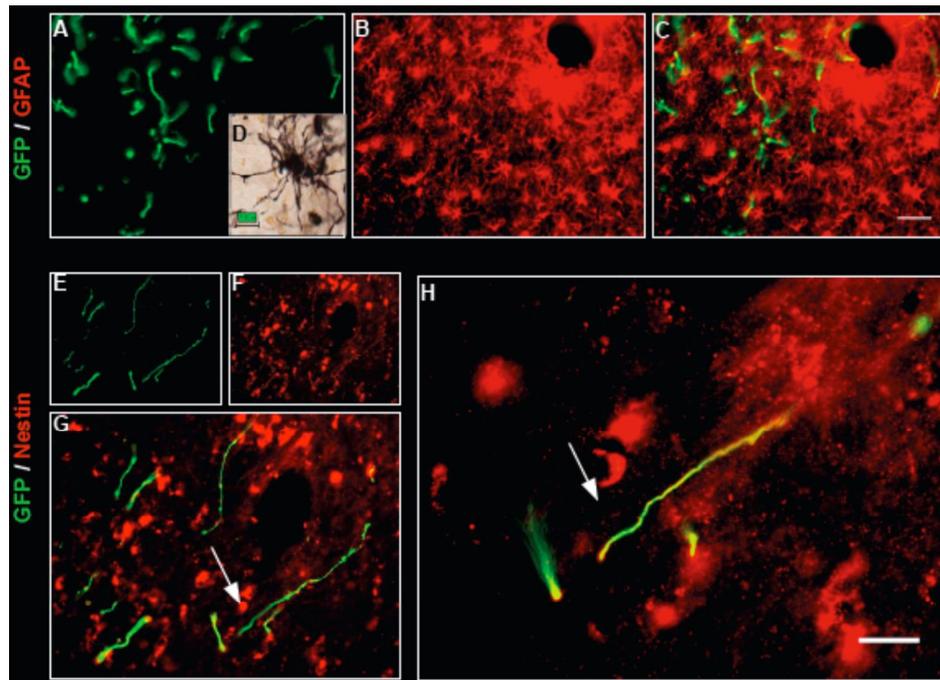


**Figure 4-12. Neuritic processes pursue multiple trajectories along host circuitry.** In some instances, GFP-ir/pan-NF-ir fibers (green/red) were found exiting rostrally into host tissue and coursing back caudally towards the SN, resembling a circuitous loop [Figure 4-6 A-F]. NF-ir was most abundant at the distal tips of GFP-ir projections (arrows) and became spotty towards the soma. Interestingly, NF staining revealed that the direction of GFP-ir fibers often favored that of host fiber-tracts and suggests local signaling molecules may override GDNF chemotaxis. DAPI staining [D] demonstrates no morphological aberrations to host cytoarchitectural arrangements were induced by hfNPC grafts.

#### 4.3.12 Graft Lineage Specification:

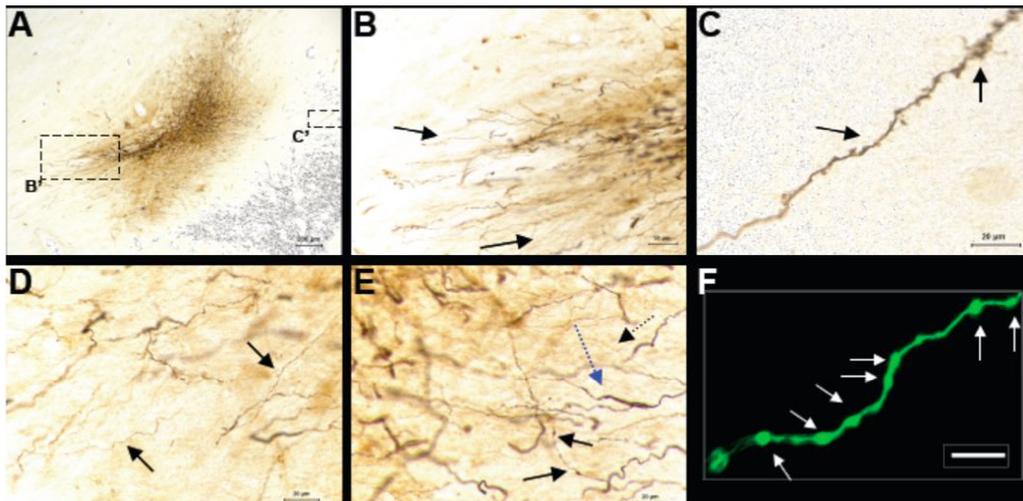
The lineage fate of grafted hfNPC was analyzed by IHC-IF for neuronal and glial lineage-related markers. After 11-months, 12-17% of grafted GFP-ir neurites co-localized with the undifferentiated lineage filament Nestin [Figure 4-13 E-H], with a lesser proportion

(3-9%) also expressing Vimentin (data not shown). Interestingly, 67-87% of GFP-ir cells co-localized with pan-neurofilament protein (NFM) [Figure 4-12 A-F] however they did not express significant levels of the mature neuronal protein, MAP-2, (~1% of GFP-ir/NFM-ir cells). In addition, few ( $\approx$  .1-1.0%) donor-derived GFP-ir cells co-expressed the DA marker, tyrosine hydroxylase (TH) [Figure 4-11 A-F]. Some GFP-ir/TH-ir cells developed larger soma, but without marked arborization of processes, suggesting that these cells were of an immature DA phenotype. Interestingly, grafted cells no longer expressed GFAP or Tuj1/beta-3-tubulin, indicating downregulation of these proteins post-transplantation, even in astrocyte rich regions of the brain [Figure 4-13 A-C].



**Figure 4-13. Engrafted GFP-ir cells downregulate immature filamentous proteins in-vivo.** Eleven month's post-transplantation, some GFP-ir donor cells (green) had the characteristic morphological appearance of astrocytes [D]; however, surprisingly, grafted cells did not co-localize with GFAP (red), even in regions where many endogenous astrocytes were present [A-C] and despite high expression in-vitro. Analysis of the “immature” neural marker filament Nestin (red), revealed co-localization with GFP-ir donor cells (green) only at the distal ends of some fibers [E-H], further indicating a switch from expression of stem cell to neuronal related proteins.

In one animal, GFP-ir cells were also found within the host ventricular lumen and SVZ at the dorsal region of the needle tract [Supplementary Figure 4-2]. Cells remaining at the ventricular surface retained a type-B/C cell phenotype, displaying either elongated radial processes resembling radial glia or bipolar migratory cells. Some of these cells were also Vimentin-ir (data not shown), indicating that HFB-2050 hfNSC were capable of integrating into the adult primate SVZ and likely incorporated into the host SVZ niche, thus further supporting HFB-2050 as a bonafide forebrain SVZ hfNPC. Furthermore, some GFP-ir neurites were also found lining branched vascular walls and expressed complimentary markers in-vitro.



**Figure 4-14. Morphological characteristics of 11-month grafts suggest differentiation into multiple neuronal phenotypes.** Histological enhancement of GFP-ir fibers revealed extensive fibrillar growth reminiscent of mature monoamine neurons [A, B, D-F]. Extremely fine GFP-ir fibers (arrows in [B] and [D]); dashed-black arrow in [E]) adorned with varicosities (solid-black arrows in [E], white-arrows in [F]) projected within the SN and were extremely dense at the ventro-caudal region [B] and dorsal-rostral of the graft [B]. Some GFP-ir processes were thicker and/or smooth (blue arrow) [E] and may reflect non-terminal ends of axons, while other longer processes terminated in, “spine-like” structures (black arrows, [C]).

Many longer (>800um) GFP-ir neurites exhibited morphological features of a variety of more mature, monoamine neuronal phenotypes<sup>252</sup>. Very fine, densely packed, small (0.3-.06 um) beaded varicose GFP-ir fibers, characteristic of DA neuron terminals<sup>253-256</sup> were found as

diffuse spotted masses within the SN and protruding from the dorsal-rostral and rostral-caudal regions of the graft [Figure 4-14 A, B, D, E]. In addition, many GFP-ir cells bore both thick and thin smooth processes [Figure 4-14 D, E] similar to non-terminal ends of monoamine axons, as well as those bearing both small and large beaded processes typical of the catecholamine (DA and norepinephrine (NE)) and 5-HT neurons, *in-vivo*<sup>257-281</sup>. Many processes of variable thickness had abundant round or fusiform enlargements, “varicosities”, spaced at variable distances and number per unit length [Figure 4-14 F]. These structures are morphologically consistent with pre-synaptic sites for amine synthesis, storage, and release, analogous to synaptic boutons found at axo-dendritic junctions. Furthermore, many longer processes terminated in highly branched, “spine-like” structures [Figure 4-14 C].

#### **4.3.13 Fluorogold Tract Tracing:**

To determine if GFP-ir neuritic processes innervated striatal targets, we analyzed retrograde transport from striatal terminals to the SN using Fluorogold (FG). Densely labeled FG-ir endogenous cells were found extensively throughout the injected striatum [Supplementary Figure 4-3 A], however relatively few (1-3%) endogenous TH-ir cell bodies in the SN displayed marked FG-ir [Supplementary Figure 4-3 B]. No GFP-ir/FG-ir donor cells were found in the SN, suggesting a lack of striatal innervation after 11 months *in-vivo*. In addition, GFP-ir processes were never found near the most caudal striatal regions arguing against striatal innervation or NS reconstruction. These data also suggest that FG may not be an efficient retrograde tracer in the primate or that endogenous TH+ nigral cells were deinnervated by MPTP exposure at their striatal terminals, the latter of which can be negated as we saw many spared endogenous TH+ projections throughout the NS.

#### 4.3.14 Safety Concerns and Immunological Response:

In order to address safety concerns involving cellular transplantation of fetal CNS tissue<sup>282</sup>, we analyzed brains for signs of aberrant pathology and immune response. We found no evidence for gross anatomical differences within grafted tissue by Cresyl Violet [Supplementary Figure 4-3 C] or DAPI staining. In addition, Ki-67 and PCNA immunostaining revealed no evidence for sustained proliferation within grafts at 1.5 or 11 months, suggesting that “bonafide” hfNPC do not continue to replicate long-term, initiate glioma-like tumors, or induce aberrant host pathology. Furthermore, cd11b staining indicated no major immune response activation of microglia around the grafted region at 11 months in hfNPC grafted animals or in the two animals that received killed-cell control transplants. Together, these results indicate no major anatomical changes or inflammatory reaction in animals with successful grafts, unsuccessful graft (one of five animals injected with HFB-2050-GFP), or controls that emulate an unsuccessful graft (dead cells). Occasionally, red blood cells were found within the injection site, indicating that a small amount of blood likely entered through the needle tract and into the VM during surgical injection. In addition, FG markedly damaged striatal tissue directly adjacent to the injection and was often accompanied by a small striatal hemorrhage.

All subjects were video monitored extensively both pre-MPTP injection and throughout the entire experimental timeline to ensure good health. Although animals selected for this study were deemed behaviorally asymptomatic for both PARK Score and Healthy normal behaviors, we continued to monitor them for negative side-effects. No animals regressed statistically for the entire post-surgical intervention period for both parameters; however, two hfNPC transplanted animals displayed improved scores for signs of healthy behaviors that neared statistical significance. Interestingly, these animals also displayed the highest survival and most extensive neuritic output, suggesting that these grafts may be

functionally active. At no point did any participants show adverse behaviors or signs of distress. These data support the safety of hfNPC grafts in the primate brain.

#### 4.4 Discussion

The results demonstrate that human fetal NPC xenografted into the primate VM are able to generate new immature neuronal phenotypes and project neuritic processes along the host NS in the dopamine-depleted midbrain. Previous studies have demonstrated axonal outgrowth along the nigrostriatal pathway and other brain regions in rodents using human or pig neuroblasts<sup>90,100-102,105,106,283-285</sup> indicating the adult NS as a permissive environment for regenerative growth. In addition, exogenous GDNF has been shown to enhance survival and direct neuritic outgrowth from grafted cells in both the rodent and primate dopamine-depleted brains<sup>69,71,84,87-90,96,286,287</sup>. Furthermore, striatal over-expression of GDNF has been shown to enhance endogenous axonal regeneration in the lesioned NS as well<sup>75,287</sup>

In a series of studies, we demonstrated that hfNPC, derived from the forebrain germinal zone, retain the capacity to engraft in the primate CNS, migrate to disease loci, and promote functional improvements in parkinsonian monkeys<sup>11,135,136</sup>. Histological analysis indicated only a small percentage of grafted cells differentiated into phenotypically DA cells (BrdU+/TH+ or DAT+), suggesting that the behavioral rescue and normalization of endogenous TH-ir cell numbers/area were related to indirect, secondary support of spared host cells. Similarly, others have shown lesion induced-migration<sup>288</sup>, differentiation into NFM-70-ir/Map2-ir neurons and GFAP-ir astrocytes, as well as extensive long-term axonal outgrowth<sup>289-292</sup> from undifferentiated hfNPC grafted into rodents. In addition, when differentiated in-vitro or transplanted into dopamine-depleted, 6-OHDA lesioned rats, differentiated cells with neuronal phenotype also expressed TH, albeit sometimes transiently<sup>289,290,292,293</sup>, suggesting that hfNPC retain the plasticity to produce disease relevant

DA neurons in-vivo; however, the extent to which these cells might fully differentiate and mature into disease-specific DA neurons, long-term, has never been fully or adequately accessed. Interestingly, hfNPC engineered to overexpress GDNF appear to remain predominantly undifferentiated (90%nestin, 5-8% GFAP, 0% and <1% NeuN discrepancy) when transplanted into rodents and one primate and do not differentiate into DA neurons<sup>294</sup>.<sup>295</sup>. While these results appear promising, it remains relatively unclear to what extent SVZ-derived hfNPC can differentiate into mature striatally integrated, A9-(Girk2-ir) DA neurons and respond to local axon guidance molecules when transplanted into the lesioned adult primate brain.

The use of sagittal plane sectioning and hfNPC engineered to express an eGFP reporter allowed us to unambiguously identify graft-derived cells and their neuritic output from endogenous cells long-term and evaluate their morphological and histological profile in relation to host architecture. Robust long-term graft survival and extensive neuritic outgrowth was evident at 11-months post-transplantation. Large, densely clustered GFP-ir grafts projected neurites in many directions from the core of the graft out into the surrounding tissue; however, the directional output was clearly polarized at anatomical loci where grafted cells overlapped with host neural circuitry. Many GFP-ir processes at the dorsal end of the graft exited at the rostral pole and positioned themselves rostrally in a parallel trajectory to host TH-ir NS or MFB fibers. Some of the processes resembled developing/regenerating neurons with bifurcate endfeet, similar to growth cones. While there were several clearly polarized areas of growth, neurites were found projecting three-dimensionally from the periphery of the graft in nearly all directions. Therefore, the grafts generally had an ovular, 3D shape (much like a watermelon) with axonal projections extending out from the periphery (rind of the melon). It appears that wherever the graft happened to overlap with endogenous circuitry, the donor axons aligned and positioned themselves in a similar orientation to the host neuronal fibers.

The specificity for host fiber trajectories suggests that local axon guidance molecules persist in a permissive environment and may directly influence hfNPC neuritic output in the dopamine-depleted primate brain. Netrins, Semaphorins, Ephrins and Slits play critical roles during the development of midbrain DA neurons<sup>296-310</sup>, but it is still relatively unclear to what extent these axonal guidance cues persist into adulthood in the primate brain and if they retain their instructive patterning cues long-term. Supporting previous studies utilizing more mature neural substrates, hfNPC also display directionally polarized outgrowth along host NS trajectories, specifically when the position of the graft was such that it intersected along or adjacent to endogenous fiber tracts. Importantly, the results demonstrate that a partially intact endogenous neural circuitry is likely to facilitate (and may be necessary for) NS reconstruction utilizing homotopic neural grafts. In addition to maintaining instructive guidance cues, endogenous DA fibers likely secrete neurotrophic growth promoting factors like BDNF as well. In this study, animals with partial lesions (loss of approximately 50% TH-ir) were specifically chosen to allow spared nigrostriatal fibers to facilitate directional outgrowth through the NS pathway, as seen in the 6-OHDA-lesioned mouse<sup>90</sup>. Fully lesioned animals with little to no DA nigrostriatal fibers remaining likely would not retain the axonal guidance molecules or produce trophic stimuli necessary for directed outgrowth. Therefore, as a whole, it appears that the partially lesioned primate brain retains a permissive environment to harbor neuronal differentiation and outgrowth of undifferentiated hfNPC.

The most plausible explanation for this targeted axonal outgrowth is that hfNPC retain their developmental plasticity and are able to utilize preserved trophic signaling cues from positions surrounding the source of their target in the adult brain. Newly grafted pioneer-like axons<sup>311-316</sup> may use existing spared adult axons as tracts, utilizing endogenous TH-ir fibers as nutrient and matrix rich scaffolding for generating new tissue. Interestingly DARPP32-ir medial spiny striatonigral projection neurons extend long fibers along congruent trajectories to

TH-ir nigrostriatal fibers<sup>317-321</sup>. These neurons are a potent source of GDNF for NS neurons and thus may facilitate a permissive growth environment for new graft-derived axonal processes. Grafts of DA neuroblasts have been shown to project axons that course in direct apposition to striatonigral DARPP-32-ir fiber bundles in the adult 6-OHDA lesioned adult mouse brain<sup>90</sup>. Furthermore, in the study, overall graft survival, efficiency of NS reconstruction, and degree of neuritic branching were enhanced by the addition of exogenous striatal AAV2/5-GDNF. Limited tissue constraints did not allow for co-analysis of GFP-ir neurites with DARPP-32 in the present study; however it is possible that a similar pattern might be found for hfNPC as well.

