

# Lineage restriction of adult human olfactory-derived progenitors to dopaminergic neurons<sup>\*</sup>

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## ABSTRACT

Human adult olfactory epithelium contains neural progenitors (hONPs) which replace damaged cellular components throughout life. Methods to isolate and expand the hONPs which form neuronspheres *in vitro* have been developed in our laboratory. In response to morphogens, the hONPs differentiate along several neural lineages. This study optimized conditions for the differentiation of hONPs towards dopaminergic neurons. The hONPs were treated with Sonic hedgehog (Shh), in the presence or absence of retinoic acid (RA) and/or forskolin (FN). Transcription factors (Nurr1, Pitx3 and Lmx1a) that promote embryonic mouse or chicken dopaminergic development were employed to determine if they would modulate lineage restriction of these adult human progenitors. Four expression vectors (pIRES-Pitx3-Nurr1, pLN-CX2-Pitx3, pLN-CX2-Nurr1 and pLNCX2-Lmx1a) were transfected into the hONPs, respectively. Transcription factor expression and the rate-limiting enzyme in dopamine synthesis tyrosine hydroxylase (TH) were detected in the transfected cells after 4 month-selection with G418, indicating transfected hONPs were stably restricted towards a dopaminergic lineage. Furthermore, a dopamine enzyme immunoassay (EIA) was employed to detect the synthesis and release of dopamine. The most efficient transfection paradigm was determined. Several neurotrophic factors were detected in the pre-transfected hONPs which have potential roles in the maintenance, survival and proliferation of dopaminergic neurons. Therefore the effect of transfection on the neurotrophin synthesis was examined. Transfection did not alter synthesis. The use of olfactory progenitors as a cell-based

therapy for Parkinson's disease (PD) would allow harvest without invasive surgery, provide an autologous cell population, eliminate need for immunosuppression and avoid the ethical concerns associated with embryonic tissues. This study suggests that specific transcription factors and treatment with morphogens can restrict human adult olfactory-derived progenitors to a dopaminergic neuronal lineage. Future studies will evaluate the utility of these unique cells in cell-replacement paradigms for the treatment of PD like animal models.

**Keywords:** Human Olfactory Epithelium; Progenitors; Dopaminergic Neurons; Parkinson's Disease

## 1. INTRODUCTION

Parkinson's disease (PD) remains one of the leading causes of chronic degenerative neurological disability, which affects more than 6,000,000 people world-wide, with approximately 60,000 new cases diagnosed each year in the United States [1]. The incidence rises with age, being approximately 1:1000 overall and 1% of the population over the age of 60 and 4% in those over 80 years. Unfortunately, the mortality rate of PD has increased steadily in recent years [2,3]. PD is characterized by the extensive loss of dopaminergic (DA) neurons in the substantia nigra (SN) in the midbrain [4]. Currently the principle treatment for PD is oral L-3, 4-dihydroxyphenylalanine (L-dopa) [5], which is the precursor of dopamine that can pass the blood-brain-barrier [6]. L-dopa promotes symptomatic relief, but with time becomes less effective for two reasons: 1) During the progression of the disease the neurons become less sensitive to the drug [7] and 2) L-DOPA does not prevent or rescue the DA neurons from degeneration [8,9].

Recent research has attempted to find cell populations that can be used to replace lost or degenerating dopaminer-

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gic neurons [2,10,11]. The basic concept of cell replacement therapy is to restore function lost as a result of the disease in the central nervous system (CNS) by replacing degenerating or lost cells with viable functional cells. Recent studies also suggest that the engraftment of stem cells or progenitors can up-regulate or enhance existing endogenous progenitor populations [12-14]. Studies have employed neural cell grafts obtained from the fetal ventral mesencephalic (VM) dopaminergic neurons [15-20]. However, this frequently resulted in significant dyskinesia [21-24]. Even when clinical improvements were achieved in the absence of dyskinesia, the amount of tissue required for each PD patient necessitated a minimum of 4-5 fetal brains [25]. This requirement increased the possibility of viral or bacterial infection and significantly limited the utility of this approach. In addition the number of surviving neurons was highly limited as the majority of the engrafted cells died in the initial days following transplantation [15,20,24]. The limited supply of fetal VM cells coupled with their poor graft viability severely limited the therapeutic utility of this population for the treatment of PD. Therefore, an alternate expandable source of dopamine cells has become a major research focus [26-29].

Stem cells are undifferentiated cells with an unlimited capacity for self-renewal and the potential for lineage restriction (maturation) into one or more specific cell types, depending on their origin and the micro-environmental signals that they receive [28,30]. These characteristics make stem cells an attractive target population for PD cell replacement therapy [31-34]. Human embryonic stem cells (hESCs), lineage-restricted towards dopaminergic neurons when transplanted into a rodent model of PD, provide a significant relief of symptoms. However, with time, animals engrafted with hESCs have frequently developed teratomas [35]. Clearly an alternate approach is warranted.

Human olfactory epithelium (OE) is a unique source for neural progenitors that can be harvested by minimally invasive endoscopic nasal surgery without a craniotomy. Furthermore since no demonstrable olfactory deficits result from OE biopsy [36], the tissue can be used to generate an autologous progenitor population from patients with PD. An autologous cell source provides total histocompatibility and thus eliminates the need for immunosuppressive therapy as well as long waiting lists for available matched tissue. Previously our laboratory developed methods for the isolation and culture of a neurosphere forming population [37]. To date more than 150 patient-specific cell lines of human olfactory neural progenitors (hONPs) have been established from primary cultures of human adult olfactory epithelium isolated from cadavers [37] and patients undergoing endoscopic sinus surgery [36]. Our studies have

shown that the hONPs have the potential to differentiate along several neural lineages following exposure to environmental signals *in vitro* [38].

The objective of this study was to determine if hONPs could be lineage restricted towards dopaminergic neurons and if so to optimize the methodology. Molecular techniques were applied for the transfection of Nurr1 [34, 39], Pitx3 [40,41] and Lmx1a [42], transcription factors which promote dopaminergic differentiation. The transfection effects of different paradigms were evaluated and compared.