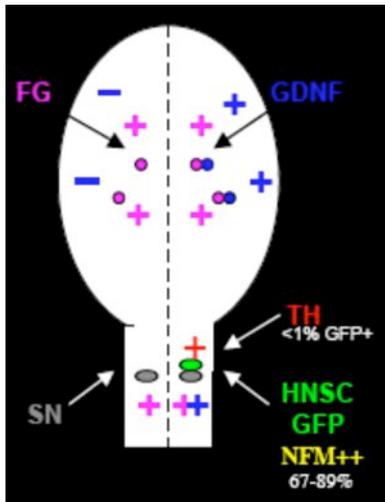
Congruent with studies in rodents and primates<sup>90,322</sup>, we found that AAV5-GDNF most efficiently transduced striatal projection neurons, providing GDNF neurotrophic support to surrounding striatal tissue, as well as being anterogradely transported downstream along the striatonigral pathway to relevant targets including the SN. Others have shown a similar pattern of anterograde transport from the caudate to the globus pallidus, internal globus pallidus, and substantia nigra pars reticulata after 3 months using AAV5-GFP in *Cynomolgus* monkeys<sup>322</sup>. In addition, the percentage of transduced GFAP-ir glial cells in the present study ranged from at most 10% to less than 5% compared to NeuN-ir neuron structures, in agreement with *Cynomolgus* monkeys transduced with AAV5-GFP<sup>322</sup>. Interestingly, far fewer GFAP-ir cells were found transduced with AAV5-GDNF than previously reported using AAV5-GFP in *vervet* monkeys, where nearly half of the transduced cells in the striatum were GFAP-ir<sup>323</sup>. Large host cell bodies within the SN were also transduced, indicating some degree of retrograde transport as well. Thus, donor and endogenous cells are simultaneously exposed to GDNF at multiple levels: within the SN enhancing donor survival within the GFP-ir graft site, along GFP-ir neurites adjacent to GDNF transduced striatonigral projections, as well as locally within the striatal target to enhance target innervation. Therefore, GDNF acts to

enhance endogenous striatonigral projections that deliver GDNF downstream, preserving spared host TH-ir fibers and promoting graft survival and possibly a permissive growth environment to establish new NS projections.

Interestingly, some NFM-ir fibers appeared to project with circuitous trajectories. These cells generally coursed dorsal-rostral along or adjacent to NS fibers, then appeared to turn back caudally, coursing with a ventral trajectory in a similar manner as developing striatonigral projections. Newly developed pioneer-like axons and existing NS and striatonigral axons may be playing dual roles in these circumstances. Donor axons may be receiving multiple signaling cues, first locally in the SN to project in accordance with NS axonal signaling cues then once reaching and interacting with existing striatonigral projection neurons, turning back under the influence of their opposing guidance molecules.

Multiple neuronal phenotypes with morphological parameters relevant to monoamine neurons were present, displaying ramifications that closely resembled that of host circuitry, specifically those of the intrinsic DA pathways. Fine, densely packed, GFP-ir processes with small beaded varicosities were found at the ventral region of the graft coursing in a ventral-caudal trajectory within the host SN. These thin fibers were often bundled together with a diffuse or dotted staining pattern, reminiscent of DAT bearing terminals, *in-vivo*<sup>259-268,270,271,273-275</sup>. Morphologically, DA, NE, and 5-HT monoamine neuron terminals are generally indistinguishable without the aid of histochemical enhancement, bearing abundant varicosities spaced somewhat irregularly throughout the length of the fiber<sup>259-263,266-269</sup>. We found many GFP-ir fibers coursing from the periphery of the graft with both small and large oval to round varicose enlargements. In general, varicosities varied in length, and width, and were not spaced with any marked regularity. These structures highly resemble pre-synaptic boutons and are the site for amine synthesis, storage, and release of amines *in-vivo*. In contrast, many processes were long and generally smooth at non-terminal ends (closer to soma), but often

terminated in highly branched structures, also characteristic of monoamine axons *in vivo*<sup>259,261,266,267</sup>. Other fibers terminated in highly branched spine-like structures, indicative of inhibitory complexes.



**Diagram 4-3. Schematic representation of 11-month histological data.** After 11-months *in vivo*, robust expression of AAV5-GDNF (blue +) was confirmed throughout most of the striatum, as well as within the ipsilateral SN, but never expressed in the contralateral hemisphere. HfNPC-eGFP (green oval) differentiated into a predominantly NFM-ir phenotype, but with a very small number of GFP-ir/TH-ir cell bodies (red +). Fluorogold (pink +) was found surrounding the injection site but rarely noticeably transported into the SN (gray oval) and never found within grafts themselves.

Immunohistochemically, the majority of grafted cells appeared to be transitioning from immature (Nestin+) precursors and preferentially differentiating into highly arborized, NFM-ir neuronal phenotypes within the dopamine depleted VM [Diagram 4-3]. While many GFP-ir cells morphologically resembled monoamine neuronal phenotypes, most donor-derived cells lacked expression of several characteristic neuronal markers, including TUBB3 and MAP-2(a+b). However, a few donor derived GFP-ir cells possessed characteristics of immature DA cells, including TH-ir soma, providing evidence that hfNPC may develop into lineage-appropriate DA cells when placed homotopically into the SN. Interestingly, a small percentage of larger cells located at the periphery of the caudal region of graft were GFP-ir to

a lesser extent and had a morphology reminiscent of host nigral DA neurons. While these cells were excluded from analysis, it is very possible that these are an example of donor cells that have turned down expression of the eGFP transgene upon differentiation and maturation in-vivo. Previous reports have demonstrated similar results utilizing the CAG-promoter in rodents and primates; therefore, future studies utilizing a secondary reporter under the control of the TH-promoter similar to those used in the mouse<sup>90</sup> will be necessary to further eliminate these apparent discrepancies and determine whether SVZ-derived hfNPC can truly recapitulate a relevant A9-DA neuron in-vivo. Limited tissue did not allow for analysis of DA subtype, specifically to determine if any TH-ir donor cells had matured into A9 (Girk2-ir) or A10 (Calbindin-ir) regionally specific DA subtypes. Given the substantial 11-month period of time between transplantation and histological analysis, as well as the facilitated support of exogenous GDNF along disease relevant tracts (NS), it is reasonable to suggest that transplanted SVZ derived hfNPC do not generate *significant numbers of new mature* DA neurons, even in the presence of neurotrophic chemotaxis. Furthermore, there was no conclusive histological evidence that GFP-ir/TH-ir co-labeled cells retained morphological parameters of size or axonal profile consistent with mature A9-DA neurons.

A distinct possibility remains that grafted human cells still need longer to mature and at 11 months are just now becoming receptive to chemotaxic guidance cues. At first glance, this scenario appears unlikely, given such a long period of time. Undifferentiated hfNPC should have fully matured by eleven months in-vivo; however, it may be true that human cells simply need a far greater period of time to fully develop and only once upon reaching a certain level of maturation, will they activate necessary signaling programs to correctly respond and project processes in accordance with permissive endogenous cues. In previous rodent and primate studies, hfNPC generally retained Nestin-ir and/or GFAP-ir up to 20 weeks post-transplantation<sup>288,290,324</sup>, suggesting that they had limited capacity to differentiate into mature

neuronal phenotypes; however, given that grafted cells retained a relatively immature but pro-neuronal immunohistochemical profile after 11-months in the current study, it is certainly plausible that these cells require a more extensive period of time to fully develop.

In theory, as cells continue to mature and begin utilizing axonal guidance molecules from existing host circuitry, they may actually be capable of integrating striatal targets, similar to results seen utilizing fetal VM neurons<sup>89</sup>, where outgrowth was found coursing to the most caudal portion of the caudate, roughly 5-7mm. In this theoretical model, regeneration and NS reconstruction would occur simultaneously utilizing two distinct complementary mechanisms. In the first phase, exogenous GDNF promotes regeneration and protection of spared endogenous DA neurons and their striatal projections via mechanisms discussed above while also promoting a survival advantage to grafted hfNPC through a permissive neurotrophic environment. During this extended period of time, host neurocircuitry begins to regenerate as well as produce potent stores of local trophic factors that may act on developing grafts, preferentially exerting DA signaling cascades and enacting parallel receptor programs, thus priming grafted cells to now respond to axonal guidance molecules being displayed by locally restored circuitry. In the second phase, newly committed neuronal processes would then utilize rejuvenated host circuitry and local guidance cues to properly course the NS pathway and reinnervate the GDNF-rich striatum.

Surprisingly, GFP-ir cells did not express GFAP, although many had a morphological profile consistent with that of astrocytes. In previous studies, where hfNPC were placed into the striatum, we found that grafted cells migrated extensively and differentiated into a small percentage of astrocytes<sup>136</sup>. Similarly, hfNPC grafted into normal or 6-OHDA lesioned rats differentiate into astrocytes<sup>7,288,289,291,292</sup> and migrate extensively when implanted ectopically into the striatum of fully lesioned rats<sup>292</sup> or transplanted homotopically into the SN of striatal partial-lesion rats<sup>288</sup>. Interestingly, homotopic grafts of both undifferentiated and

predifferentiated (TUBB3-ir) hfNPC retained expression of both Nestin and GFAP 12-weeks post-transplantation in 6-OHDA partial-lesion rats. The data suggest that the signaling cues at the local transplantation zone as well as lesion zone may profoundly impact the fate of grafted cells. In this case, hfNPC were placed directly adjacent to the lesion zone (SN) of partial-lesion primates, and we saw no migration or astrocytic differentiation. Therefore, the striatal environment likely maintains astrocytic differentiation cues, whereas the VM appears to preference towards neuronal phenotypes. These studies highlight significant differences in the fate of undifferentiated hfNPC in relation to their site of transplantation as well as the species into which they are delivered. Secondly, the degree of lesioning and persistence of existing circuitry are likely to have a profound impact on donor cell fate. Fully lesioned, behaviorally parkinsonian animals have extensive degeneration of both DA circuitry as well as a general breakdown of the cytoarchitectural milieu, whereas asymptomatic animals retain enough DA transmission to maintain most healthy daily behaviors. With only partial degeneration, asymptomatic animals may not require secondary glial support or enact the same signaling mechanisms employed by a fully lesioned brain.

Safety considerations and immunological response to grafted cells are important obstacles to consider for any cell-based therapy. Host tissue within the transplantation zone displayed no aberrant pathology by Cresyl Violet or DAPI stain. In addition, grafted cells were negative for the proliferative markers, Ki-67 and PCNA, confirming previous findings that grafted hfNPC exit the cell-cycle shortly after transplantation<sup>292</sup>. Interestingly, after 1.5 months, we found a few cells within the core of the graft with morphological characteristics of late stage symmetric division; however, these cells did not stain positive for PCNA or Ki-67 as expected. These results are consistent with those seen in the rat<sup>292</sup>, where some hfNPC replicated several times before exiting the cell-cycle, but never maintained long-term sustained proliferation. Staining for inflammatory activation revealed little to no infiltration of

CD-11b-ir microglia or apparent long-term immune response of grafted cells. Overall, the results demonstrate that “bonafide” hfNPC do not proliferate long-term, initiate tumor like growth, or induce aberrant host pathology. Furthermore, hfNPC appear to be well-tolerated by the endogenous environment without major immunological rejection long-term or adverse effects derived from the transplantation procedure itself.

It is still relatively unclear whether undifferentiated NSC or lineage specific post-mitotic DA precursors/neurons represent the ideal substrate for neural regeneration in PD. It is important to examine each individually defined resource and test its potential separately from the others. Careful and meticulous side-by-side characterization of NSCs or DA progenitors derived from fetal cadavers, hESC, and induced pluripotent stem cells (iPSC) may ultimately lead to successful treatments. Biochemical and gene expression arrays indicate that HFB-2050 *SVZ-derived* hfNPC retain multiple characteristic consistent with a neural stem/precursor phenotype. Immunocytochemical analysis indicated that these cells exist as a heterogeneous population of neural stem and progenitor cells, highly resembling the neurogenic niche in-vivo. Interestingly, almost all hfNPC are Vimentin-ir, Nestin-ir, GFAP-ir, Sox-2-ir, Ki-67ir replication competent, and readily incorporate the thymidine analogue BrdU; however, upon exposure to an extracellular matrix-rich environment, a greater percentage of type-B, radial glial like cells were multi-nucleated and aligned symmetrically, suggesting they may be the true hfNSC that give rise to and support their slightly differentiated  $BLBP^{low}/DCX+$ , type-C and type-A hfNPC counterparts. Moreover, there is no evidence to suggest these cells are not still fully plastic and can revert to either lineage in response to local signaling molecules that may govern the overall percentages of each cellular sub-population. In fact, it has been shown that secondary type-C transit amplifying cells can function as multipotent NSCs, in-vitro, when exposed to appropriate mitogenic stimuli<sup>148,325</sup>, suggesting the cell fate determination transition from type-B to type-C cell may not be a unilinear event in-vitro, but rather a plastic

bi-directional spectrum of differentiation, poised to respond to local signaling molecules<sup>326,327</sup>. It is therefore likely that mature, high density, HFB-2050 MAN cultures are intrinsically interconnected networks comprised of both populations co-existing as one functional unit.

HfNPC (SC-23: derived from a premature infant of about 23 weeks gestation)<sup>150</sup> were sorted by immuno-flow cytometry for CD9, a marker associated with pluripotency in hESCs, revealed both positive and negative-staining cells in standard cultures (Phillip H Schwartz, CHOC; personal communication). A few days after sorting and separately replating each population, the morphology of the CD9+ cells differed from the CD9- population; a promising result in hopes of separating the stem from progenitor and differentiated cells. After two weeks in culture; however, the cultures appeared morphologically indistinguishable by phase microscopy. In addition, flow cytometry showed an equal distribution of CD9+ and CD9- cells in both cultures, suggesting that they had both "re-set" to the original distribution found in the initial parent culture. Over several weeks, daughter cultures inherently repopulated as a heterogeneous mixture; therefore, cells appeared to reorganize and adjust accordingly, demonstrating the inherent plasticity hfNPC possess. Interestingly, these cells were extremely similar to HFB-2050 in both mRNA and miRNA expression<sup>156,157</sup> and have similar morphological features in-vitro.

#### **4.5 Conclusion**

These results provide evidence that hfNPC can engraft and survive long-term in the MPTP lesioned primate SN while their distant striatal target is releasing an elevated concentration of GDNF. We demonstrate for the first time, extensive neuritic outgrowth from undifferentiated hfNPC in the primate dopamine-depleted midbrain. Specifically, smooth and fine varicose GFP-ir processes were found to course along trajectories congruent to endogenous neural circuitry suggesting that grafted hfNPC, much like their VM counterparts,

can be placed into appropriate homotopic loci and develop and respond appropriately to axonal guidance signaling molecules in the degenerative adult primate brain. The results also confirm previous reports in other regions of the CNS, indicating that the adult brain retains permissive instructive cues for repair and reconstruction<sup>90,100-103,106,110,328</sup>. In these studies, fate committed DA-neuroblasts were capable of innervating multiple highly specific short and long-distance target regions in the lesioned adult CNS.

Similarly, undifferentiated hfNPC demonstrate the capacity to differentiate and utilize local signaling cues retained within spared endogenous nigrostriatal circuitry. Interestingly, after 11-months, grafted cells remained somewhat immature and still appeared to be developing and growing in relation to host axon guidance cues, indicating that human stem cell derived neurons likely need a much longer time to fully develop and mature than rodent or primate fate committed DA-neuroblasts within the primate brain compared to rodent brain. Human cells have a much longer developmental timescale compared to rodents in-vivo; therefore, it is likely that they also remain receptive to growth factors and guidance cues throughout this longer process as well. Furthermore, undifferentiated stem cell derived grafts are likely to give rise to multiple alternative cell types, which may explain why outgrowth was somewhat promiscuous and did not appear to be quite as target-restricted compared to DA-neuroblasts, which have been shown to direct circuit-specific projections toward denervated targets along relevant nigrostriatal fiber tracts (medial forebrain bundle and internal capsule). It is still relatively unclear how long it would take in-vivo and to what degree SVZ-derived hfNPC can differentiate into fully differentiated A9-specific DA neurons and integrate distant targets. In the present study, we found ~.01-1% of GFP-ir donor cells also co-expressed TH, confirming our previous findings where AAV5-GDNF was not used combinatorially. GDNF, therefore, did not appear to exert a significant effect on DA differentiation, but rather improved graft survival and rescue of endogenous NS fibers. The experiments further support

the rationale that early initial functional improvements in parkinsonian primates reported previously<sup>11</sup> were likely due to trophic support and restoration of spared endogenous host-neurons not from direct cell replacement. Cell replacement strategies utilizing undifferentiated NPCs, therefore, may likely require several years to fully mature when directed by host tissue. In contrast, fate-committed DA-neuroblasts readily differentiate into TH-ir neurons and have been shown to accurately and efficiently reconstitute new nigrostriatal projections along existing host circuitry. It is likely that hESC and iPSC derived NSC and their differentiated DA precursors will also retain many of these valuable characteristics.

Great strides have been made in neural induction and maturation of A9-DA neurons from hESC and iPSC, in-vitro<sup>12-14,182,188,329-338</sup>. Additionally, the identification of new extracellular markers for FAC sorting specific populations of progenitors<sup>339,340</sup> will be crucial to predictably establish and reliably expand transplantable populations in a clinical setting. With the advent and present onslaught of iPSC biology, as well as the initiation of new fetal DA cell clinical trials in Europe, cellular transplantation appears to have come full circle. It may be time to test each specific cellular substrate side-by-side, to determine if we can make a more therapeutically relevant, renewable cell in large quantities, in-vitro and determine which stage of differentiation provides the most functionally relevant substrate for treatment of Parkinson's disease.

## **4.6 Methods**

**4.6.1 Derivation of hfNSC/hfNPC HFB-2050:** Primary neural stem/progenitor cell line (HFB-2050) was derived from the neuroectoderm-derived forebrain ventricular zone of a 13-week human fetal cadaver as previously described<sup>11,142,341-343</sup>. Cultures with normal karyotype and sustained proliferative capacity for greater than 20 population doublings were maintained as multilayer adherent networks (MAN) in *NB-B27 complete media* containing: Neurobasal

medium (Gibco) supplemented with B-27/without Vitamin A (Gibco), Glutamax (Gibco), heparin [8 $\mu$ g/ml] (Sigma), Normocin (InvivoGen), FGF-2 [20ng/ml] (Peprotech), and LIF [10ng/ml] (Millipore). Stable populations of hfNPCs were frozen as aggregates in 1 ml aliquots, approximately 10-15 cells in diameter as previously described<sup>142</sup> and stored as large batches for future use as a library of early passage biological replicates.