Several studies have shown that neurotrophic factors, such as brain-derived neurotrophic factor (BDNF), ciliary neurotrophic factor (CNTF), neurotrophin-3 (NT-3), etc. are important for the survival and function of dopaminergic neurons in CNS [43-47]. Recent studies also indicate that the neurotrophins have the potential to optimize the local micro-environment of the damaged area, and thereby induce endogenous stem cells to replace or rescue degenerating neurons [48,49]. HONPs derived from adult human olfactory epithelium have been shown to produce and release neurotrophins [10,50,51], which could further support their use in a cell-based therapy for PD. Therefore, this study also evaluated the ability of pre and post transfected hONPs to synthesize key neurotrophins.

## 2. MATERIAL AND METHODS

### 2.1. Cell Culture

The two patient-specific olfactory progenitor lines used in this study were obtained from adult olfactory epithelium harvested from a 42-year-old female patient and a 20-year-old male via endoscopic biopsy [37]. The tissues were cultured to allow the emergence and harvest of hONPs as previously described [36,52]. The hONPs were thawed from frozen stock that was maintained in liquid nitrogen and cultured in minimal essential medium (MEM) with 10% heat inactivated fetal bovine serum (FBS, GIBCO, Grand Island, NY) (10% OE) for one week. The hONPs were adapted to serum-free growth media via serial dilution of serum every day for 4 days until the cells were finally cultured in DFBNM (DMEM/F12 supplemented with 1% B27 and 0.5% N2 and 100 µg/ml gentamycin (GIBCO, Grand Island, NY)) [52]. Parallel independent experiments were performed on hONP lines from the two different patient lines. Since equivalent results were achieved, data from only one line has been presented.

### 2.2. Construction of Expression Plasmids

The mouse Nurr1 cDNA was cloned into the pLNCX2 expression vector (Clontech) between ClaI. Similarly,

the rat Pitx3 and mouse Lmx1a cDNA were inserted into pLNCX2 vector between ClaI. For the Nurr1 and Pitx3 co-expression vector, Nurr1 cDNA was cloned into pIRES (Clontech) between XbaI and SalI, and pitx3 was inserted between EcoRI. The pLNCX2 and pIRES expression vectors served as controls (**Figure 1**). All expression vectors were verified by extensive DNA sequencing.

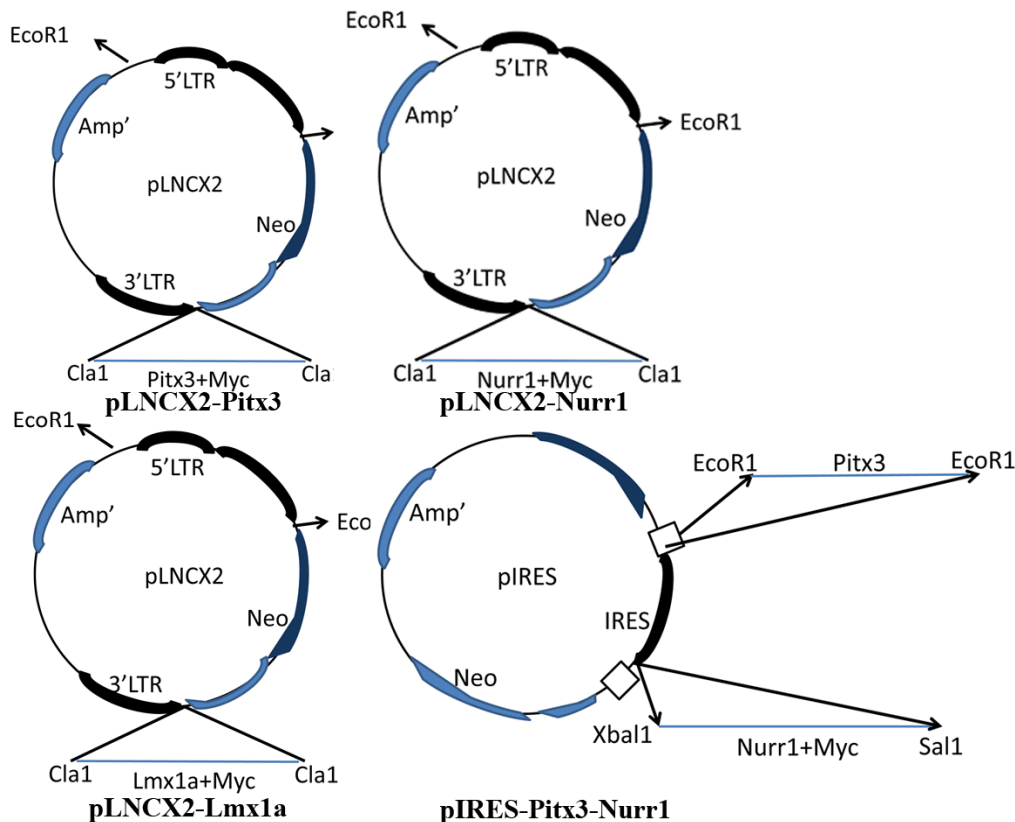
### 2.3. Transfection and Selection

All plasmid constructs were introduced into the hONPs by liposomal transfection. The cells were plated on glass coverslips in six-well plates ( $5 \times 10^4$  cells per 35 mm well) in DFBNM without antibiotics 1 day before transfection. hONPs were transfected with each plasmid (4  $\mu$ g/well) for 24 hours according to the manufacture's protocol (Lipofectamine 2000, Invitrogen, Carlsbad, California). One day after transfection, the cells were fed with 10% FBS in MEM and selected with G418 (400  $\mu$ g/ml; Invitrogen, Carlsbad, California). The selection pressure was kept for up to 4 months to insure a purified stably transfected cell population. Immunocytochemistry and Western blot analysis were applied to detect several dopaminergic neuronal markers. After a four-month selection, the trans-

fected hONPs were frozen in liquid nitrogen for additional four-six months of storage. After removal from cryostorage and several days' recovery in MEM with 10% FBS at 37°C, the dopaminergic lineage restriction was probed with immunocytochemistry and Western blot analysis.

### 2.4. Treatment with Morphogens

The hONPs were treated with Sonic hedgehog (Shh) in the presence or absence of retinoic acid (RA, 1  $\mu$ M) and/or forskolin (FN, 5  $\mu$ M) [52]. Highly purified Shh (kindly provided under a Material Transfer Agreement with Curis and Wyeth, Inc.) was applied to hONPs and compared to a commercially available control sample obtained from Sigma to determine the extent to which purification of Shh can affect the expression of tyrosine hydroxylase (TH). The hONPs were plated on glass coverslips in six-well plates ( $5 \times 10^4$  cells/35 mm well) in DFBNM and treated with medium containing various concentrations and combinations of RA, FN, and Shh for 7 days (CO<sub>2</sub> atmosphere at 5% and temperature of 37°C). Treatment with Shh included several concentrations: 0.25 mg/ml (Shh0.25), 0.1 mg/ml (Shh0.1), 0.05 mg/ml (Shh0.05), 0.025 mg/ml (Shh0.025) in the presence or absence of



**Figure 1.** Construction of expression plasmids.

1  $\mu$ M retinoic acid (RA1) and/or 5  $\mu$ M forskolin (FN5). After treatment, the TH expression was determined at 1 - 7 days *in vitro* by immunocytochemical analysis. Once the optimized environment for inducing dopaminergic neurons was determined, the medium containing the optimized combination was applied to stably transfected hONPs to further improve the yield of these neurons.

## 2.5. Immunocytochemistry

The hONPs ( $5 \times 10^4$  cells/well) were plated on 35 mm round glass coverslips in six-well plates (Becton, Dickinson and Co.) and incubated at 37°C in 5% CO<sub>2</sub>/95% air for 24 hours and treated with RA, FN, and Shh or transfected and selected for different periods of time prior to fixation for immunofluorescence. 4, 6-diamidino-2-phenylindole dihydrochloride (DAPI) (1:1000, 2 mg/ml, Molecular Probes, Eugene, OR) was applied in culture for 30 minutes at 37°C for vital nuclear staining. The coverslips were rinsed with cytoskeletal buffer (CB) twice and fixed in 3% paraformaldehyde in CB (10 minutes). 0.2% Triton X-100 (10 minutes, Sigma) in tris buffered saline (TBS) was applied and cells were incubated (1 hour) in 3% bovine serum albumin (BSA) in TBS. Primary antibodies were applied overnight (4°C). After 30 minutes washing (10 minutes each, 3 times) in TBS, the cells were incubated with secondary antibodies: Texas-red conjugated goat anti-rabbit immunoglobulin G (IgG), Texas-red-conjugated goat anti-mouse IgG, Cy2-conjugated goat anti-mouse IgG and/or Cy2-conjugated goat anti-rabbit IgG (all diluted 1:600, Cy2, Jackson Immunology Research Laboratories; Texas red, Molecular Probes). The coverslips were rinsed in TBS for 30 minutes (10 minutes each, 3 times) and mounted on slides. The slides were examined with confocal microscopy. All experiments were repeated a minimum of two times to ensure the specificity of staining; only one set of data has been presented since similar results were obtained.

## 2.6. Western Blot Analysis

Western blot analysis was used to further examine and confirm the immunofluorescence studies. Proteins from hONPs transfected with control vectors, as well as hONPs transfected with the vectors plus each combination of transcriptions factors (pLNCX2-Pitx3, pLNCX2-Nurr1, pLNCX2-Lmx1a, pIRES-Pitx3-Nurr1), cultured in DFBNM, selected in all groups were collected in cell lysis buffer (Sigma, St. Louis, MO). After 15 minutes of incubation on ice, samples were centrifuged for 30 minutes (4°C) and the protein concentration of each supernatant was determined. The protein samples (20  $\mu$ g/well) were electrophoresed on 10% SDS-polyacrylamide gels along with standardized-molecular-size marker proteins

in an adjacent lane and transferred from gel to nitrocellulose paper. Nonspecific binding was blocked (1 hour) with 5% nonfat dry milk in TBS-Tween (TBST) buffer. Blots were incubated (4°C overnight) in primary antibodies (anti-TH, MAB; anti-actin, MAB). Blots were washed three times for 10 minutes in TBST, after which they were incubated (1 hour, room temperature) monoclonal horseradish peroxidase-labeled anti-mouse IgG (1:2000). ECL Western blotting detection reagents (Amersham Biosciences) were used to identify bound antibodies. Densitometry of the protein bands was carried out on a high performance chemiluminescence film (Amersham Biosciences). Data was analyzed using the Image-J software programs supplied by the NIH official website (<http://rsb.info.nih.gov/ij/>).

## 2.7. Dopamine Assay

Stably transfected hONPs were plated into flasks (25 cm<sup>2</sup>, Corning) at  $10^5$  per flask before they were adapted to the absence of serum via serial dilution of serum every day for 4 days until the cells were finally cultured in DFBNM, which was collected daily after the serum was totally eliminated from the medium. The DFBNM collected from each restricted hONP line was then concentrated to 1/50 volume respectively by centrifugal filters (Amicon Ultra-15, Millipore). The differentiated hONPs were then collected and lysed (lysis buffer, Sigma). Dopamine expression was analyzed quantitatively in the concentrated medium as well as in the cell lysates with a dopamine enzyme immunoassay kit (Dopamine EIA, Immuno Biological Laboratories, Inc.), according to the manufacture's protocol.

## 2.8. Neurotrophin Assay

Pre- and post-transfected hONPs were plated into flasks (25 cm<sup>2</sup>, Corning) at  $5 \times 10^5$  per flask and cultured in 10% OE media for two days before they were adapted to the absence of serum via serial dilution of serum every day until they were finally cultured in DFBNM. The differentiated hONPs were then collected and lysed (lysis buffer, Sigma). Neurotrophins were detected in the extracted protein with different enzyme-linked immunosorbent assay (ELISA) kits (BDNF, Chemicon; CNTF, Quantikine; NT-3, Chemicon) respectively, according to the manufacture's protocol. The ELISA absorbance (OD) was obtained with a microplate spectrophotometer (Spectra-max Plus), and the results were plotted and calculated with the compatible software (Softmax Pro).