#### **4.6.2 Establishment and Maintenance of hfNSC/NPC Multilayer Adherent Network**

**(MAN):** Frozen HfNPC aggregates were “quick thawed” at 37°C from identical previously frozen aliquots containing 2-3x10<sup>6</sup> cells, washed immediately in NB-B27 complete media, centrifuged for 3.5 min at 300 rcf, resuspended in 8-ml fresh NB-B27 complete media, and plated at *uniform* density into vented Falcon T-25 cm<sup>2</sup> flasks. Flasks were incubated at 37°C, 5% CO<sup>2</sup>, *undisturbed*, to allow cellular clusters to form homogenous, “low affinity” adherence, replication competent MAN *colonies* as described previously<sup>142,344</sup>. After 72 hours, metabolized media was gently aspirated and replaced with 8-ml fresh NB-B27 complete media to expand hfNSCs colonies and allow individual cell clusters to extend processes and begin migrating into adherent *networks*. This process was repeated every 3 days to refresh mitogens and remove toxic metabolites. Within 7 DIV, cellular clusters were “moderately” adherent, highly migratory, and came into direct contact with adjacent neighboring clusters forming the “webbed” structural foundation for the multilayer network. After 2-weeks in culture, these “monolayer-like” colonies continued to expand into contact with each other forming a uniform, densely webbed, interconnected, migratory, *multilayer adherent network* of hfNPC.

**4.6.3 FACS Analysis:** HFB-2050-eGFP hfNPC were enzymatically dissociated into single cells with Accutase (Millipore), allowed to equilibrate for two hours and resuspended at 1x10<sup>6</sup>

cells/ml as previously described<sup>142</sup>. For lentiviral labeling efficiency, eGFP epifluorescence was analyzed in live cells without antibody enhancement. BrdU labeled cells were fixed in 4%PFA, incubated in 2N HCL @ 45°C for 30min, and stained by methods described below with AlexaFluor mouse-anti-BrdU-488 (1:250, Molecular Probes, Eugene OR). GFP+ or BrdU:488-ir cells were analyzed on a fluorescent activated cell sorter (FACS). For detection of BrdU, data was acquired with a FACSCanto (BD Biosciences, San Jose) flow cytometer and analyzed with FACSDiVa version-4 software. For GFP sorting, cells were sorted on a FACSVantageSE DiVa sorter with FACSDiVa version-4 software at 15psi with a 100-um ceramic nozzle and a drive frequency of 31,000 drops per second. BrdU-488 was excited at 488nm and detected with a 530/30 bandpass filter and PI was detected with a 670 longpass filter. 30,000 cell events were collected in each data file. Events were gated in forward and side scatter to eliminate debris and dead cells. Aggregates were gated out using pulse shape parameters. Fluorescence was determined and gated in comparison to appropriate controls: (unlabeled GFP-negative cells for GFP labeling efficiency) or (labeled + unstained, unlabeled alone, unlabeled + antibody; for BrdU labeling efficiency). All comparisons were carried out on cells at an identical stage of maturity and prepared in parallel. To assure specificity, sorted cells were verified visually by immunofluorescence microscopy (see below) and by FACS reanalysis. All sorts were repeated at least three times.

**4.6.4 Microscopic Confirmation of eGFP & BrdU Labeling:** Flow sorted cells were collected and reanalyzed with standard immunofluorescence microscopy. Briefly, cell suspensions were gently triturated and plated onto standard microscope slides or hemacytometer, coverslipped, and visualized with an Olympus IX-71 fluorescent microscope. Cells in ten random fields of ten coverslips per sample were counted in three experimental replicate samples.

**4.6.5 Immunocytochemistry (ICC)-Immunofluorescence (IF):** Routine IF techniques were applied to dissociated cells as well as cells grown on coverslips or TC-treated plastic chambers/wells as previously described<sup>142</sup>. Briefly, cells were fixed in cold 4% PFA (Electron Microscopy Sciences, Hatfield, PA USA) for 15-20min at 4C, then permeablized (non-surface antigens) in 0.1%TritonX-100 (Sigma-Aldrich, St Louis, MO USA) for 20min, and blocked in 3-5% Normal Donkey Serum (Jackson ImmunoResearch Laboratories, Inc., West Grove PA, USA) + 1-3% BSA (Sigma-Aldrich, St Louis, MO USA) in 0.2% TritonX-100 for 1hr at RT. Primary antibodies were diluted in 1:1 (blocking solution: dPBS + 0.1% TritonX-100) and incubated on cells overnight at 4C. Subsequently, cells were washed thoroughly, followed by secondary antibody incubation with species appropriate fluorescence-labeled AlexaFlour conjugates; Donkey anti-mouse, rabbit, or goat 488 (green), 555 or 568 (near red), 647 (far-ir) (Molecular Probes, Eugene OR, 1:500). Where applicable, nuclei were visualized with Hoechst 33342 (Molecular Probes, 6 ug/ml) (blue). Primary antibody was omitted from specified controls. Coverslipped samples were mounted on permafrost-coated slides in Vectashield (Vector Labs).

**4.6.6 Q-PCR:** RNA was extracted from cell pellets using the SV Total RNA Isolation System (Promega). Following purification, the extracted RNA was treated with RQ1 RNase-Free DNase (Promega) to ensure complete removal of any contaminating genomic DNA. In preparation for real-time PCR amplification, first-strand cDNA was synthesized from 100ng of each RNA template, using the ImProm-II(tm) Reverse Transcription System (Promega). Following thermal inactivation of the reverse transcriptase, the cDNA was used as the template in quantitative PCR amplifications. Amplification of pluripotency-associated transcripts was performed as two-color duplex reactions, with each pluripotency-associated

gene (labeled in FAM) duplexed with a reference transcript (GAPDH, labeled in JOE). Amplification of neural-associated transcripts was performed as monoplex reactions, with the reference transcript (GAPDH) also amplified as a monoplex. Quantitative PCR was performed using the StemElite(tm) Gene Expression System (Promega) and a LightCycler 480 instrument (Roche). Analysis of qPCR data was performed using the Plexor(r) Analysis Software (Promega). Primer sequences were developed and provided by Promega. Additional information or specific sequence inquiries available by request to Promega. During the amplification, the point where the decrease in fluorescence crosses the detection threshold is the CT; samples are compared using this number. A smaller CT value denotes a higher level of gene expression. CT was determined and normalized to the housekeeping gene, GAPDH; average CT=19.5.  $\Delta CT = (CT\text{-gene of interest}) - (CT\text{-GAPDH})$ . Significant expression differences between samples are defined as a difference in  $\Delta CT > 2$ .

**4.6.7 Immunoblotting:** Undifferentiated HFB-2050 MAN cultures were rinsed five times in cold dPBS and immediately lysed in RIPA buffer (Pierce #89900, Rockford, IL) + fresh HALT phosphatase and protease inhibitor cocktails (Pierce 78410, 78420) for 5-7 min with shaking. When lysis was complete, homogenate was pulled through an insulin syringe 3-5 times, sonicated for 5sec, and centrifuged for 10 min to clear debris. A small aliquot was removed and placed on ice for protein quantification (BCA Assay, Pierce) and the remaining cleared lysate aliquoted and stored at -20C. 5-30ug of solubilized protein from each sample were mixed 1:4 with sample buffer and NuPAGE reducing agent and heated to 70C for 10 min according to manufacturers instructions (Invitrogen, NuPAGE electrophoresis system). Samples and molecular weight standards (New England Biolabs, Ipswich, MA) were loaded onto precast NuPAGE Novex 4-12% Bis-Tris gels (Invitrogen) and electrophoresed in NuPAGE MES SDS Running Buffer + Antioxidant at 200V, 35min. After electrophoresis, the

proteins were electrically transferred to Immobilon-FL membranes at 30V for 1hr in NuPAGE Transfer Buffer + Antioxidant. After transfer, blots were incubated in PBS with 0.1% Tween-20 containing 5% Odyssey Blocking Buffer (LI-COR) or 5% nonfat dry milk. Blots were then incubated with the appropriate primary antibody (Nestin-human Ms 1:2000; Chemicon; GFAP Rb Dako 1:3000; BLBP Rb Chemicon 1:2000; DCX Gt SantaCruz 1:1000; Tuj1 Ms 1:2000; Covance) plus 0.1% Tween-20 and 3% Odyssey Blocking Buffer at 4°C overnight. Subsequently, fluorescently-conjugated secondary antibodies (AlexaFluor-680, Rockland-IR800, Ms, Rb, Gt; 1:10,000) were applied in 0.1% Tween-20 and 0.01% SDS, and immunoreactive bands were visualized using the Odyssey infrared imaging system (LiCorr Biosciences) with GAPDH (Chemicon 1:1000) serving as a loading control.

#### **4.6.8 Stereotactic Surgeries, Transplantation of hfNSC, and GDNF Vector Delivery: St.**

Kitts (African) green monkeys (*Chlorocebus sabaues*) were anesthetized with ketamine (10 mg/kg IM) and sodium pentobarbital (15-25 mg/kg IV), as previously described<sup>142</sup>. Briefly, animals were given atropine, intubated and the head was prepped, shaved, and mounted into a stereotactic frame using sterile technique. The animal was monitored for vital signs, EKG, pO<sub>2</sub>, and temperature and given a 30-50 ml/hour intravenous infusion of Lactated Ringer's solution. Each animal received 300,000 Units of Flocillin (penicillin G benzathine and penicillin G procaine) before surgery. After the animal was anesthetized to a level that showed no deep pain responses, the scalp was incised. Small 1-2 mm holes in the cranium were drilled at the desired coordinates (David Kopf, Tujunga, CA). A 22-gauge needle connected to a Hamilton (Reno, NV) syringe was lowered through drilled holes to the desired depth, and left *in situ* for 2 minutes pre- and post-injection, and in cases where the depth of the needle was varied, the needle was left *in situ* for 2 minutes before each withdrawal in an upwards direction.

Two days prior to transplantation, HFB-2050-eGFP MAN cultures were enzymatically dissociated into single cells with Accutase and replated at low sphere forming density to deter mass aggregation. Ten minutes prior to transplantation, donor cells were gently re-dissociated into a single-cell suspension at [50,000 cells/ul] in dPBS and aseptically loaded into a 22-gz needle attached to a 20 ul Hamilton syringe (Reno, NV) for stereotactic injection as previous. Six animals received undifferentiated HFB-2050-eGFP hfNSC, two received unmodified wild-type HFB-2050 hfNSC, and two received killed hNSC-eGFP (controls for transgene fusion and host immune response). HfNPC were grafted into the rostral mesencephalon immediately dorsal to or within the host SN using Kopf Instruments stereotaxic devices (Tujunga, CA) to target empirical stereotactic coordinates determined based on previous studies<sup>89</sup> [AP 11.1, Lateral 3.5, and Vertical 9.6 mm from actual ear bar zero] and driven by a microperfusion pump (Stoelting Instruments, Wood Dale IL) at a rate of 1 ul/min.

For vector delivery, 100 µl Hamilton syringe was driven by a perfusion pump delivering vector at 1 µl per minute (Stoelting Instruments, Wood Dale IL). The vector dose was  $1 \times 10^{12}$  viral genomes per milliliter. Two targets were injected within the hemisphere ipsilateral to the hfNPC injection; the rostral caudate nucleus (CD) and post commissural putamen (PM). After injection of vector was completed for each target, the needle was slowly withdrawn (1 mm/min) over a period of five minutes. The measurements used for the target regions of the brain were as follows: Caudate: AP: 21.1, Lateral: 4, Vertical relative to ear bar zero: 19; Putamen: AP: 21.1, Lateral: 10, Vertical relative to ear bar zero: 18.9. All animals were immunosuppressed for the duration of the experiments.

**4.6.9 GDNF Viral Vector Synthesis:** The methods for vector production have been published elsewhere<sup>345</sup> and are briefly, as follows: A single-stranded rAAV vector was constructed as

follows: Digest the plasmid pSub201 with XbaI and HindIII restriction endonucleases to remove the rep and cap fragments, and gel purify the 4000-bp plasmid backbone containing the AAV2 wt-ITRs. The desired transgene expression cassette was inserted between the XbaI sites to construct the single-stranded rAAV vector plasmid. For pSub201, HindIII was used in the digest to cut the rep and cap fragment in half for easy isolation of the plasmid backbone.

Self-complementary rAAV vectors were constructed as follows: The plasmid pHpa-trs-SK was digested with Acc65I and SalI to remove the CMV-GFP-poly(A) gene expression cassette and gel purify the 4166-bp plasmid backbone containing the mutated AAV2 ITR (left ITR) and wt-ITR (right ITR). Replace it with the desired transgene expression cassette between Acc65I and SalI. The size of the resulting rAAV construct (including two wild-type ITRs, which are each 145 bases for single-stranded rAAV genome, or two wild-type ITR and a mutated ITR, which is 120 bp, for self-complementary rAAV genome) should be between 3.4 and 4.8 kb, with the optimum size being that of the wild-type AAV genome of 4681 bp. Therefore, the size of the transgene cassette in the cloning plasmid, which includes the gene of interest and the promoter sequences necessary for its expression, spans between 3400 and 4400 bp (for single-stranded rAAV) or 1500 and 1950 bp (for self-complementary rAAV). Filler sequence was included if the transgene cassette is <3000 bp. The AAV ITRs are unstable in *E. coli*.

A large-scale plasmid preparation (at least 1 mg) of the rAAV vector and the suitable AAV helper plasmid and pXX6 plasmids was purified by double CsCl gradient fractionation. AAV serotype 5 was then generated by using different AAV helper plasmids of the pXR series (e.g., pXR1 for generating AAV serotype 1 capsids). For pXR plasmids, restriction digests tested plasmids upon arrival prior to transfer into bacteria (e.g., DH10b) and large-scale plasmid preparation. The restriction digest pattern was: pXR1: BamHI and SphI digest yielded 2949-bp and 4516-bp fragments; pXR2: BamHI and NcoI digest yielded 420-bp,

2722-bp and 4320-bp fragments; pXR3: BamH1 digest yielded 1528-bp and 5937-bp fragments; pXR4: BamH1 and SphI digest yielded 2051-bp and 5411-bp fragments; pXR5bam: BamH1 digest yielded 212-bp, 1443-bp and 5761-bp fragments. When growing the rAAV vector plasmid, the culture was not allowed to remain in stationary phase too long, as ITR deletions occur more rapidly after the culture has proceeded past log phase. The pXX6 plasmid was also unstable in bacteria. After receiving the plasmid, pSub201, pHpa-trs-SK, and XX6 was transformed into SURE bacteria (Stratagene) and the integrity of the miniprep plasmid was checked by restriction digest, and the integrity of AAV ITRs by SmaI restriction digest, followed by electrophoresis. After selection of an intact clone, individual glycerol stocks were prepared. The plasmid preparations were analyzed to be free of contaminants (ethidium bromide, CsCl, and RNA) before use.

#### **4.6.10 General Methods Used for Animal Care and Systematic MPTP Treatment: All**

animal work complied with National Research Council guidelines and was approved by Axion Research Foundation's animal care and use committee. The study was performed in accordance with U.S. federal guidelines and with the approval of the BIMR Institutional Animal Care and Use Committees (IACUC). Mature adult male African green monkeys (*Chlorocebus sabaues*, St. Kitts) were singly housed with natural daylight light/dark cycle at 17° N. Latitude. Monkeys were systemically treated with MPTP HCL (RBI, Natick, MA) for a five-day period (total dose 2 mg/kg of body weight)<sup>346-348</sup> at least five weeks before surgical injections with a rAAV-hu-GDNF serotype5 vector and hfNPC as described below. Four doses were administered with a spacing of approximately twelve hours between each dosing for the first three days, with the fifth dose being administered on the morning of the fifth day. MPTP was handled using published guidelines<sup>349</sup>. Behavioral scoring before and after MPTP injection confirmed a moderately low PFS (non-parkinsonian) for all animals selected

indicative of low to moderate pathological deficit. These animals were selected based on their likelihood to harbor a relatively intact NS unit, while having been previously exposed to pathological insult and related toxic microenvironment.

**4.6.11 Fluorogold Injection:** Ten to twelve days prior to sacrifice, previous striatal targets used for GDNF delivery were injected bilaterally with 10 ul of the retrograde transport tracer, Fluorogold to interpret the extent of donor-derived striatal integration in long-term animals. Methods used to deliver viral vector were identical for fluorogold.

**4.6.12 Sacrifice & Histological Preparation:** Animals were sacrificed according to IACUC guidelines and regulations after 1.5 months (short-term) [1 hNSC-WT, 1 hNSCeGFP] or 11 months (long-term) after grafting [1 hNSC-WT, 5 hNSC-eGFP, 2 hNSC-Kill]. Animals were deeply anesthetized with ketamine (10mg/kg, i.m.) and sodium pentobarbital (50mg/kg i.v.) until complete loss of corneal reflexes. Whole brains were collected following cardiac perfusion with cold heparinized saline and cold 4%PFA, then suspended in 4%PFA at 4°C for deep fixation. Following fixation, brains were allowed to “sink” in 30% sucrose preservative then transected at the mid-sagittal plane into two hemispheres and sectioned serially on a vibratome at 50um in the parasagittal plane to include the entire striatum and SN up through the extent of the lateral ventricle into the midline. Free-floating sections comprising every sixth section were stored in buffer at 4°C.