## 3. RESULTS

Cryopreserved vials of the two representative hONP



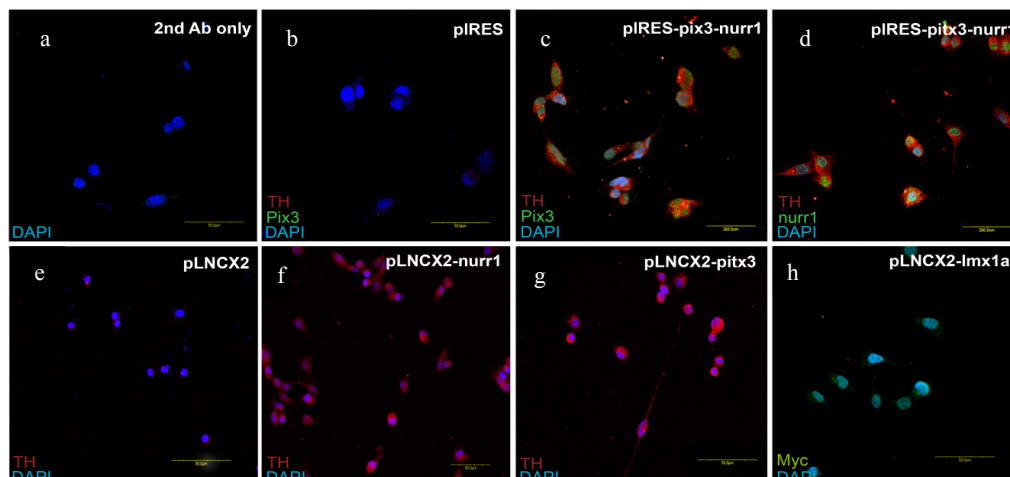
lines were obtained from storage and grown for 1 - 2 weeks prior to the start of these experiments to insure equivalent passage (4 - 8) and sufficient cell numbers for the following studies.

### 3.1. Transfection of Olfactory-Derived Progenitors (hONPs) to Achieve Dopaminergic Lineage Restriction

HONPs were obtained from previously frozen stock with low passage number (4 - 8) and maintained in MEM 10 medium during their recovery period. These mitotically active cells divided every 18 - 20 hour which typically required passage three times per week as previously described. The heterogeneous nature of the hONP population prior to transfection was determined by immunocytochemistry. No reactivity was observed for Pitx3, Nurr1, Lmx1a with pre-transfected hONPs and only a few (5 - 10%) of them were positive for the dopamine precursor, TH, when treated conditionally [53]. Low passages (Passage 4 - 8) of hONPs from 2 different patient-specific cell lines were employed in parallel transfection experiments. To examine the phenotypic expression of hONPs after transfection and selection, representative cultures as well as their respective pre-transfection controls were evaluated. Non-transfected hONPs or those transfected with lipofectamine alone died within 1 week after selection with 400  $\mu$ g/ml G418. In contrast, 30% of the transfected cells (both with the concerned genes and the control vectors) survived under the selection pressure. Transfection with control vectors, single genes, or Pitx3-Nurr1 combined resulted in no morphologic changes compared to the typical pretreated hONPs. However, the transfected hONPs divided more slowly,

with a new doubling time of three to four days, which required a feeding schedule of only twice a week and necessitated passage every 9 - 10 days. Immunofluorescent analysis of the transfected populations demonstrated that hONPs were stably transfected and TH expressed.

- Human olfactory derived hONPs were transfected by pIRES-Pitx3-Nurr1 to restrict them towards DA neurons. The vector alone was employed as a control. To obtain a purified population of restricted cells the transfected population was maintained in G418 for selection. Although only several weeks of selection produced relatively pure populations, an interval of four months was employed to insure stability and purity. HONPs remained TH positive after transfection of pIRES-Pitx3-Nurr1, whereas the transfection of control vectors exhibited no phenotypic changes, demonstrating that hONPs can be restricted towards dopaminergic neurons (**Figure 2**).
- HONPs were transfected with pLNCX2-Nurr1, pLNCX2-Pitx3, pLNCX2-Lmx1a or the vector alone as a control. The transfected cells were exposed to G-418 for selection for periods up to 4 months. HONPs were TH positive after transfection of pLNCX2-Nurr1 and pLNCX2-Pitx3, whereas the transfection of control vectors resulted in no phenotypic changes. Therefore pLNCX2-Nurr1 or pLNCX2-Pitx3 can be employed to lineage restrict the hONPs towards dopaminergic neurons. In contrast, the hONPs transfected with pLNCX2-Lmx1a remained unreactive for TH, although positive of myc, which demonstrated the successful incorporation of the plasmid (**Figure 2**).



**Figure 2.** Immunocytochemical analysis. HONPs transfected with pIRES-Pitx3-Nurr1, pLNCX2-Pitx3 or pLNCX2-Nurr1 were tyrosine hydroxylase (TH) positive after 4 months selection with G418 (c, d, f, g), while the lines transfected with pIRES or pLNCX2 were TH negative (b, e). HONPs transfected with pLNCX2-Lmx1a were Myc positive, demonstrating that the plasmid was transfected into the nucleus (h).

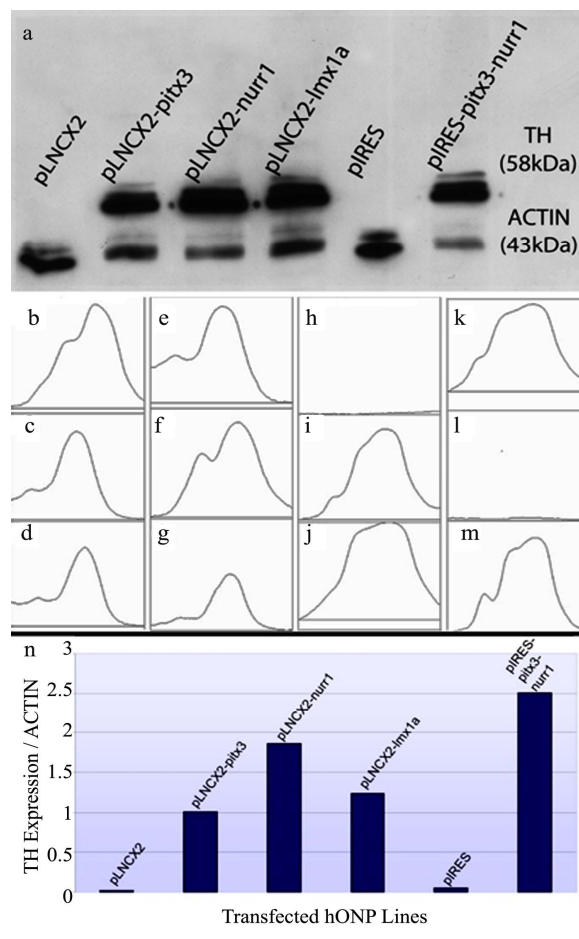
- Western blot analysis was employed to confirm quantitatively the immunocytochemical studies of the transfected hONP populations. The following transfected lines were analyzed for TH expression: hONPs transfected with pIRES-Pitx3-Nurr1, pLNCX2-Nurr1, pLNCX2-Pitx3 and pLNCX2-Lmx1a all of which were TH positive, which indicated their potential to release dopamine. In contrast, the hONP populations' transfected with the control vectors (pIRES and pLNCX2) did not express TH.  $\beta$ -actin, a protein that is widely expressed in all mammalian and avian cells was used as a reference protein for the comparison of TH expression by the various lines. Image-J was applied for the data analysis. Each curve from B to M in **Figure 3** illustrates the density of bands evident on the western gel (**Figure 3a**), and the area that each curve was measured. The bars in picture N represent the ratio of TH expression and ACTIN expression in the cell line. hONPs transfected with pIRES-Pitx3-Nurr1 exhibited the highest ratio for the TH and ACTIN expression, while the cells transfected with the control vector (pLNCX2 or pIRES) had the least TH staining (**Figure 3(b-n)**). These results demonstrate that individual transcription factors have unique abilities in promoting the dopaminergic restriction of hONPs.

### 3.2. Transfected hONPs Remain Restricted to Dopaminergic Lineage after Removal from Cryostorage

After a 4-month selection, the dopaminergic lineage restricted cells were cryopreserved in liquid nitrogen for additional 4 - 6 months. Following their removal from cryostorage and several days' recovery in MEM10 at pIRES-Pitx3-Nurr1 to restrict them towards DA neurons. The vector alone was employed as a 37°C, all but one of the transfected hONP populations survived under the selection pressure of 400  $\mu$ g/ml G418, demonstrating that these cells were stably transfected and retained their potential for long term storage and clinical application. Immuno-cytochemistry and Western blot analysis was applied to these previously stored populations to examine their TH expression. The hONPs transfected with pLNCX2-Pitx3, pLNCX2-Nurr1 and pIRES-Pitx3-Nurr1 remained healthy and TH positive under the pressure of selection, while the pLNCX2-Lmx1a transfected line did not (**Figure 4**).

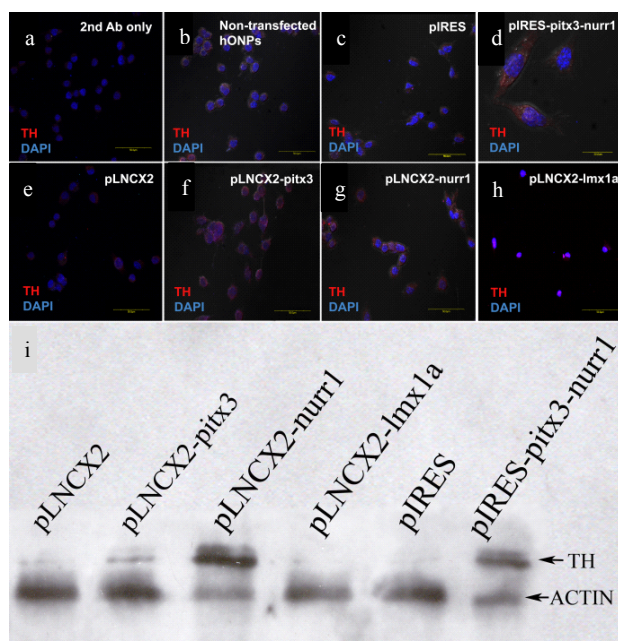
### 3.3. Lineage Restricted hONPs Produced and Released Dopamine

After removal from the cryostorage, dopamine production was detected in the hONP lines which were stably transfected with concerned genes, while the cells transfected

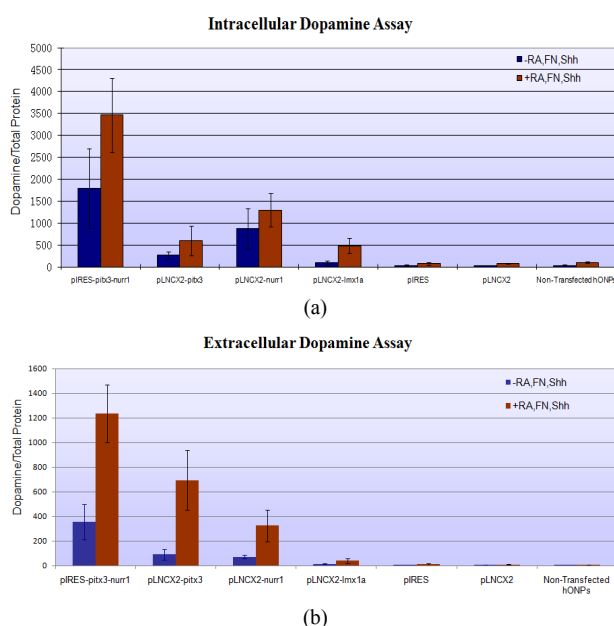


**Figure 3.** a. Western blot analysis. b-g. Scanning densitometry demonstrates ACTIN-expression in a hONP line of pLNCX2, pLNCX2-Pitx3, pLNCX2-Nurr1, pIRES and pIRES-Pitx3-Nurr1 respectively. h-m. Densitometry of TH-expression as shown in A. N. Histogram demonstrating the ratio of TH/action produced by each population.