**4.6.13 Immunohistochemistry:** Free-floating sections were treated with standard IF techniques. Briefly, sections were incubated (non-surface, intracellular antigens) in 0.3%TritonX-100 (Sigma-Aldrich, St Louis, MO USA) for 20min, and blocked in 5% Normal Donkey, Horse, or Goat Serum (Jackson ImmunoResearch Laboratories or Vector Labs

Burlingame, CA) + 1% BSA (Sigma-Aldrich) in 0.3% TritonX-100 for 1hr at RT. Primary antibodies (see below) were diluted 1:1 (blocking solution: dPBS + 0.3% TritonX-100) and incubated for 24-72hrs at 4C. Primary antibody was omitted from specified controls. Sections were then washed thoroughly, followed by incubation with species appropriate secondary antibodies. For IF: fluorescence-labeled AlexaFlour conjugates; Donkey anti-mouse, rabbit, or goat 488 (green), 555 or 568 (near red), 647 (far-ir) (Molecular Probes, Eugene OR, 1:500). Where applicable, nuclei were visualized with Hoechst 33342 (Molecular Probes, 6 ug/ml) (blue). For immunoperoxidase staining: sections were treated by standard protocols using the Vectastain ABC-Elite kit with Nickel-DAB chromagen substrate enhancement. Secondary antibodies used were biotinylated horse anti-mouse IgG (1:500) (Vector, BA2000) and biotinylated horse anti-goat IgG (1:250) (Vector, BA9500). Sections were rinsed, mounted onto gel coated slides, and coverslipped in PVA DABCO mountant for Nickel-DAB or mounted on permafrost-coated slides and mounted in Vectashield (Vector Labs) for fluorescence.

**4.6.14 GFP-ir Graft Quantification:** One series (every sixth section) was immunolabeled for GFP with a mouse monoclonal or rabbit polyclonal antibody against GFP (Molecular Probes; 1:250-1:500) and treated and enhanced with the ABC+DAB method described above. We quantified the volume of tissue occupied by GFP immunoreactivity using the Aperio Scanscope XT (Aperio) at 40X magnification and analyzed for total GFP-ir area. Every 6<sup>th</sup> section was analyzed using the optical fractionator probe. The area containing concentrated GFP immunoreactivity was outlined at 20x magnification and GFP-ir cells were analyzed at 20-40x. Non-specific staining was extracted from the analysis by removing the highlighted area from the total area per image. The volume of spread of concentrated GFP immunoreactivity

was determined by multiplying the area of the outline by the individual section thickness and the sampling interval.

**4.6.15 Imaging and Processing:** Tissue sections and in-vitro samples were examined under either a standard light microscope or one of several fluorescence microscopes (Olympus IX71, Leica DMI400B) and images captured with a Hamamatsu-ORCA-ER C47/42 camera using ImagePro 6.2 or Metamorph 7.5.6. All confocal data was taken from a Radiance 2100/AGR-3Q BioRad Multiphoton Laser Point Scanning Confocal Microscope and acquired with Lasersharpe 2000 V6.0. Image J v1.38 was used to process original images in the following order: application of look-up-table (red or green), scale bar applied, smoothed with the 2-D or 3-D hybrid median filter, segmented with multi-thresholder (maximum entropy or mixture modeling methods), and then combined into a RGB image. Confocal z-stacks were arranged into 3D reconstructions and analyzed using Volocity 5 software. Figures were arranged in Adobe Photoshop or Illustrator (Adobe Systems, San Jose, CA).

**Primary Antibodies**

Rabbit-anti-Ki-67	(1:500)	(AB15580-100 AbCam)
Goat anti-PCNA	(1:200)	(SC-9857 Santa Cruz)
Mouse anti-Sox2	(1:200)	(MAB2018 R&D)
Goat anti-Vimentin	(1:200)	(AB1620 Chemicon)
Rabbit anti-GFAP	(1:500)	(Dako)
Rabbit anti-Nestin	(1:200)	(MAB5922 Chemicon)
Mouse anti-Nestin Human Specific	(1:200)	(MAB5326 Chemicon)
Mouse anti-Tuj-1	(1:500)	(MMS-435P Covance)
Mouse anti-Beta-III-tubulin	(1:250)	(MAB1637 Chemicon)
Mouse anti-NeuN	(1:100)	(MAB377 Millipore)
Goat anti-Doublecortin	(1:100)	(SC8066 SantaCruz)
Mouse anti-PSA-NCAM	(1:100)	(MAB5324 Chemicon)
Rabbit anti-BLBP	(1:300)	(AB9558 Chemicon)
Mouse anti-MAP2	(1:250)	(M4403 Sigma)
Mouse anti-Neurofilament (PAN)	(1:250)	(18-017 Invitrogen)
Mouse anti-TH	(1:5,000)	(MAB 318 Chemicon)
Rabbit anti-TH	(1:500)	(PRB-515P Covance)
Goat anti-GDNF Biotinylated	(1:500)	(BAF 212 / R&D)
Rabbit anti-GFP	(1:500)	(A11122 Molecular Probes)
Mouse anti-BrdU-488	(1:500)	(A21303 Molecular Probes)
Goat anti-Ang-1	(1:300)	(SC-6319 Santa Cruz)
Rabbit anti-Tie-2	(1:300)	(SC-324 Santa Cruz)
Rabbit anti-p-Tie-2	(1:300)	(SC-130607 Santa Cruz)
Mouse anti-Cd11b	(1:100)	(CD11b00 CALTAG Labs)

**Microscopes**

		Excitation	Dichroic	Emission
<b>Leica DMI4000B</b>	DAPI	320-410	400	430-500
	GFP	460-500	505	512-542
	RFP	534-558	565	560-640
<b>Olympus IX71</b>	DAPI	360-370	400	435-485
	GFP	470-490	500	510-560
	RFP	540-580	595	600-660

**Radiance 2100 Confocal**

Collected wavelengths: 520-560

520 Long Pass = Block filter

560 Short Pass = Emission filter

Collected wavelengths: 580-700

580 Long Pass = Block filter

Open = Emission filter

Photomultiplier Tubes  
 488 channel, 100%  
 568 channel, 100%  
 No mixing between PMTs

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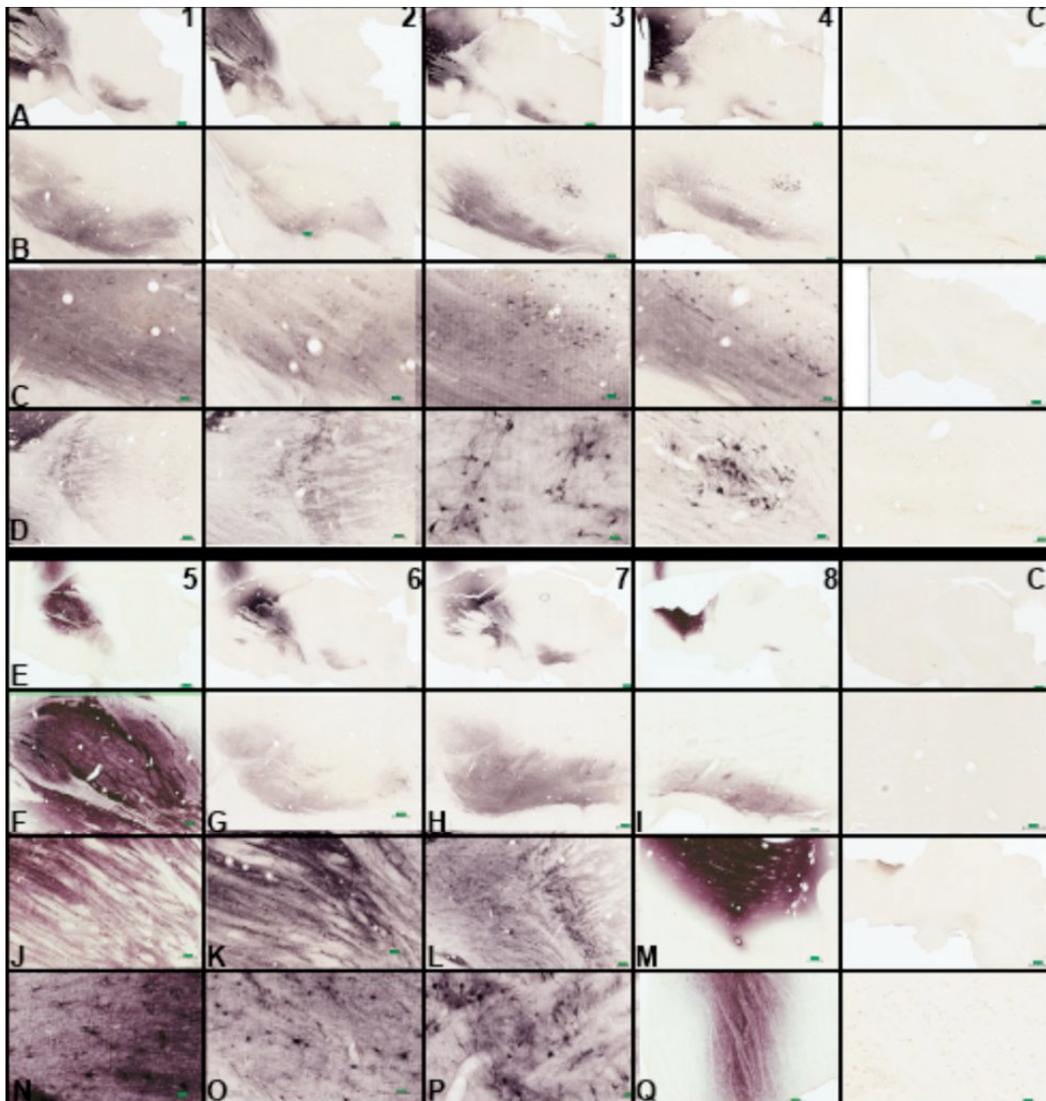
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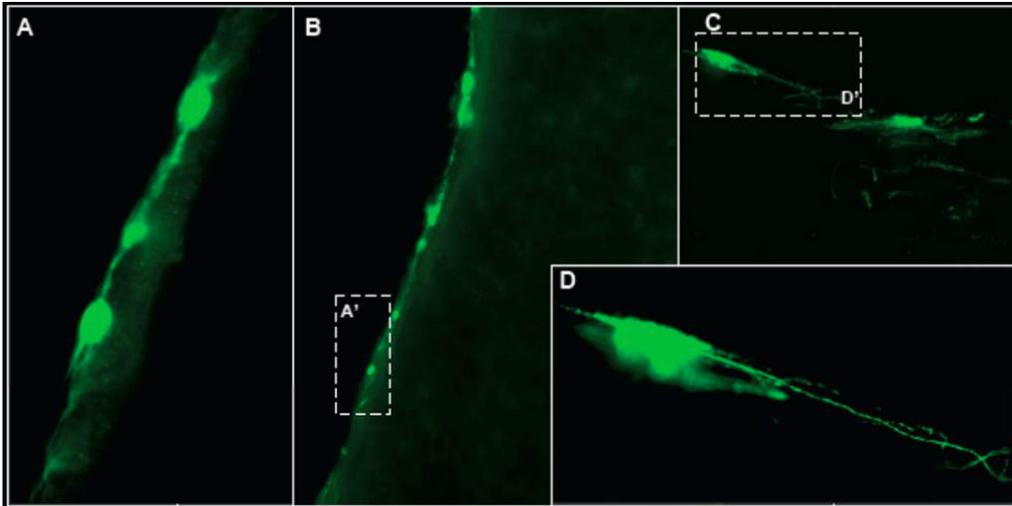
#### 4.9 Supplementary Information:

**Supplementary Table 4-S1.** Experimental subjects and treatment guide.

<b>Monkey</b>	<b>hfNSC subtype</b>	<b>Reporter</b>	<b>FG</b>	<b>Sacrifice</b>
W843	2050-NS-eGFP	GFP	NO FG	1.5 Months
W820	2050-WT-3BIC-3.3	NO GFP	NO FG	1.5 Months
W883	2050-WT-3BIC-3.3	NO GFP	NO FG	11 Months
X955	HFB-2050-WT-DEAD	NO GFP	FG	11 Months
X850	HFB-2050-eGFP-DEAD	GFP	FG	11 Months
W841	2050-p32-eGFP	GFP	FG	11 Months
X896	2050-3BIC-3.2-GFP	GFP	FG	11 Months
Y092	2050-3BIC-3.2-GFP	GFP	FG	11 Months
X740	2050-NS-eGFP	GFP	FG	11 Months
X577	2050-p32-eGFP	GFP	FG	11 Months



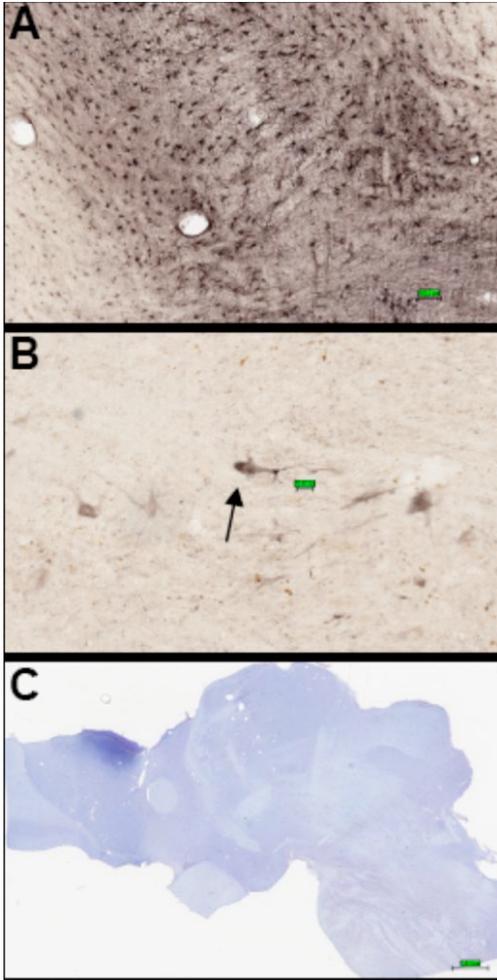
**Supplementary Figure 4-S1. GDNF expression at multiple anatomical levels 1.5 months post-injection.** Robust exogenous GDNF was detected throughout the striatum [Columns 1-8 A,E; Columns 1,2, row D; J, K, M, and Q] and ipsilateral SN (Columns 1-4, rows B and C; Column 3, row D, Columns 6-8, rows G-I] throughout multiple anatomical planes within the same animal [Columns 1-8], but never on the contralateral side [Column C]. Robust GDNF-ir cells were also found near the locus coeruleus [Column 4, row D] and dorsal extents of the internal capsule. [L, P]. Most GDNF-transduced striatal cells had a morphology indicative of medium spiny projection neurons [N, O]



**Supplementary Figure 4-S2. Incorporation of GFP-ir hfNPC into host ventricular wall.** In one animal, residual donor cells (green) retracted dorsally through the needle tract into the ventricle. Some of the cells which engrafted near the ventricle incorporated into the lumen [A-D] and appeared to retain morphological characteristics consistent with the SVZ-NSC niche.

**Supplementary Table 4-S2.** GFP-ir whole-graft quantification in the four animals that received successful hfNPC transplants (0-5, with 5 being the highest rank).

	<b>X577L</b>	<b>X740L</b>	<b>Y092R</b>	<b>X896R</b>
<b>GFP-ir Rank</b>	5	5	4	2
<b>SN (mm<sup>2</sup>)</b>	9.834	13.341	8.674	7.108
<b>Needle(mm<sup>2</sup>)</b>	22.704	28.186	21.254	0
<b>Area (mm<sup>2</sup>)</b>	32.538	41.527	29.928	7.108
<b>Volume (mm<sup>3</sup>)</b>	1.6269	2.07635	1.4964	0.3554



**Supplementary Figure 4-S3. Fluorogold (FG) and Nissl at 11-months post-transplantation.** Striatal tissue adjacent to injection sites displayed robust FG-ir cells [A]; however, few FG-ir were found in the corresponding SN [B], labeling at best 5% of the spared endogenous TH-ir population. No FG-ir/GFP-ir cells were detected after 11-months suggesting non-innervation of the striatum at this time-point. [C] A representative example of Nissl staining demonstrates that hfNPC grafts do not induce aberrant pathology or alter host neuronal organization.

## CHAPTER 5:

### CONCLUSIONS AND FUTURE DIRECTIONS

#### 5.1 Summary

This dissertation describes a clinically relevant paradigm for hNSC transplantation in the dopamine-depleted non-human primate brain. Several conclusions can be ascertained from this data. Contrary to previous reports utilizing suspension culturing conditions, human fetal subventricular-zone derived NSC can be maintained as multilayer-adherent-networks and expanded long-term (at least three years) without senescence or karyotypic aberration, in sufficient numbers to treat multiple patients over time. Previous groups have reported mass senescence in fetal cortical derived neural stem cells after just sixty passages, indicating that these cells would not have the capacity to be expanded indefinitely to treat multiple patients long-term. Furthermore, Clive Svendsen's group has shown that these cortical progenitors also accumulate chromosomal abnormalities, specifically chromosomes in 7 and 19, when cultured long-term. I have managed to overcome both of these obstacles utilizing the novel multilayer adherent network culturing system, thus providing a clinically relevant neuroectoderm derived undifferentiated fetal NSC population.

Transplantation of these undifferentiated fetal neural stem cells into rodent and non-human primates indicates they may retain developmentally appropriate cues and respond to local trophic and guidance molecules in vivo. Specifically, I have shown recovery in behavioral indices of parkinsonism in MPTP-lesioned vervet monkeys as well as long-term survival and axonal outgrowth up to eleven months post-injection when transplanted concomitantly with exogenous GDNF. Prior to these studies, it was unknown whether

undifferentiated human fetal NSC would even engraft into the CNS, let alone incorporate within the dopamine-depleted primate brain and provide functional improvement in severely lesioned animals. Previous reports by Svendsen's group indicated that fetal NSC do not migrate or significantly differentiate when transplanted striatally into parkinsonian rodent models. In stark contrast, we found extensive migration of fetal NSC derived from the subventricular zone of the lateral ventricles when transplanted into the striatum. Interestingly, migration was never found when cells were transplanted into the substantia nigra, indicating regionally specific cues are present within the striatum to induce not only migration but also glial differentiation of donor cells into astrocytes.

Previous reports have suggested that hfNSC do not differentiate in the primate brain, specifically when induced to express GDNF. In contrast, development of GFP-labeled hfNSC allowed for the unequivocal determination of "monoamine-like" differentiated morphologies with extensive varicose projections. I found morphological evidence consistent with neuronal differentiation including branching dendrites with spine-like structures as well as elaborately ramified fine-beaded structures. Unfortunately, standard biochemical analysis for mature neuronal marker proteins did not confirm the histological results that indicated extensive differentiation *in vivo*. It is possible that these grafted human cells simply need more time or more instruction *in vitro* or *in vivo* to assume appropriate biochemical indices of mature neuronal differentiation. Additional staining with 5-HT and GABA were inconclusive and unfortunately limited tissue did not permit for further testing.

Anatomically, the data were quite impressive, as historically, the biology of the NS unit is heavily rooted in morphological anatomical analysis (immunohistochemistry isn't always perfect and often depends on specific fixation techniques). Unfortunately, there appears to be a relative disconnect between what is found morphologically compared to biochemical readouts. Most classic neuropathologists would stop at this point and declare

these grafts as neuronally mature, however there appears to be a disconnect between anatomy and true molecular maturation, therefore, extreme caution should be warranted when making assumptions about the maturity of grafted cells in relation to their overall anatomical appearance. Either way, this is the first report to our knowledge of human fetal neural stem cells engrafting long-term and projecting extensive neuritic fibers within the normal or lesioned adult primate brain, and provides validation that human NSC preparations retain the capacity to undergo programs of early neuronal differentiation *in vivo*.

To determine if these morphologically relevant fibers are truly neuronal in function, a more thorough analysis of grafted cells is absolutely necessary. Specifically, electron microscopy would be ideal to determine if the “bouton-like” enlarged fusiform structures that highly resemble sites of synaptic neurotransmission are truly integrated and appear functionally active. If so, we would expect to find evidence for pre- and post-synaptic contact structures as well as vesicles carrying neurotransmitters at the pre-synaptic sites. In addition, determining neuronal functionality relies heavily on electrophysiological techniques to measure pre- and post-synaptic transmission. As neuronal precursors begin to mature, they develop the potential to carry “mini-spikes” of electrical activity and will eventually fire full-on action potentials as they mature into neurons. A combination of specific dopamine agonists/antagonists could be used to selectively inhibit or excite grafted cells and determine if they respond in a similar manner as dopamine neurons *in vivo*. Utilizing these techniques, we could determine what specific types of neuronal properties grafted cells retain, specifically, whether they are of an excitatory or inhibitory nature and if so, which specific types of neurotransmitters they have the capacity to respond to. Moreover, dopamine neurons, like many other mature neuronal sub-types, typically retain a characteristic set of electrophysiological properties that may distinguish them from other cells thus allowing for a more unambiguous determination of cellular properties within grafted cells (Reviewed by

Grace and Bunney in *Electrophysiological Properties of Midbrain Dopaminergic Neurons*, in *Neuropsychopharmacology* Volume 5).

Combined results appear to confirm previous studies indicating that SVZ-derived human fetal NSC lack the inherent potential to significantly differentiate into mature A9-DA neurons *in vivo*. In both transplantation paradigms, only 1-3% of the total grafted cells expressed tyrosine-hydroxylase after several months *in vivo*, indicating that even in the presence of neurotrophic stimuli known to induce dopaminergic neurogenesis (i.e., GDNF), SVZ-derived fetal NSC cannot be induced to differentiate preferentially into A9 dopamine neuron phenotypes, further indicating regional specificity within CNS derived stem cell preparations. These data are highly congruent with those reported by Ole Isacson's group, clearly showing that SVZ-derived neural progenitor cells do not retain the capacity for midbrain-specific neurogenesis, whereas ventral mesencephalic precursors readily express tyrosine hydroxylase and secrete dopamine *in vitro* and *in vivo*. One possibility is that these cells simply need more time to mature in the primate brain compared to the rodent brain, as described by Lars Olsen (personal communication) who has found congruent morphological data when transplanting similar cells into rodent substantia nigra. Another equally plausible explanation is that these cells are only capable of expressing midbrain-associated proteins, specifically tyrosine hydroxylase, transiently. The only way to accurately assess this question is through multiple time points to determine at what time grafted cells begin to express tyrosine hydroxylase and to what extent they maintain expression past 11-months *in-vivo*. In addition, it would be extremely beneficial to determine whether grafted cells that appear dopaminergic and co-express tyrosine hydroxylase actually make and secrete dopamine. While technical limitations do not allow for such biochemical analysis, future studies would benefit by techniques that identify the presence and/or absence of specific enzymes involved in catecholamine synthesis such as DOPA decarboxylase, dopamine-B-

hydroxylase, and phenylethanolamine N-methyl transferase to determine specifically which types of monoamine neurons may be present.

Interestingly, many NFM-ir neurites had morphological parameters consistent with developing axons and appeared polarized parallel to endogenous neural circuitry when grafted cells were in close proximity or overlapped, demonstrating for the first time that undifferentiated hfNSC retain developmentally relevant programs of early differentiation and may respond to host signals in the dopamine-depleted primate brain. As previously indicated, a full time-course with multiple short and long-term time-points is necessary to determine if eGFP-ir graft derived fibers are still elongating and extending into host tissue. Interestingly, staining for the growth cone associated protein (GAP-43) expressed at the distal tips of developing neurons was negative, once again indicating that biochemical and morphological differentiation status do not necessarily overlap.

In addition to severely Parkinsonian monkeys, I also examined a large number of hNSC-transplanted monkeys treated with lower doses of MPTP to induce significant (50-80%) depletion in the DA system, but few functional effects. (If functional impairments are not essential, we attempt to minimize incapacitating the monkeys and having to provide special care). These animals were observed and scored on the same behavioral rating system that also produces a sum factor "healthy behavior" analogous to the "activities of daily living" part of the UPDRS scale. It includes such items as walking about the cage, eating, drinking, looking out of the cage, threatening other monkeys, self-grooming, picking at the cage, etc. Monkeys decrease the amount of healthy behavior in response to many different types of disturbance, drugs, or illness, including MPTP-induced Parkinsonism.

Data from 42 monkeys were analyzed which had received hNSCs injected into the striatum and/or SN (not including the stage 4 animals reported in Chapter 4). Normal and healthy behaviors were not reduced by the surgery or the hNSC implantations suggesting that

the cells were exerting no behaviorally apparent toxicity. Rather, normal behaviors appeared to trend toward and increase in these animals and no abnormal behaviors such as dyskinesia or dystonia were seen. Although, with these monkeys, we cannot infer any functional improvement associated with the NSCs, at least they appear to do no harm or cause any apparent behavioral toxicity. The pathological results of the monkeys reported here demonstrate that no tumors or overgrowth has ever been seen under any stem cell transplantation paradigm we have implemented with these cells. These results are paramount for safety and efficacy of future clinical trials.

One of the most difficult obstacles to overcome with any research involving large animals, specifically in this case, non-human primates, lies in attaining a large enough experimental group of animals to attain significance and control for all experimental variables. In this case several historical controls were necessary to compare GDNF-injected animals. Ideally, each experimental group would be assigned the same number of replicate animals, however, in this case, expenses and ethical considerations did not allow for a complete set of experimental controls. Specifically, no comparisons could be made between animals injected with neural stem cells alone and those co-injected with GDNF as the original hfNSC were not engineered to express the GFP-reporter. As a result, we have no way to accurately compare the effects of exogenous striatal GDNF on total graft survival or the degree to which GDNF enhanced neuritic fiber outgrowth in comparison to transplanted cells alone without exogenous growth factor support.

## **5.2 Conclusions and Future Directions**

Significant progress has been made towards testing the hypotheses of the specific aims, showing survival, migration, and conversion to the DAergic phenotype in the nigrostriatal system of normal and diseased monkeys, normalizing effects on host brain

morphology, as well as reversal of parkinsonism induced by MPTP through multiple homeostatic mechanisms. Specifically, we have shown survival of hfNSCs after implantation into normal and DA-depleted monkey striatum, including differentiation into TH-expressing cells and other neural cell types using multiple methods for identifying hNSCs and their progeny. We found functional improvements in severely affected MPTP-parkinsonian monkeys compared with sham-operated monkeys. We also found that hNSCs "normalized" host systems by returning them toward normal cell numbers, cytoarchitecture, size, and function.

Experiments during this dissertation have supported the original hypothesis that human neural stem cells can reverse dysfunction in the neurotoxic model of parkinsonism produced by MPTP in monkeys and suggested that a combination of endogenous effects and cellular replacement may be responsible for functional improvements in severely parkinsonian monkeys. The effects we have observed appear to fall into 2 broad categories (1) the impact the abnormal milieu has upon the stem cell (e.g., shifting its differentiation fate and neuritic output along the nigrostriatal pathway) & (2) the stem cell's impact upon the host (e.g., providing neurotrophic & neuroprotective support to endogenous dopaminergic (DA) neurons & their projections, restoring homeostasis to the cytoarchitecture including the size-to-number ratios of endogenous cells, preserving compromised neural circuits, and possibly "circuit restoration" achieved by a small percentage of stem cells that acquire characteristics of dopamine neurons. However, significant questions remain as to how these functional effects occur and, more importantly, can be enhanced.

While the recovery is very impressive and without apparent toxicity or dyskinesias, it is not yet complete and few hNSC derived dopaminergic cells survived, even after GDNF enhancement for eleven months. We believe that the multiple effects of stem cells on the nigrostriatal system are interesting and should be further investigated, specifically the

relationship between transplanted cells and the specific axonal guidance molecules along spared endogenous circuitry. Additional studies confirming these results should also include a TH-promoter driven reporter construct integrated into donor hfNSC to verify the potential of these cells to express tyrosine hydroxylase and determine to what extent the TH-ir donor-derived cells express A9-DA neuron specific *Girk2* at time points beyond 1-year. Importantly, there was evidence for transgene silencing of eGFP in larger TH-ir cells that retained morphological features consistent with endogenous mature dopamine neurons (large cell body and axonal arborization). The data indicate that we still may not be appropriately accessing the actual dopaminergic potential of grafted cells and further argue for a more detailed analysis before a definitive conclusion can be made. In addition, a side-by-side comparison with fetal DA-neuroblasts as well as embryonic stem cell derived and induced pluripotent stem (iPS) derived NSC and their DAergic derivatives to determine which tissue source provides the most therapeutic cellular substrate is essential to the advancement of cellular transplantation based therapies.

Most prior studies have focused on the concept that the *host* environment — as it changes during the course of development and aging, or after injury or degeneration — influences the transplanted stem cell, as exemplified here by the emergence of some donor-derived TH+ and DAT+ cells. Based upon past reports of the small difference in DA depletion that appears to underlie large functional differences, even a 5% elevation in DA may be sufficient to produce significant functional improvement, either from stem cell-derived DA neurons or from augmentation of host DA pathways via trophic/neuroprotective effects. We believe our data suggest that the primate CNS may benefit (1) from human stem cell-derived replacement of missing cell types (including neurons) and (2) from the supportive effects of other stem cell-derived progeny that may be necessary for promoting optimal recovery via the

variety of mechanisms described in this dissertation (trophic, homeostatic and guidance) as well as others that remain to be elucidated.

Given the analysis we have performed utilizing undifferentiated human fetal NSC in the non-human primate brain, and the extensive knowledge our colleagues and others have attained transplanting fetal DAergic tissue, it is still relatively unclear which neural substrate may provide the most therapeutic substrate to restore DAergic tone in Parkinson's disease. While NSC have the advantage of being expandable nearly indefinitely, DAergic neuroblasts appear to functionally integrate the striatum and developmentally reconstruct the nigrostriatal pathway in response to GDNF chemotaxis in the rodent and monkey. Unfortunately, fetal grafts may also induce dyskinesias in patients over time given their apparent vigor and sustained uncontrolled production of dopamine, therefore, a more slowly developing substrate may, with time, eventually be a more therapeutic option clinically.

The degree of degeneration within the patient may also have a dramatic impact on how neural grafts survive and respond in vivo. Currently, neurologist are only able to make a correct diagnosis for Parkinson's disease based on motor deficits (Tremor, Rigidity, Akinesia, Postural Instability). These characteristic motor behaviors do not become clinically diagnosable until greater than 80% of the dopaminergic tone has been lost, therefore a typical Parkinson's disease patient may not have enough time to wait for a NSC graft to mature and develop into relevant cell types in-vivo. When diagnostic tools have improved and biomarkers are available to diagnose and treat Parkinson's disease earlier, before massive degeneration has already taken place, perhaps NSC therapies aimed at neuroprotection can provide therapeutic value for endogenous dopamine neurons. Therefore, under current clinical circumstances, the data described here argue against the use of undifferentiated hfNSC as a therapeutic option for cell replacement in Parkinson's disease. Specifically, simple neuralization (in this case by natural development in vivo) of stem cells does not appear to be

sufficient for in-vivo differentiation into dopaminergic cells. Rather, neural stem cells will most likely require further developmental cues, maturation, and specification into DA precursors in vitro to attain neural replacement. An intriguing question remains as to whether all undifferentiated neural stem cell preparations maintain similar characteristics in vivo.

Future studies will be necessary to determine exactly where the differentiation “sweet-spot” lies for creating safe, transplantable cells with high survival and sustained controlled release of dopamine in-vivo. A useful approach might be to systematically determine the phenotypic characteristics of ventral mesencephalic cells that have previously been shown to reduce behavioral deficits in parkinsonian primates pre-clinically. It is known that only a small portion of these cells are responsible for motor recovery, whereas others are expected to be involved in deleterious mismatch synapses inducing dyskinetic behaviors. Developing techniques to mimic phenotypic characteristics of the beneficial cellular components of these grafts may ultimately reshape how we prepare and differentiate cells in vitro for further optimization of a transplantable therapeutically relevant neural substrate capable of restoring dopaminergic tone in Parkinson’s disease patients. With the advent of human embryonic stem cells, iPS derived neural cells, and most recently induced neural (iN) cells, we are now equipped with more tools than ever before to create and test neural transplantation therapies for PD, however it should be noted that extreme caution is warranted for any therapy utilizing undifferentiated, non-midbrain specified neural stem cells.

## APPENDIX A:

### CULTURE SYSTEM FOR RODENT AND HUMAN OLIGODENDROCYTE SPECIFICATION, LINEAGE PROGRESSION, AND MATURATION

#### Summary:

Here we document protocols for the production, isolation, and maintenance of the oligodendrocyte phenotype from rodent and human neural stem cells. Our unique method relies on a series of chemically defined media, specifically designed and carefully characterized for each developmental stage of oligodendrocytes as they advance from oligodendrocyte progenitors to mature, myelinating oligodendrocytes

#### Introduction:

Here we document a protocol for the production, isolation and maintenance of the oligodendrocyte (OL) phenotype from rodent and human neural stem cells (NSC). Our unique method relies on a *series of chemically defined media*, specifically designed, and carefully *characterized for each developmental stage* of OL, as they advance from OL progenitors (OLP) to mature myelinating OL [Figure A-1] (Neman and de Vellis, 2008). Providing hNSC with the nutrients specifically required at a particular moment in OL development, our system allows for *the propagation of OL at a desired stage* from OLP to mature premyelinating OL. Therefore, lineage progression can be manipulated by controlling the duration of a given developmental stage as needed, in a more “natural” manner, and without using gene transfer (Park et al., 2002b; Kim, 2004; Müller et al., 2006; Ahn et al., 2008), co-cultures, or undefined substrates such as, cell line-derived conditioned medium or animal serum.



## I. RODENT NSC ISOLATION AND MAINTENANCE

### *Materials:*

Remarks: All dissection instruments, plastic, and glassware must be sterile.

Animals: Time-pregnant embryonic day 14-16 (ED14-16) Sprague-Dawley rats.

(Charles- River, Wilmington, MA)

Dissection Instruments: Refer to numbers in [Diagram 1].

- |   |                                |
|---|--------------------------------|
| 1) Mayo Scissors                                  | 14010-17 FST                   |
| 2) Lister Scissors                                | 14131-14 FST                   |
| 3) Blunt-Pointed Forceps                          | 08-887 Fisher                  |
| 4) Iris Scissors                                  | 14060-09 FST                   |
| 5) Moma Iris Forceps                              | 11373-12 FST                   |
| 6) Dumont #7 Forceps                              | 11297-10 FST                   |
| 7) 140 $\mu\text{m}$ and 230 $\mu\text{m}$ sieves | Collector, E-C Apparatus Corp. |
| 8) Syringe  | 520673 Jone Kendall            |

Bovine Serum Albumin (BSA)

Conical tubes

Flasks (T-12.5  $\text{cm}^2$ , T-75  $\text{cm}^2$ )

Hank's Balanced Salt Solution (HBSS)

Hemocytometer

Isoflurane

Petri Dish (Bacterial grade, non TC-treated)

PSA-NCAM Coated Dishes (See Support Protocol 1)

Sterile surgical gauze

STMc media

14-gauge Penetration needle, 14-825-15 Fisher

### **Step 1. Isolation of Neural Stem Cells from Newborn Rat Brain**

#### *Procedure:*

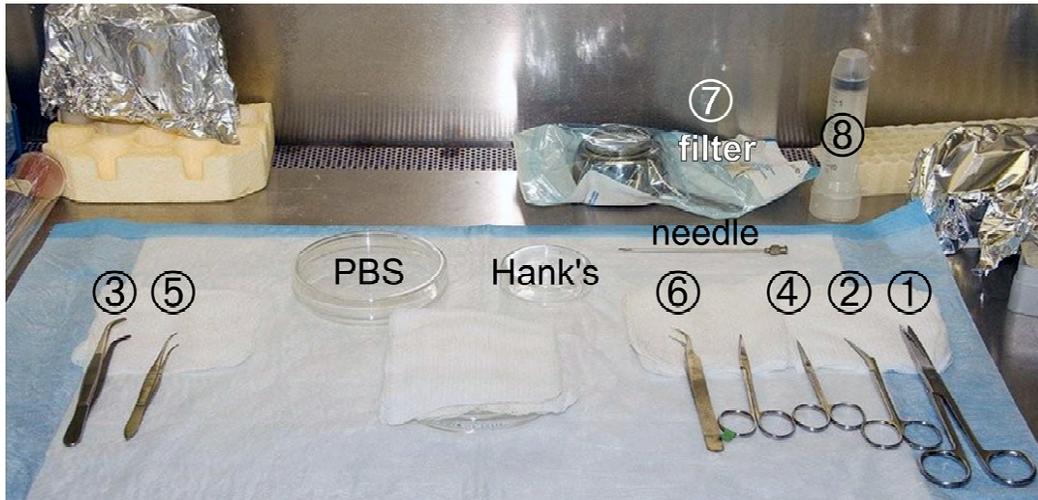
1. Prepare the work area and sterile tools in a biosafety hood [Diagram 1].
2. Euthanize the rat by isofluorane inhalation.
3. Extract the placenta containing embryos and place in STM Basal + 1% BSA.
4. Remove the cerebellum.
5. Dissect the brain of each embryo and place in STM complete (STMc) medium.
6. Separate the cortex from the rest of the brain and remove the meninges with forceps.
7. \*Combine the tissue of all the brains without meninges and mechanically dissociate with the needle by gently aspirating the brain pieces (10 times) and releasing the suspension slowly with the needle attached to the wall of the tube (try to minimize foaming).
8. \*Recover the supernatant with the cells in suspension and transfer to a 15ml tube.
9. \*Add 2-4ml of complete medium to the chunks left over in the dissociation tube and dissociate again 5 to 8 times.
10. Filter the suspension of dissociated cells through the 230  $\mu\text{m}$  and 140  $\mu\text{m}$  sieves to remove cell clusters.
11. Rinse the sieves sequentially with basal STM + 1% BSA at room temperature, and add this medium to the tubes containing the cells.