with control vectors and the non-transfected hONPs didn't produce dopamine. The dopamine level of each sample was then divided by the concentration of protein in each specific hONP line to calculate the efficiency of dopamine production. Among all the 4 gene transfected lines, hONPs transfected with pIRES-Pitx3-Nurr1 exhibited the most efficient dopamine formation (**Figure 5(a)**). Spent medium was collected 4 days after culturing the lineage restricted hONPs. This medium was then concentrated to 1/50 volume respectively, and dopamine E. I. A. was applied to detect the dopamine release (extracellular levels). Data were calculated in the same manner as the intracellular dopamine analysis. Lower levels of dopamine were detected in the concentrated media compared to the corresponding analysis of the cell lysis. The greatest level of dopamine release was detected in pIRES-Pitx3-Nurr1 transfected hONPs compared to the other restricted cell lines (**Figure 5(b)**).



**Figure 4.** Immunocytochemistry (a-g) and western blot analysis (i) demonstrating that hONPs transfected with pLNCX2-Pitx3, pLNCX2-Nurr1 and pIRES-Pitx3-Nurr1 remain healthy and TH positive following removal from cryostorage under selection pressure (d, f, g). In contrast, the Lncx2-Lmx1a transfected line no longer expressed TH (h).



**Figure 5.** Histograms demonstrating the ratio of dopamine formation (pg/100 µl) to total protein concentration (mg/ml) of cells transfected with pIRES-Pitx3-Nurr1, pLNCX2-Pitx3, pLNCX2-Nurr1, pLNCX2-Lmx1a, pIRES, pLNCX2 and non-transfected hONPs. HONPs transfected with pIRES-Pitx3-Nurr1 exhibited the highest levels of intracellular and extracellular dopamine production. Dopamine production and release were enhanced in hONPs treated with the morphogens.

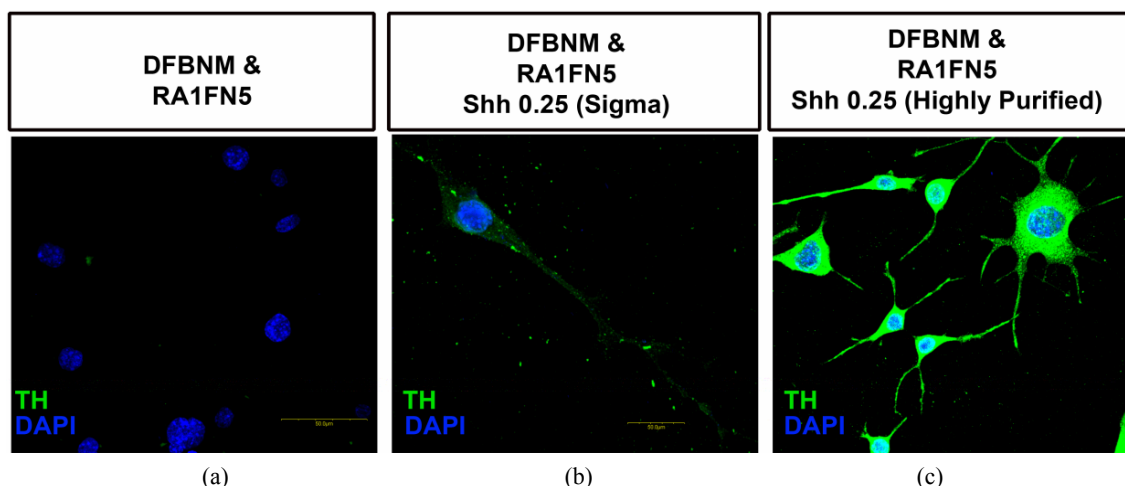
### 3.4. The Effect of Morphogens on Tyrosine Hydroxylase (TH) Expression, Dopamine Formation and Release

HONPs were cultured in DFBNM along with RA (1 µM), FN (5 µM) and either of two different sources (purities) of Shh for four days. Both Shh treatments resulted in greater expression than in those cultured solely in DFBNM. TH expression was greater in the cells that were treated with highly purified Shh than the commercial product obtained from SIGMA when applied for same period of time (**Figure 6**).

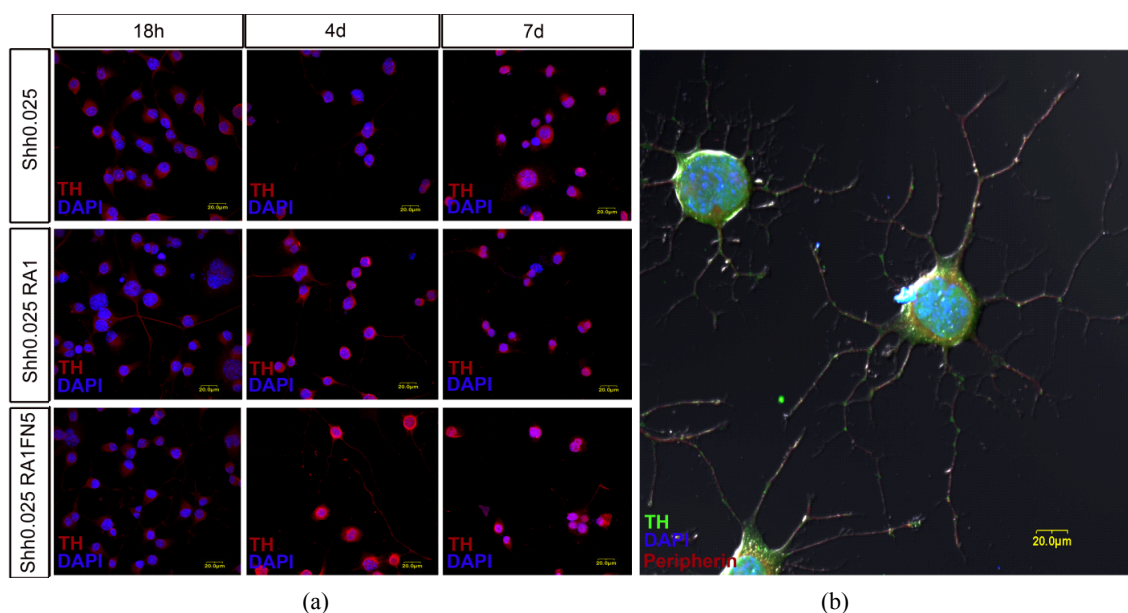
HONPs treated with RA1FN5 and highly purified Shh expressed seemingly more intensive TH reactivity in the positive cells (**Figure 7(a)**). Therefore, the concentration of Shh was reduced to determine the lowest concentration of Shh that could drive the hONPs towards dopaminergic neurons. In contrast to the response when a high level of Shh was applied, the reduction of the Shh to 0.025 mg/ml applied with RA (1 µM) & FN (5 µM) did not produce an immediate response. The hONPs became TH positive only after 18 hours of treatment with highly purified Shh; however, they were healthy and maintained TH expression for longer periods. The application of RA and FN promoted an even greater expression of TH (**Figure 6 A**). Therefore, the optimal conditions for restricting the hONP lineage to dopaminergic neurons (under these defined conditions) was determined to be DFBNM supplemented with RA1FN5Shh0.025 (**Figure 7(b)**).