12. Collect the cells by centrifugation at 45 RCF for 8 min.
13. Gently discard the supernatant.
14. Resuspend the cell pellets in 4 ml of STM complete (freshly prepared), and gently dissociate the pellets.
15. Assess cell viability (see next section), count cells using hemacytometer, and plate onto fresh PSA/NCAM coated dishes ( $2 \times 10^6$  cells/100 mm dish).
16. Incubate plated cells overnight at 37°C with 4.5% CO<sub>2</sub> and 95% humidity. *Note:* Younger cells do not yet express PSA-NCAM and will remain floating as small clusters, whereas the older cells will attach overnight.
17. \*\*On the next day, recover the non-attached cells, pellet as previously described, and remove and save the conditioned media (CM).
18. Resuspend the pellet and dissociate in 4 ml of fresh STMc medium by passing through a 14-gauge needle eight times. Complete the volume to 8ml with conditioned medium and plate on additional anti-PSA-NCAM coated plates. *Alternatively:* Place 1.0 ml of the cell suspension in a 75ml Erlenmeyer [EM] flask in 25ml of STMc, and incubate with shaking at 37°C [Figure A-2].
19. Feed cells every other day by removing 1/3 of the culture medium and adding the same volume of fresh STMc.
20. Switch the cells from 4.5% to 5.0% CO<sub>2</sub> only if your cells are growing slowly. Leave them at 4.5% if they maintain the color of the medium red/orange.

**Notes:** A chart of the STM cell isolation method is illustrated in [Figure A-2].

\*Steps 7-9 can be substituted by dissociating the cells 2.5 min using a Stomacher 80 (Seward, London, UK).

\*\*Collect, filter, and save the conditioned medium at 4°C for immediate use (or frozen for later use). CM is an excellent supplement to start NSC cultures from frozen stocks. Allow cells to grow to 70-90% confluency.



**Diagram A-1.** Instruments required for dissection. #1, Scissors for decapitation; #2,scissors to cut the head skin to expose the skull; #3, forceps to hold the head in place as you cut the skin and cut the skull cartilage with scissors, #4 to expose the brain. Some users prefer the curved scissors, #2 instead of #4. Use the same scissors to transfer the brain to the Petri dish containing PBS. #5 & #6, Forceps to hold the brain in place and remove the meninges (respectively). After removal of the meninges place the brains in Hanks solution while dissecting the rest of the brains. #7, filter mesh to filter the cell suspension after dissociation. #8, 20ml sterile syringe and sterile dissociation needle.

### Cell Viability Assay

Cell viability can be determined with the SYTOX blue nucleic acid stain (Molecular Probes, Eugene, OR). Cells with compromised plasma membranes are labeled by SYTOX binding to nucleic acids and detected by fluorometry.

*Materials:*

PBS

SYTOX blue nucleic acid stain

Tris-Buffered Saline (TBS)

*Procedure:*

1. Harvest and wash cells with 1× TBS
2. Incubate for 12 min in 1 μM SYTOX in PBS
3. Remove the solution and wash the cells with 1× TBS, five times
4. Determine the *number of positive cells per random field* and record as a percentage of the total number of cells.

## **Step 2. NSC Propagation and Maintenance**

### ***Propagation of Rodent NSCs as 2D-cultures***

NSCs can be propagated in bi-dimensional (2D) or three-dimensional (3D) cultures. When attached, NSCs (2D cultures) tend to grow faster and are therefore ideal to create a large cell stock quickly before starting specific studies. In addition, we have developed a new method for expansion and maintenance of human NSC in NB-B27 (see Media Preparation) as multilayer adherent network (MAN) cultures, with increased proliferation rates compared to standard sphere forming assays (Wakeman et al., 2009b). In order to accommodate for the difference in basal media, human NSC can either be initially derived in STMc media (replacing NB-B27), or previously established cultures may be slowly transitioned away from

the basal NB-B27. Simply substitute 25% STMc for 1 week, followed by successive weeks at 50%, 75%, and finally 100% STMc after one month.

*Materials:*

Cell Scraper

Conical tubes

Flasks (T-12.5 cm<sup>2</sup>, T-75 cm<sup>2</sup>)

Hank's Balanced Salt Solution (HBSS)

Petri Dish (Bacterial grade, non TC-treated)

PSA-NCAM Coated Dishes (See Support Protocol 1)

STMc media

*Procedure:*

1. When confluency has been reached, remove the supernatant conditioned media (CM), add 5ml of Hank's Buffered Salt Solution (HBSS) without Ca<sup>2+</sup> or Mg<sup>2+</sup>, detach the cells with a cell scraper, and transfer to a 15ml conical tube (1 to 3 Petri dishes).
2. Rinse the Petri dish once with 2ml HBSS and centrifuge at 45 RCF for 8min.
3. Resuspend the cell pellet in 3 ml STMc, dissociate gently using a 14-gauge needle, and centrifuge at 45 RCF for 5min.
4. Save the CM (as above), resuspend cells in fresh STMc + CM (2:1), and re-plate. If you have repeated this process several times and the cell pellet is 0.5ml volume or larger, divide the cell suspension into two parts.

5. One part of the suspension will be used to start a frozen stock (see next section for cryopreservation). The second half of the cell suspension is further dissociated using a needle (as described above, however, the sieves are not necessary).
6. Count the number of cells/ml and adjust the volume (to 15ml) with freshly prepared STM containing 1/3 self-conditioned-STM + 2/3 of fresh STMc.
7. Plate the cells in five T-12.5 cm<sup>2</sup> cell culture flasks coated with anti-PSA/NCAM as described.
8. Feed cells with 1/3 of self-conditioned-STM + 2/3 of fresh STMc every other day until they reach 80 to 90% confluency, and repeat the process to increase the number of cells.
9. When four or more T-12.5 cm<sup>2</sup> flasks reach confluency, harvest the cells as described above, and seed the equivalent content of cells from three T-12.5 cm<sup>2</sup> flasks into one T-75 cm<sup>2</sup> flask (coated with anti-PSA/NCAM). Feed the cells with 1/3 self-conditioned-STM + 2/3 of fresh STMc in a total volume of 10ml/ flask.
10. \*After 1-2 days, the culture medium should be red. If the medium is turning orange, add 3ml of STMc and repeat this step as needed. Add 3ml of STMc every day only if the medium changes color.
11. \*\*When cells reach confluency, you may elect to freeze the contents of one T-75 cm<sup>2</sup> flask (see cryopreservation section).
12. After accumulating at least 10 to 15 vials of cryopreserved NSC in a frozen stock, NSCs may be grown as neurospheres (3D-cultures) for slower growth, allowing more time to devote to experiments.

**Notes:**

\* If cells seem not to grow but look healthy, or if the culture medium is not red but rather purple, you will need to remove 1/2 of the plating medium and complete to 10ml volume of the mixture (1/3 self-conditioned-STM + 2/3 of fresh STMc freshly prepared). If the opposite is true and the culture medium turns orange overnight, the cells have proliferated heavily, and you will need to replace the culture medium and seed more T-75 cm<sup>2</sup> flasks (per three T-12.5 cm<sup>2</sup>).

\*\* When propagating cells to create frozen stocks, we strongly recommend maintaining a “mother flask/dish” by scraping most, but not all of the cells attached to the flask. After removing the detached cells, feed the mother flask with fresh medium and CM (1:1) to ensure continuity of these cultures (in case re-plated cells do not look healthy, grow slowly or die).

### ***Formation, propagation and maintenance of Neurospheres (3D cultures)***

Suspension aggregate, or “neurosphere”, 3D cultures are an alternate strategy to propagate NSCs at a slower pace than attached cells, while preserving most of the standard characteristics of a proper NSC. NSC suspension cultures are started from freshly dissociated NSC (2D cultures) and grown in Erlenmeyer flasks (EM) to prevent attachment and encourage free-floating spherical growth.

#### ***Materials:***

Conical tubes

Flasks (T-12.5, 75 cm<sup>2</sup>)

Glass Erlenmeyer flasks (EM) 25 ml or 50 ml with cap.

Hank's Balanced Salt Solution (HBSS)

Petri Dish (Bacterial grade, non TC-treated)

PSA-NCAM Coated Dishes (See Support Protocol 1)

STMc media

*Procedure:*

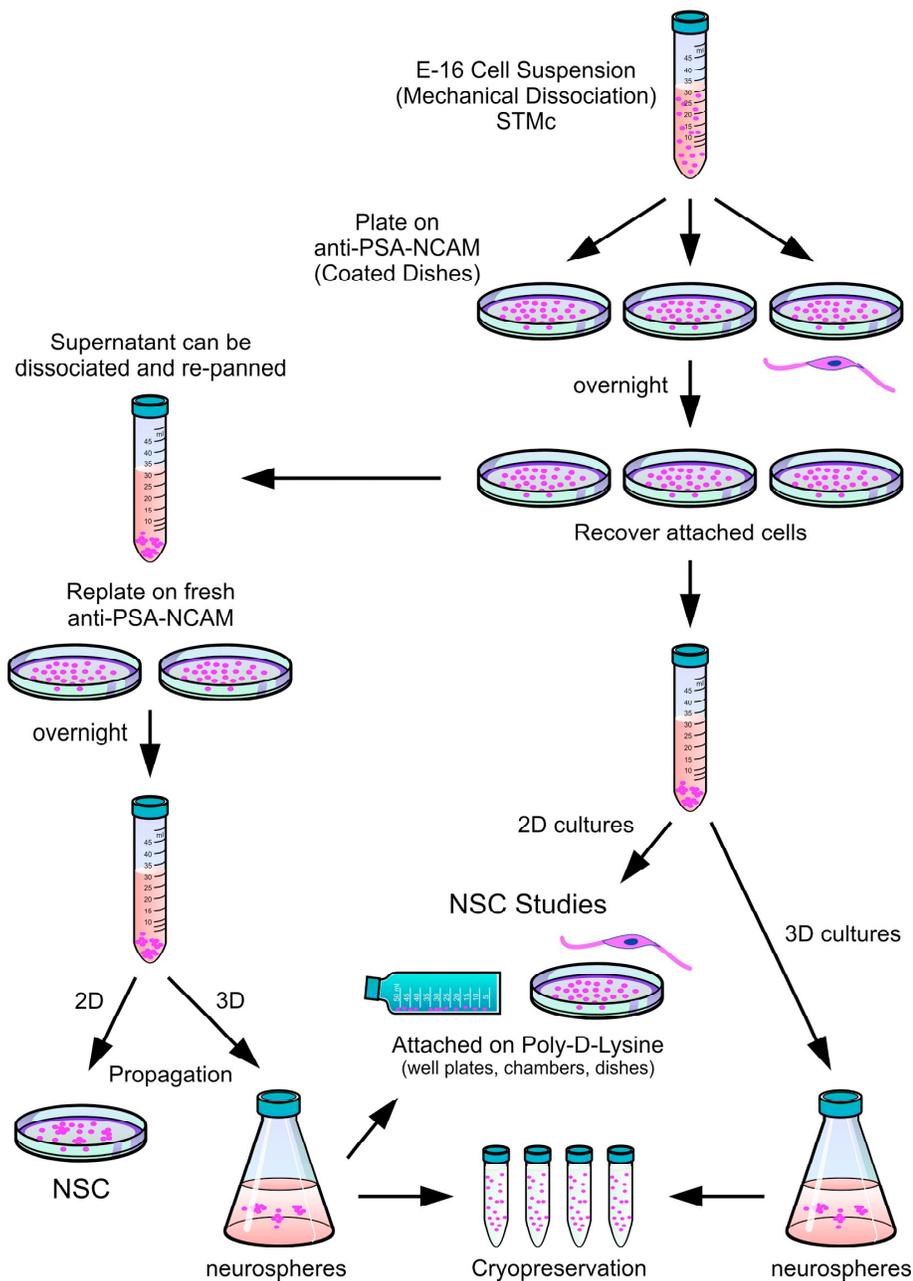
1. Place 15ml of fresh STM + 3ml of self-conditioned medium (CM) into the EM.
2. Add 2ml of the cell suspension (freshly harvested from 2D cultures), and close the top of the flask partially to allow for O<sub>2</sub>/CO<sub>2</sub> exchange.
3. Place the EM, continuously shaking, at 90 rpm in the incubator. If placing a shaker in the incubator is not an option due to safety regulations, the cell suspension may be placed directly into two Petri dishes (bacterial grade) to prevent cell attachment.
4. Add 1.5ml fresh STMc, every other day, and dissociate routinely (3 times gently) with a syringe and needle (as described for pellet dissociation in the previous section) to keep the spheres at a small size. This process allows for increased sphere formation without the negative potential for spontaneous differentiation. It also permits cells more exposure to the fresh nutrients in the culture medium, allowing for preservation of "stemness" in all cells.
5. When the culture medium turns orange overnight, it is time to renew the culture medium and split the cells. Collect the contents of the EM flask and place into one 50ml conical tube. Centrifuge for 6 min at 45 RCF.
6. Slowly collect the CM, filter (0.22 μm), and save for replating cells.

7. Resuspend the pellet in 4ml of STM to dissociate larger spheres, \*(at this point they should all be easily dissociated). Add freshly prepared STMc + 1/3 self-conditioned medium.
8. Seed cells on desired containers for experiments, or continue to propagate NSCs as 2D or 3D cultures.

**Notes:** A diagram for the propagation of NSC in 2D & 3D cultures is shown in [Fig. A-2].

\*If some spheres remain large in spite of repeated dissociation, use the sieves to eliminate the clumps instead of drastically dissociating them. This step will prevent significant cell death at the time of re-plating.

## Rodent Neural Stem Cell Preparation



**Figure A-2. Rodent Neural Stem Cell Preparation.** Following dissection, the cell suspension is plated on anti-PSA/NCAM antibody coated dishes and allowed to adhere. The process can be performed repeatedly to increase the numbers of neural stem cells, as 2D or 3D “sphere” cultures. (shown in the left side of the diagram). Alternatively, cells can be propagated and immediately used for cell culture experiments (as shown on the right side of the diagram). While we prefer to use committed OL-progenitors for cell transplants, other investigators also use uncommitted progenitors for grafting.

**Support Protocol 1:****Immunopanning**

The following method was developed based on published work (Wysocki and Sato, 1978 and Williams and Gard, 1997) to isolate the rodent NSC population from the other cell populations in the brain during initial plating. We also use anti-PSA/NCAM coated dishes to propagate rodent NSC in 2D cultures (Espinosa et al., 2002). Please refer elsewhere for specific methods on the selection of human NSC during primary derivation (Wakeman et al., 2009a).

*Materials:*

Anti-PSA-NCAM

Bovine Serum Albumin (BSA)

Flasks (T-12.5 cm<sup>2</sup>, T-75 cm<sup>2</sup>)

PBS

Petri Dish (Bacterial grade, non TC-treated)

Tris (pH 9.5)

*Procedure:*

1. Prepare the immunopanning cocktail by mixing: Tris (pH 9.5) +1% BSA + 50 µg/ml anti-PSA/NCAM.
2. Coat the bottom surface of non tissue-culture grade Petri dishes (351029) with the anti-PSA/NCAM mixture (4 to 5ml/per dish).
3. Incubate for 30 min at 37°C.
4. Wash Petri dishes 3 times with PBS and once with PBS + 1% BSA just before using.  
Do not allow the plates to dry.
5. Extra plates may be covered with foil and stored at 4°C for up to 10 days.

### ***Support Protocol 2***

#### **Cryopreservation / Thawing of NSC stocks.**

We recommend collecting cells for frozen stocks at low passage number. Human NSC are cryopreserved using modified methods found elsewhere (Wakeman et al., 2009a). In addition, the method formerly described for rat and mouse NSC (Espinosa et al., 2002) can also be used to stock human NSC.

##### ***A) Freezing NSCs***

###### ***Materials:***

Cryogenic slow-freezing chamber

Cryogenic freezing vials

HBSS

STM media

14-gauge needle

*Procedure:*

1. Allow NSCs to grow to 70-90% confluency. Remove all of the cell culture medium, add 5ml of HBSS to each Petri dish or 10ml to T-75 cm<sup>2</sup> flasks, and detach cells by gently scraping the culturing surface.
2. Centrifuge the cells at 45 RCF for 8 min and resuspend in 3 ml STM media.
3. Gently dissociate cells using a 14-gauge needle, pellet at 45 RCF for 8 min with a centrifuge, and discard the supernatant.
4. Gently resuspend the pellet from one (100mm Petri dish or T-75 flask) in 1ml of serum-free freezing medium.
5. Transfer the contents to a 1.2ml cryovial, and place the vial(s) in a cryogenic freezer container overnight for slow freezing.
6. Next day, place the vials in liquid nitrogen for long-term storage.

B) *Thawing NSCs*

*Materials:*

Anti-PSA-NCAM coated plates

CM and STM media

NSC Cryovial

*Procedure:*

1. To “reanimate” NSCs, defrost cryovials quickly in a 37°C water bath, and transfer the contents of the vial to a 2ml tube containing 1ml of STMc + CM at 37°C.
2. Centrifuge gently at 35 RCF for 5-7 min.
3. Remove the supernatant, add fresh plating medium (STMc + CM) and remove a small aliquot to test the initial cell viability (as described above).
4. Count the number of viable cells in the tube (approx.  $1 \times 10^6$  cells expected).
5. Plate cells onto anti-PSA/NCAM coated surface (Petri dishes or T-75 cm<sup>2</sup> flasks, plate the equivalent of 1 vial/T-75 cm<sup>2</sup> flask). If the yield is lower, utilize T-25 cm<sup>2</sup> flasks to increase the cell density necessary for healthy growth. Seeding low-density cultures in large containers decreases the proliferation rate and might be detrimental to the culture.
6. To propagate NSCs after replating, proceed as described in step 2 for “propagation and maintenance of NSCs”.

*Basic Protocol 2:***OL commitment in 2D and 3D cultures**

During development, the nutritional and environmental needs of cells change as they lose multipotency and become lineage restricted. The present system is based on the modification of nutrients contained in the cell culture medium and the percentage of CO<sub>2</sub> needed to optimize and direct lineage restriction towards the OL phenotype. Like NSCs, OLs can be propagated in 2D and 3D cultures. When attached (2D cultures), OLs grow

faster, and thus ideal to create an OLP cell stock quickly before starting specific *in vitro* cell culture or *in vivo* transplantation studies. A diagram of the following steps can be found in [Figure A-3].

*Materials:*

Media: STM-CM, OSM, GDM, OLDEM

Anti-IgM coated plates

Anti-PSA-NCAM coated plates

BFGF

Cell Scraper

Erlenmeyer (EM) flask

HBSS without  $\text{Ca}^{2+}$  or  $\text{Mg}^{2+}$

Petri dishes

14-gauge needle

15  $\mu\text{m}$  sieve

24-well plate

4.5%  $\text{CO}_2$  incubator

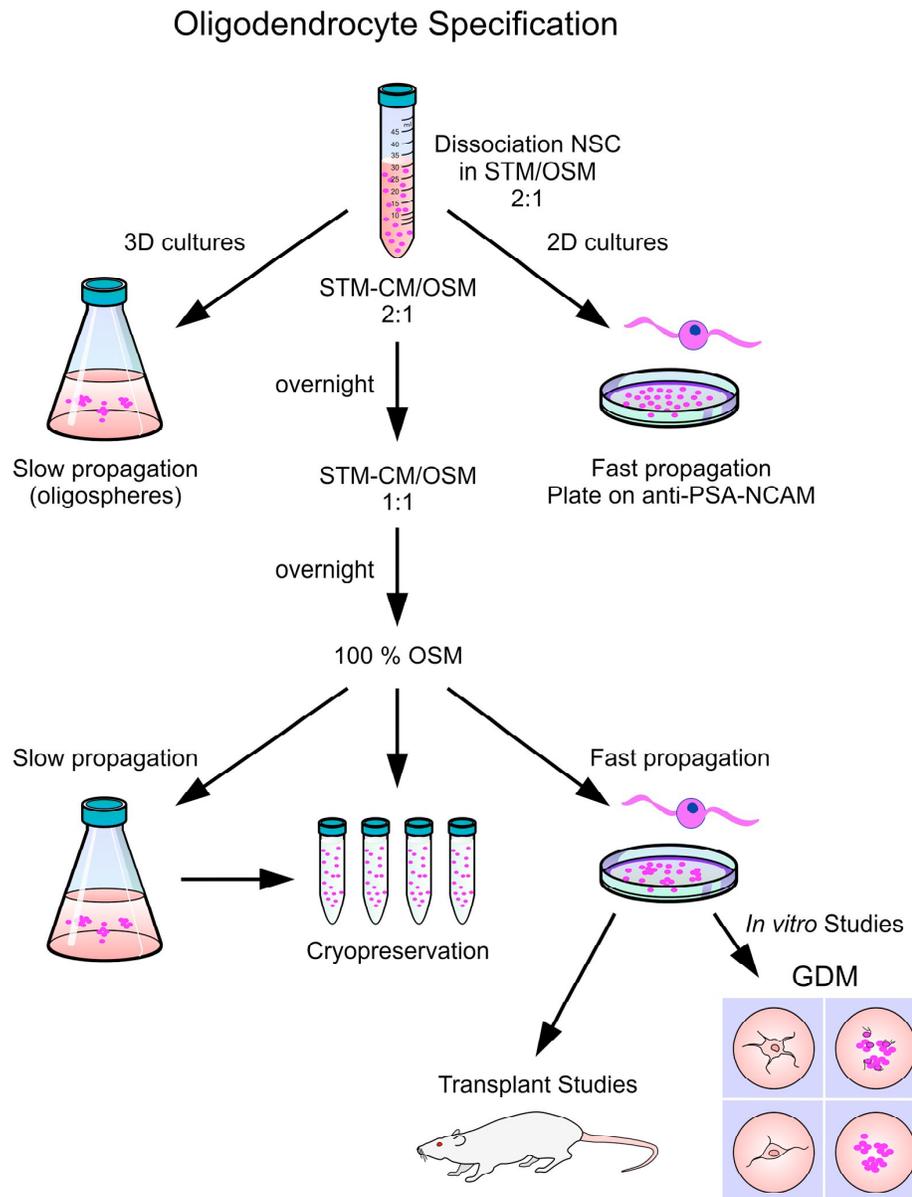
*Procedure:*

1. When NSCs reach confluency, remove the supernatant (CM), and add 5ml of Hank's Buffered Salt Solution (HBSS) without  $\text{Ca}^{2+}$  or  $\text{Mg}^{2+}$ .
2. Detach the cells with a cell scraper, transfer into a 15ml tube (1 to 3 Petri dishes), rinse once with 2ml of HBSS, and centrifuge at 45 RCF for 8min.

3. Resuspend the cell pellet in 3 ml OSM media and gently dissociate (3X) using a 14-gauge needle. Centrifuge at 45 RCF to pellet the cells.
4. Resuspend the cells in fresh OSM + STM-CM and seed cells on anti-IgM coated dishes or flasks (use the same procedure as for anti-PSA/NCAM), and maintain the cells as described in *Basic Protocol 1* but using OSM/self-CM (2:1) respectively. From this point on, the \*CO<sub>2</sub> concentration in the incubator should remain at 4.5%.
5. Feed the cells with 1/3 self-conditioned-OSM + 2/3 fresh OSM every other day until they reach 80-90% confluency. This process can be repeated several times to attain a large number of cells for freezing (if desired).
6. Alternatively, to grow OL spheres to create/enrich a frozen stock of OLP, place the equivalent of 2mm<sup>2</sup> (pellet size after dissociated and in suspension) in a 25ml Erlenmeyer flask with 15ml of OSM + self-CM (2:1). If the pellet is 4mm<sup>2</sup>, use a 50ml Erlenmeyer flask. Prepare the cell suspension and place in a total volume of 25ml of OSM + self-CM (2:1). Feed OL-spheres with fresh OSM every other day by adding 3 ml of freshly prepared OSM (no CM). When spheres start to become larger than 2mm, gently dissociate 1-2 times in the same flask with the 14-gauge needle using a 12ml syringe (sterile).
7. When the culture medium starts to turn orange, recover and centrifuge the spheres, and split into more Erlenmeyer flasks. These may be used for experiments or cryopreserved as previously described (cryopreservation section).

\*We recommended pre-calibrating the percentage of CO<sub>2</sub> one day before plating the cells. If the incubator is shared with other people or needed at 5% for NSC propagation and maintenance, we recommend using T-flasks for 2D cultures instead of Petri dishes. Close the cap completely and then open it ¼ of a turn before placing in the incubator at 5% CO<sub>2</sub>. For

propagation and maintenance of OL-spheres, the Erlenmeyer flask should also be kept open just enough to ensure O<sub>2</sub> / CO<sub>2</sub> exchange. When using 4.5% CO<sub>2</sub>, loosen the caps of the flasks until half-way open.



**Figure A-3. Oligodendrocyte Specification.** The transition of NSC to commit to the OL lineage is brief but sequential, rather than abrupt. In order for cells to survive, they must acclimate to their new environment. OLP can be propagated to create frozen stocks as 3D “oligosphere” cultures (shown on the left of the diagram) or frozen without propagation (as shown in the sequence in the center of the diagram). OLP can also be propagated in 2D cultures for cryopreservation, for specific cell culture experiments, or for cell replacement therapies (as shown in the right side of the diagram).

### *Support Protocol 3*

#### **Oligodendrocyte lineage progression and maturation**

The nutritional needs for a committed cell within the OL lineage differ considerably as they progress and mature to the next developmental stage. These cells need to start synthesizing enzymes and proteins related to myelination, therefore, the energy demand is enormous compared to their earlier stage where migration and proliferation are the basic functions. The culture medium “GDM” (glial defined medium) was first designed to maintain O4+, GC+/-, CNP+/- cells (for details see Pre-OL in [Figure A-1]). Later, we realized that GDM also induced the transition of OLP to pre-OL (Espinosa et al., 1997).

#### *Materials:*

Media: OSM, GDM, OLDEM

bFGF

Petri dishes with cells

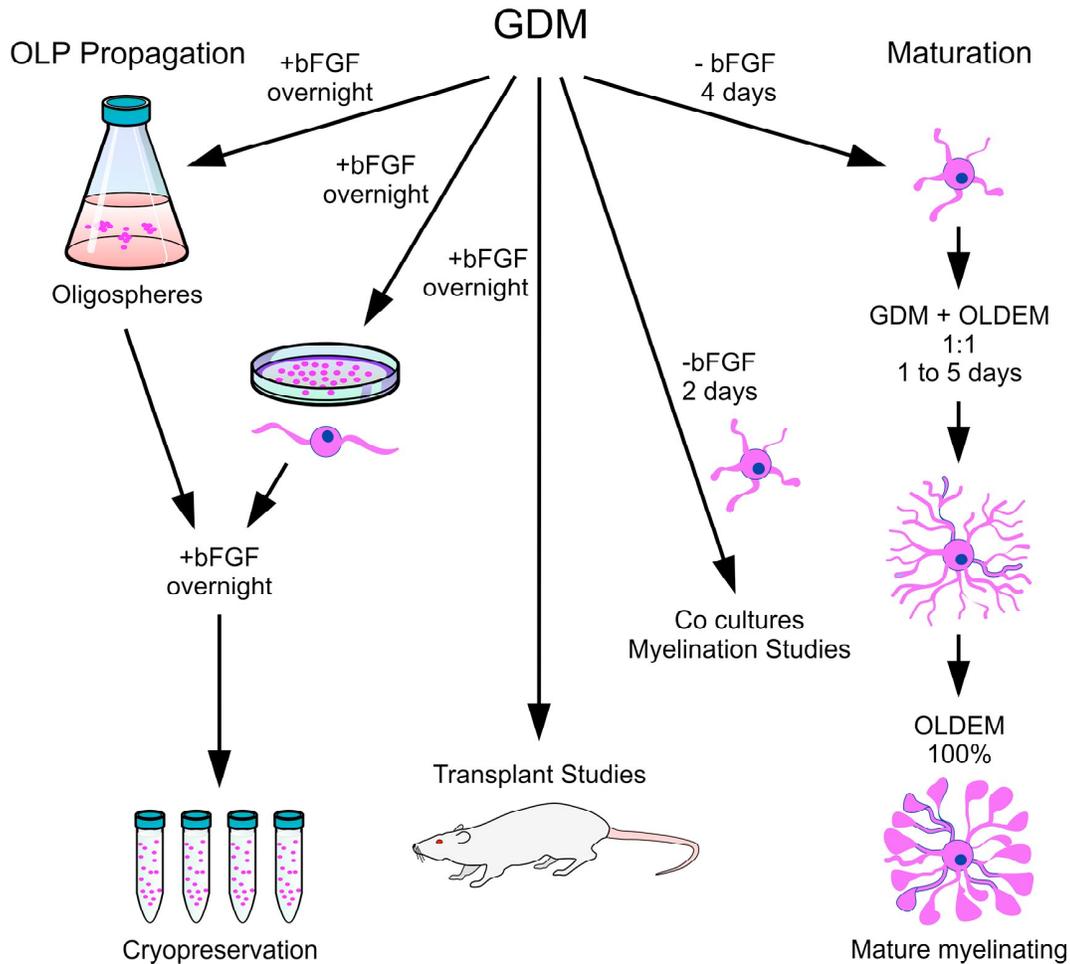
Poly-d-lysine coated wells/plates

24-well plate

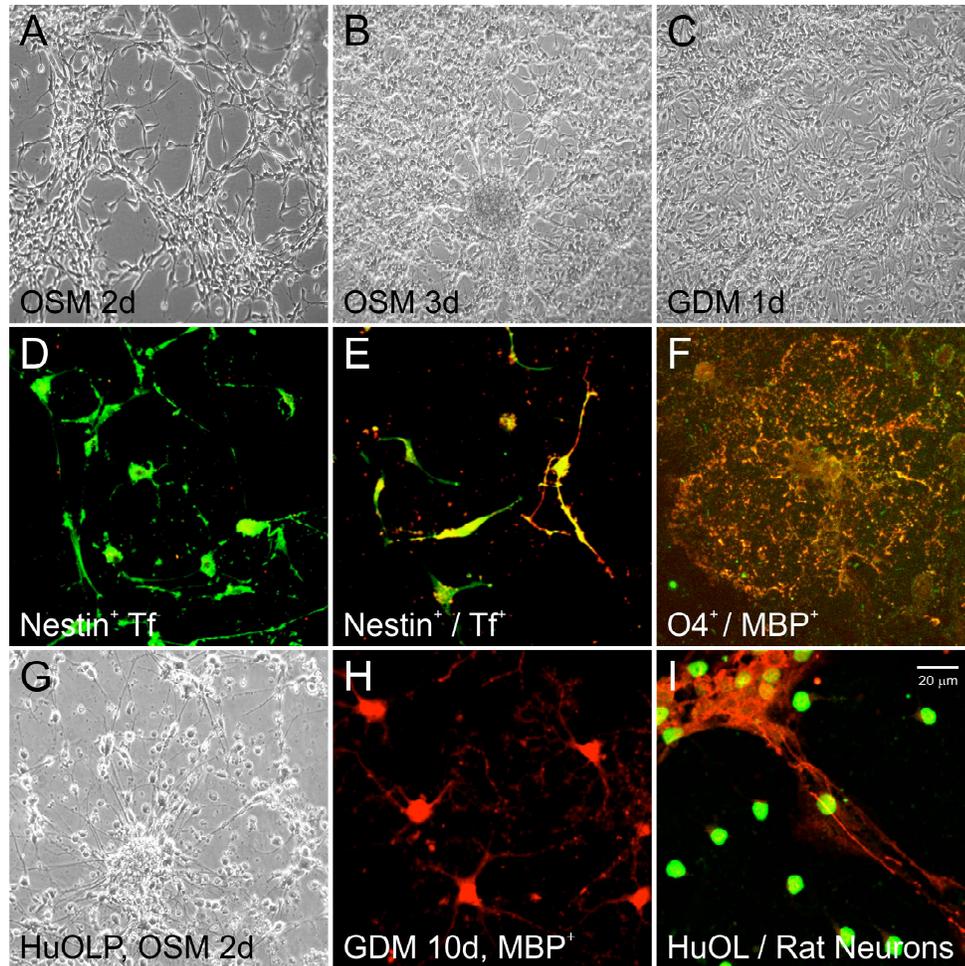
#### *Procedure:*

1. In order to obtain “pre-OL” (along the OL lineage), plate OLPs using OSM (as in Basic Protocol 2).
2. Next day, remove ½ of the volume of the plating medium (OSM) and add the same volume of GDM. As in previous steps, they may be propagated as OL-spheres (3D) or as 2D cultures on anti-IgM coated flasks, Petri dishes, or directly on cell culture grade plastic.
3. To obtain more OLP/pre-OL, cells are grown as 2D or 3D cultures in the presence of bFGF [Figure A-4]. Cells will remain at the same stage as the parent cells by adding fresh GDM + [20ng/ml] bFGF. For cell replacement therapies, we suggest using cells at this stage (1-2 days after plating without bFGF) as cells are still highly motile and readily migrate within the host post-natal and/or adult rodent brain and/or spinal cord.
4. To enhance maturation of cells into the next developmental stage, OL are cultured as 2D cultures in GDM for at least 2 days (if plated in GDM without bFGF), or 4 days (if plated in GDM + bFGF) without further bFGF supplementation [Figure A-5].
5. After exposure to GDM, cells express myelin enzymes and proteins, and they display multipolar, branched cell processes, but not a myelin-like membrane. In addition, OL maintained in GDM for at least 4d (without bFGF or any other factors) can be further induced to a fully mature myelinating stage.
6. To fully mature OL, plate as 2D cultures onto poly-d-lysine coated wells/plates or Petri dishes in GDM/OLDEM (OL maturation medium) 1:1 for 1-5 days, followed by 100% OLDEM thereafter [Figure A-4].

## Oligodendrocyte Lineage Progression and Maturation



**Figure A-4. Oligodendrocyte Lineage Progression and Maturation.** After commitment of NSC to the OL lineage, cells are propagated at the OLP stage to create a frozen stock (steps indicated on the left portion of the flow chart), or processed further for transplantation studies (as shown by the middle arrow on the diagram). To allow OLP to further mature along the OL lineage and become myelinated, cells are transitioned into OLDEM for at least 48h. Once OL have reached this stage of maturation, they are excellent for cell culture studies but are not recommended for cell grafting as detachment from the substrate can damage the numerous delicate cell processes.