Stably transfected hONPs were treated with a cocktail of RA1FN5Shh0.025 to determine if a combination of genetic modification and morphogen exposure would increase intracellular and intercellular dopamine levels. Spent medium was collected four days after morphogenic treatment and concentrated to a 1/50 volume. The treated lineage restricted hONPs were also collected. Dopamine E. I. A. was applied to both cell lysis sample and concentrated medium. Dopamine formation efficiency was calculated as previously described. HONPs transfected with pIRES-Pitx3-Nurr1 were the most efficient population with respect to dopamine formation and release after morphogenic treatment (**Figures 5(a)-(b)**). Compared to intracellular and extracellular dopamine levels of the lineage restricted hONPs in the absence of morphogens, dopaminergic expression was greatly enhanced in the stably transfected hONPs in the presence of the combination of Shh, RA and FN (**Figures 5(a)-(b)**). These studies suggest that treatment with morphogens can play an important role in dopamine formation and release by the lineage restricted hONPs.





**Figure 6.** HONPs treated in DFBNM with a highly purified Shh(c) exhibited greater reactivity to tyrosine hydroxylase (TH) than those treated with commercially available Shh (b) for 3 days in the presence of RA and FN.



**Figure 7.** (a) HONPs cultured in DFBNM supplemented with 0.025 mg/ml of Shh, in the presence or absence of retinoic acid (RA) (1  $\mu$ M) and forskolin (FN) (5  $\mu$ M) for days indicated; (b) HONPs were tyrosine hydroxylase (TH) positive following 7 days treatment with RA1FN5Shh.

### 3.5. Stably Transfected and Pre-transfected hONPs Produce Neurotrophins (BDNF, CNTF and NT-3) at Equivalent Levels

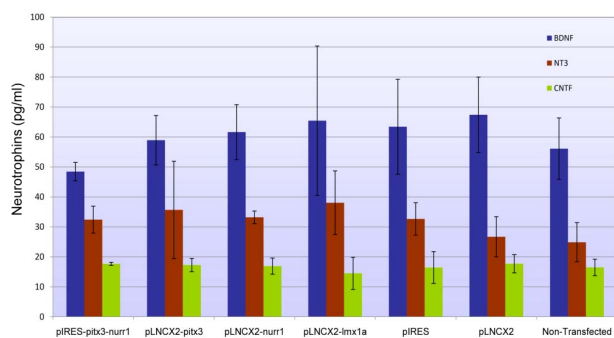
The non(pre)-transfected hONPs were found to produce neurotrophic factors such as BDNF ( $56.09 \pm 10.24$  pg/ml), CNTF ( $18.72 \pm 1.43$  pg/ml) and NT-3 ( $24.87 \pm 6.53$  pg/ml). The stably transfected lines were examined to determine if lineage restriction to dopaminergic neurons alters the synthetic capacity and activity of these neurotrophins; no significant differences in intracellular neurotrophin (BDNF, CNTF, NT-3) levels between transfected and non-transfected hONP lines were observed ( $P >$

0.01), indicating that transfection did not alter neurotrophin synthesis (**Figure 8**).

## 4. DISCUSSION AND CONCLUSIONS

Parkinson's disease, as a neurodegenerative disease, is characterized by loss of specific dopaminergic neurons in substantia nigra [4]. Although a variety of pharmacological agents have been employed in the treatment of PD their effects are transient. "Proof of Concept Studies" with embryonic adrenal medulla cells [54] although ending in failure demonstrated the potential of cell-based replacement therapy. Recently substantial effort has been





**Figure 8.** Histogram demonstrating the neurotrophin levels in hONPs (pg/ml) transfected with pIRES-Pitx3-Nurr1, pLNCX2-Pitx3, pLNCX2-Nurr1, pLNCX2-Lmx1a, pIRES, pLNCX2 and non-transfected NSFCs. Lineage restriction did not alter neurotrophin production.

devoted to the search for a suitable cell source for a cell replacement strategy for the treatment of PD. Many studies have focused on the use of embryonic stem cells; studies utilizing embryonic cells derived from mice or porcine were found to be functional in relieving PD like symptoms in PD animal models [33,55], and positive results obtained from human oriented ES cells further advanced the use and promise of stem cells as a potential source for cell therapy for PD [56-58]. However, these studies were all generally hampered by the significant side effects due to the transplantation of ES cells, such as dyskinesias and/or the formation of teratomas [35,56, 59]. Unfortunately, low cell viability following transplantation, tissue compatibility, a limited of source and ethical concerns further diminish the therapeutic utility of ES cells. In contrast, the use of adult human olfactory epithelium derived progenitors, as a unique autologous cell source, which can be obtained with minimally invasive surgery can avoid these negative factors and also eliminate the need for immunosuppression. The studies described in this manuscript demonstrate that hONPs can be stably lineage restricted under an optimized paradigm, so that they produce and release dopamine, which makes them potential candidates for cell-based therapy for PD. Additionally, the genetic modification didn't alter the capability of hONPs to produce and release key neurotrophic factors, which have the potential to support neuronal survival, as well as rescue degenerating neurons. These factors can also provide permissive micro-environments that may induce endogenous stem cell generation and differentiation [60-62].