**Figure A-5. Oligodendrocyte specification of NSC's derived from E16 rat.** Phase contrast view of NSCs derived from E16 rat brain at passage number two (P2), [A-C]. NSCs were plated and maintained in OSM for 2 days [A], 3 days [B], or 3 days in OSM then switched to GDM for 1 day, [C]. Cells in OSM still proliferate while in OSM. When cells from [A or B] are plated and maintained in OSM on poly-d-lysine coated coverslips for one day, they start to display a bipolar or multipolar morphology, [D] and most express the immature precursor marker, nestin (green) but not Tf (red), an early marker for OL. After two days in OSM, bipolar nestin<sup>+</sup> cells co-express Tf, [E]. After four days in OSM, cells were switched to GDM for one day, developed numerous cell processes and co-expressed sulfatides (recognized by the anti-O4 antibody, green) and MBP (red), [F]. (Panels G to I are human cells) [G], Phase contrast view of human NSCs (HFB-2050) acclimated and expanded in STM, then re-plated and maintained in OSM for 2 days. [H], OL derived from human NSCs (HFB-2050) were specified to the OL lineage with OSM and maintained in GDM for 10 days. OL matured and started to express MBP (red). [I], Rat cortical neurons (NFM-200-red) were cultured for 10 days, then human OLP derived from NSC (HFB-2050) were added in co-culture for 24 hours. Human Nuclei marker (HuNu green).

7. Every 4 days, feed the cells by replacing all of the culture medium with fresh OLDEM\*. The medium should look red, not orange. If it turns orange, add more medium while feeding the cells.

\* These cells will express myelin enzyme levels comparable to those found in pure myelin within 5 days after having been introduced to 100% OLDEM. As they mature, cells will synthesize myelin-like membranes *in vitro* even in the absence of neurons. They can be maintained for various weeks if they are sub-confluent, however, if the culture becomes overcrowded, cells will deteriorate and die.

#### ***Support Protocol 4***

##### **Propagation for *In Vitro* Myelination Assays**

To perform myelination studies *in vitro*, it is recommended to start with OL plated on plastic alone (rather than poly-d-lysine) and maintained in GDM for 2 days.

##### ***Materials:***

Media: GDM and OLDEM,

Cell Scraper

Neuronal Co-culture (See Support Protocol 5 below)

25  $\mu$ m sieve

24-well plates

*Procedure:*

1. Detach cells with cell scraper and centrifuge at 45 RCF for 8 minutes in the same culture medium.
2. Remove the supernatant and resuspend the cells in GDM-CM + fresh OLDEM (1:2).
3. A single cell suspension preparation is necessary for this step. Remove any cell clusters with a 25  $\mu\text{m}$  sieve (as described previously).
4. Count the cells and adjust the cell suspension to approximately [200,000 cells/ml] of OLDEM medium.
5. Remove half the volume of culture medium from the neuronal cultures without disturbing the cells (cortical neurons or dorsal root ganglion cells).
6. Slowly add the OL suspension to one 24-well plate containing the neuronal cultures to complete the original total volume in each well.
7. Follow the co-cultures for at least 10 days. To feed, replace  $\frac{1}{2}$  CM with fresh OLDEM. If the cultures are not overcrowded, they can be kept for at least 4 weeks.

***Support Protocol 5*****Preparation of Cortical Neurons***Materials:*

Instruments, animals and materials are the same as described in Basic Protocol 1.

Animals: Time-pregnant (ED14-16) Sprague-Dawley rats (Charles-River).

Bovine Serum Albumin (BSA)

Conical tubes

Culture tubes with cap, Sterile. Fisher 17 x 100mm

Hank's Balanced Salt Solution (HBSS)

Hemocytometer

Isoflurane

Needle

Poly-d-lysine

Scissors

Sterile Surgical Gauze

230  $\mu\text{m}$  and 140  $\mu\text{m}$  sieves

24 or 12 well plates

*Procedure:*

1. Prepare the work area and sterile tools in a biosafety hood [Diagram 1].
2. Euthanize the rats by isofluorane inhalation.
3. Extract the placenta containing embryos and place into STM Basal + 1% BSA.
4. Remove the cerebellum.
5. Dissect the brain of each embryo and place into Neurobasal-N complete (see Media Preparation).
6. Separate the cortex from the brain and remove the meninges with forceps.

7. \*Combine the cortical tissue of all of the brains without meninges. Mechanically dissociate with the needle by gently aspirating the brain pieces and releasing the suspension slowly against the wall of the tube ten times (try to minimize foaming).
8. \*Recover the supernatant with the cells in suspension and transfer to a 15ml tube.
9. Add 2 to 4 ml of Neurobasal-N medium to the chunks left over in the dissociation tube and dissociate again 5 to 8 times.
10. Filter the suspension of dissociated cells through 230  $\mu\text{m}$  and 140  $\mu\text{m}$  sieves to remove cell clusters.
11. Rinse the sieves sequentially with basal Neurobasal-N + 1% BSA at room temperature and add this medium to the tubes containing the cells.
12. Collect the cells by centrifugation in the culture tubes at 40 RCF for 8 min.
13. Discard the supernatant very gently as the pellet is very loose.
14. Resuspend the pellet in 4 ml of fresh Neurobasal-N medium with a 5ml pipette by gently triturating 2 or 3 times. Complete the volume to 12ml (or the equivalent of 1 embryo/ml) with 2 parts of fresh medium and 1 part of conditioned medium.
15. Assess \*cell viability, count cells using a hemacytometer, and plate onto poly-d-lysine coated well plates.
16. Incubate plated cells at 37°C with 4.5% CO<sub>2</sub>/95% humidity and monitor with a Combustion Test Kit (Bacharach # 10-500) (most electronic panels do not give an accurate reading).

Note: Cortical neurons maintained in Neurobasal-N do not need to be fed as frequently as other cell types. Simply add 500 $\mu\text{l}$  of Neurobasal-Nc every third day. Six days after plating, remove 1/4 of the culture medium and add the same volume of fresh Neurobasal-Nc.

\*Cell viability can be determined with SYTOX as described in Cell Viability Assay in Basic Protocol 1.

### ***Support Protocol 6***

#### **Transplantation of OL Progenitors into Neonatal Rats**

Neural progenitor cells and their differentiated OL counterparts can be stereotaxically transplanted into the newborn developing rat brain relatively non-invasively as previously described (Snyder et al., 1997; Flax et al., 1998; Espinosa et al., 2002). Similar results can be obtained with variations on the transplant method that are more suitable depending on the needs of the host brain and the type of study (Yandava et al., 1999; Ourednik et al., 2001, 2002; Park et al., 2002a; Teng et al., 2002; Wakeman et al., 2006; Lee et al., 2007; Redmond et al., 2007). A selection of detailed protocols for neonatal and adult mouse transplantation are described elsewhere (Espinosa et al., 1992; 1993a,b; Lee et al., 2008; Wakeman et al., 2009a,b; Yan et al., 2004). Upon implantation into the lateral ventricles, donor cells engraft and migrate from the subventricular zone into the host RMS in much the same manner as host NSC.

*Materials:* (may vary depending on the grafting method of choice).

Aspirator tube assembly (Sigma, A5177–5EA)

Borosilicate glass tube (Sutter Instrument, B100–75-15)

DPBS

Transillumination light source

Microfuge tube with cell sample

Micropipette puller (Sutter Instrument Co., Model P-87)

Neonatal rat (P0-P5)

Warming pad

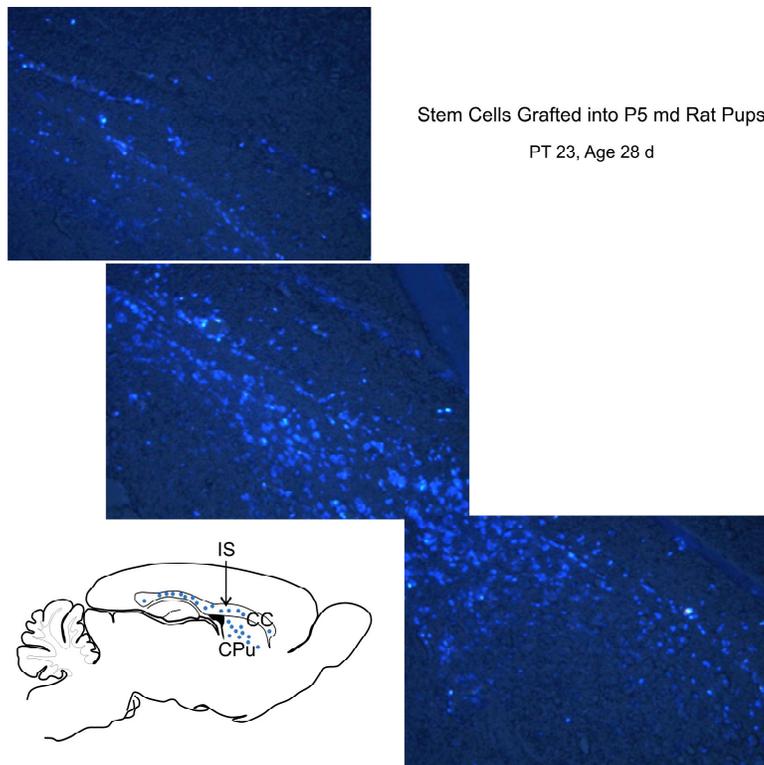
Warm Water Glove Balloon

Wet Ice

*Procedure:*

1. Anesthetize the rat pup until the animal no longer retains locomotion or responds to gentle toe and tail pinch. Carefully monitor the pup and proceed to transplantation.
2. Insert a calibrated, drawn borosilicate glass micropipette into the aspirator tube assembly, and rinse the micropipette by drawing and expelling 70% Et-OH five times followed by sterile dPBS ten times to clean the needle.
3. Gently flick sample in microcentrifuge tube prior to filling the needle, wipe the tube with 70% Et-OH, and uncap the tube.
4. Slowly draw 4-5 $\mu$ l cell suspension into the micropipette.
5. Loosely secure the head of the anesthetized pup and place directly over the light source to visualize the eyes and bregma.
6. Carefully insert the glass needle into the head at the midline between eye and bregma and slowly inject 2-5  $\mu$ l cell suspension at [5x10<sup>4</sup> cells/ $\mu$ l] into both lateral ventricles. Slowly remove the needle and check for leakage through needle tract. Repeat step 6 into the contralateral hemisphere.
7. After the injection, warm the pup by placing on a warm water balloon glove or heating pad to increase the body temperature before returning to the mother.

Note: In addition to the lateral ventricles, NSC can also be transplanted into the striatum, SN, and corpus callosum (CC) (Bjugstad et al., 2005, Redmond et al., 2007, and Bjugstad et al., 2008). Upon implantation into the CC of the host, HFB-2050 donor cells recognized by the fluorescent Fast Blue (FB) label migrated along the CC and into the CPu. [Figure A-6]. Pre-committed-OL can also be placed locally within focal sites of injury to decrease the need for extensive migration.



**Figure A-6. Stem cells engrafted into P5 rat pup.** Human OL derived from human fetal NSC were labeled with fluorescent fast blue (FB). A total of 60,000 cells were grafted into the corpus callosum (CC) of P(5) rat pups born to a md carrier mother. Twenty-three days after grafting, samples were harvested and examined. Grafted NSC survived and migrated extensively within the host brain parenchyma extending along the corpus callosum (CC) and caudate putamen (CPu). In the sketch, dots represent the location where FB+ cells were found. The sketch represents a sagittal view of the transplanted rat brain at 28 days of age, Injection Site (IS) indicates where cells were originally implanted.

***Commentary:***

**Background Information:**

The described culturing system allows for the production of relatively homogeneous primary OL cultures in adequate numbers for cryopreservation. These cell stocks can be used for basic research in further *in vitro* studies. Moreover, these cells are never exposed to animal or human sera, and therefore remain as suitable candidates for cell replacement therapies in

developmental disorders of the central nervous system (CNS) as well as neurodegenerative diseases.

Numerous methods and culture media described in the literature (even before, the times of NSC), were the basis for the optimization of the culture media formulations described here (some examples are, Botenstein and Sato, 1979; Saneto and de Vellis, 1985; Espinosa et al., 1988; Yang et al., 2005; Larsen et al., 2008., etc). Unndoubtedly, all previous reports on how to obtain and culture OL derived from NSC have also been instrumental in designing the present protocol. For example, the group of Lachapelle, and Baron-Van Evercooren described floating oligospheres derived from newborn rat brain (1996). This concept has been applied to NSC to generate OLs by Zhang (et al., 1998, Espinosa et al., 2002) and in the present protocol. Zhang and coworkers (1998) described the use of B104 neuroblastoma cell-conditioned medium (B104CM) to induce the oligodendrocyte phenotype on neurospheres and induce proliferation. This approach provides OL for many kinds of studies, but they are unsuitable as donor cells for cell replacement therapies to be used in translational studies, having used uncharacterized conditioned medium from B104 cells that have been grown in the presence of fetal bovine serum (as originally described by Louis et al., 1992). An example on the use of the protocols described can be found in Chattopadhyay et al., 2008.

#### **Critical Parameters:**

We want to emphasize that fate restriction towards commitment from NSC to OLP (as defined in “*Basic Protocol 2*”) becomes *irreversible* after NSCs have been in OSM for at least 20 hours (2D and 3D cultures). Therefore, the progeny of these cells will define a homogeneous OLP population, ideal for biochemical, toxicological and pharmacological studies, as well as an appropriate and reproducible source of committed cells to be used in cell

therapy studies. Phenotype reversal of induced OLPs may be possible with genetic manipulation but we have not attempted such studies to date.

Always monitor the concentration of CO<sub>2</sub> with a Combustion Test Kit (Bacharach # 10-500), as most electronic panels do not provide an accurate reading. The proper lineage progression relies on precise control of CO<sub>2</sub> to maintain a pH that should remain accurate and controlled.

### **Troubleshooting:**

Human NSC are more fragile than their rodent counterparts, therefore, we recommend dissociation protocols that favor as little mechanical stress as possible. In our hands, enzymatic dissociation with 2-4 ml Accutase (Millipore) @ 37°C for 3-5 minutes or light mechanical trituration through a 14-gauge needle (3-5 times) is sufficient to dissociate hNSC into single cells and small 2-6 cell clusters. Detailed methodology can be found elsewhere (Wakeman et al., 2009a,b).

### **Anticipated Results:**

OLPs obtained utilizing this system are plated on anti-PSA/NCAM plates and will attain a bipolar morphology if maintained in freshly supplemented OSM. Cells can also be plated directly onto plastic (tissue culture grade). The morphology may look more flattened or fibroblastic, but if maintained in fresh OSM, the early markers such as Olig2, Tf, PDGF-R and NG2 will be expressed. At this stage, cells are still highly motile but will migrate less if plated onto poly-d-lysine. During this time, cells attain a more mature phenotype that truly represents their *in vivo* counterparts.

Our culture media formulation includes the minimum and sufficient nutrients to support a given developmental stage, thus, cells can not be kept indefinitely in these

conditions because the substratum dictates the organization of the molecules on the cell membrane and poly-d-lysine confers a more permanent adhesion to the cells. Consequently, they would have the tendency to mature based on the signals coming from the cell membrane-substrate interaction (rev. Linnemann and Bock, 1989; Mauro et al., 1994). Unfortunately, cells will not survive or remain healthy if maintained in OSM due to a lack of nutrients to support their transition to the next developmental stage. The same concept applies to the transition to more mature OL stages. The nutrients and substrate together contribute to support cell signaling that will result in the formation of multiple cell processes followed by the synthesis of myelin components and their organization for membrane formation.

**Time Considerations:**

The initial dissection and preparation of the primary cell suspension takes approximately 2h. From the moment cells are plated on anti-PSA-NCAM (if fed regularly with fresh humoral factors), 100mm dishes can be confluent within 3-4 days. Thus, generating 20 vials of rat NSCs for cryostorage would take approximately 16 days. The generation of OLP from rNSC takes approximately 24h; yet, generating OLP in high numbers (15 vials) for storage would take 4-6 weeks. Lineage progression of rat OL towards more mature phenotypes takes approximately 48h in the specific culture medium (GDM or OLDEM). In addition, OL will still proliferate in GDM but at a much slower rate. Both, GDM and OLDEM media are favorable to protein synthesis but less favorable for cell proliferation.

Previously isolated ES cells and their NSC derivatives will need a longer period of time to provide high numbers of NSC for frozen stocks. This time will vary depending on the origin of the sample. We have had similar success directing NSC from several species, utilizing the same chemically defined media; however, incubation times may need to be increased for full maturation in higher order mammals, such as primates. Induced cells loose

NSC characteristics and acquire OLP features within 72h, yet their cell cycle is much slower and therefore, it would be necessary to propagate these cells 8 to 10 weeks to be able to create a healthy stock (6 to 8 vials) of human OLP. Previously established NSC lines (Snyder et al., 1992) can also be propagated and specified into the OL phenotype using the system described here

### **Acknowledgements:**

Appendix A, is a reprint in part of material as it appears in *Curr Protoc Stem Cell Biol*. Chapter 2:Unit 2D.4., Sep 2009 with co-authors Araceli Espinosa-Jeffrey, Seung U. Kim, Evan Y. Snyder, and Jean de Vellis.. The dissertation author was the co-primary investigator and co-author of this paper. . DRW and AEJ performed all experiments , co-wrote, and co-prepared the manuscript. EYS, SUK, and JdV supervised. Together with his coauthors, he thanks the MRRC Media Core for preparation of figures and Dr. D. Birt for photograph of the dissection set up. This work (J. de V. and A. E.) was supported in part by PPGHD065-76 and by a Pilot grant from the National Multiple Sclerosis Society PP1498. DRW thanks M. Hudson for critical review and comments. DRW is supported in part by the American Parkinson's Disease Association, HHMI Med-Into-Grad Training Fellowship, and the UCSD-NIH Training Fellowship in Clinical Genetics.

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