In the present study, several conditions have been utilized to optimize the environment for hONPs and facilitate their differentiation to dopaminergic neurons, including genetic modification and treatment with morphogens. Furthermore, hONPs have the unique potential to synthesize and release key neurotrophic molecules which

can have beneficial effects on the survival of dopaminergic neurons as well as the proliferation and differentiation of endogenous stem cell populations. These will all be discussed individually below.

#### 4.1. Pitx3 and Nurr1 Induce the DA Neuron Maturation Synergistically

The Pitx3 gene belongs to the Pitx family of transcription factor genes and has been shown to be required for the expression of TH, the precursor of dopamine, both *in vitro* and in mice from E11.5 [40]. It has been reported that Pitx3 is crucial to the formation of SN and the specification and/or the survival of the subpopulation of the DA neurons in striatum [63-65]. The earlier studies suggest that Pitx3 increased TH promoter induction in mouse and rat cell lines, but not in human cell lines [63, 66]. However, human embryonic stem cells were employed in experiments to demonstrate the regulation of TH expression by Pitx3 [67-69]. These studies suggested that pitx3 is a key transcriptional regulator of genes required specifically for the mesencephalic dopaminergic (mesDA) phenotype [69,70] and for TH expression [40, 64]. Nurr1 is a member of the nuclear receptor super family of transcription factors that is expressed in both developing and mature dopaminergic neurons in the central nervous system in mice [71]. Previous studies have shown that Nurr1 is essential to both survival and differentiation of the ventral mesencephalic dopaminergic precursor neurons [34,72]. Nurr1 has also been reported to be essential in the expression of TH, which is required for DA synthesis; and for vesicular monoamine transporter 2 (VM-AT2), which is related to DA storage; and dopamine transporter (DAT), which is crucial for DA re-uptake [72]. In addition, a recent study has shown that Nurr1 plays a previously unexpected role in protecting TH positive neurons from neurotoxicity [73]. Furthermore, Nurr1 is the only known transcription factor that is associated with the dopaminergic neurotransmitter identity in mesDA neurons [71]. Therefore, both Pitx3 and Nurr1 have been shown to be crucial to the formation of SN and the specification and/or the survival of the DA neurons in midbrain in rodents [39,74,75]. The results obtained in the present study indicate that overexpression of Pitx3 and/or Nurr1 promotes the expression of DA neuron marker, TH in human adult olfactory epithelial-derived progenitors *in vitro*. HONP lines that were stably transfected with Pitx3 and/or Nurr1 and selected for 4 months, remained healthy and TH positive following 6 months cryostorage in liquid nitrogen. Furthermore, the direct detection of dopamine production was also evaluated. Lysates of Pitx3 or Nurr1 transfected hONPs

were dopaminergic as determined by dopamine E.I.A. These results suggest that the transcription factors, Pitx3 and Nurr1, not only function as a dopaminergic promoters in chick, mouse, or human embryonic cells [41,68,71,76], but also can participate in dopamine production in adult human olfactory-derived progenitors. Based on previous studies which focused on the regulatory function of Pitx3 and Nurr1 in dopaminergic neuron promotion [63,68,70,72,74,77] and the studies described in this manuscript, we hypothesized that Pitx3 and Nurr1 may collaborate to induce a higher efficiency of dopamine production in midbrain DA neuron maturation. Previously a synergistic effect between Pitx3 and Nurr1 on TH expression has been reported, which appeared to be species dependent occurring in human but not in embryonic murine stem cells [66-78]. The current studies demonstrate that the simultaneous transfection of Pitx3 and Nurr1 into the hONPs produced higher levels of TH expression and dopamine production than transfection of either of the individual genes. We evaluated the effect of transfection on the level of the precursor (TH) and final intracellular and extracellular dopamine levels to confirm and compare the efficiency of the different transfected hONP lines. Therefore, our data, in combination with published reports in rodents [79,80] and human embryonic stem cells [67,81], indicate that Pitx3 and Nurr1 cooperatively induce the maturation of DA neurons. We extend the previous studies to show the feasibility of genetic modification of adult human olfactory-derived progenitors to promote the generation of DA neurons. These studies demonstrate that the co-expression of Pitx3 and Nurr1 will enhance significantly the lineage restriction of adult human progenitors toward dopaminergic neurons which can be employed in cell-replacement paradigms for the treatment of PD.

#### 4.2. Treatment of hONPs with Morphogens Enhances Intracellular and Extracellular Dopamine Levels

Human adult epithelial derived progenitors have the potential to differentiate along several neural lineages in response to morphogenic signals *in vitro* [82]. For example, 11.6 ( $\pm$ 1.5) % of hONPs expressed TH following a 7 day treatment of RA1FN5Shh (1  $\mu$ M RA, 5  $\mu$ M FN and 15 nM Shh), indicating that a dopaminergic lineage can be driven by exposure to these morphogens [53]. Sonic hedgehog (Shh), (RA) and Forskolin (FN) have all been shown to be crucial developmental factors that regulate neuronal specification and differentiation [83-88]. Shh has been shown to be required for the generation of ventral midbrain motor neurons [89,90] as well as dopaminergic neurons in rodents [56,58,75] and chick em-

bryos [59]. This study suggests that Shh increases the expression of TH and that the purity of Shh is an important determinant of TH expression. RA regulates neuronal differentiation in embryonic stem cells [91,92] and adult human neuronal progenitors [93, 94]. RA has several pathways through which it can effect cellular differentiation [95,96]. FN is an adenylyl cyclase activator that increases intercellular levels of cAMP that can stimulate axonal elongation [85,86] and induce embryonic rat/mouse motor neuron survival [97,98]. Following the treatment of RA and FN, the progenitor nature of hONPs is diminished, as characterized by a loss of nestin expression, and the presence of more mature neuronal markers. In this study, a combination of highly purified Shh, RA and FN was applied to the lineage restricted hONPs. The intracellular level of dopamine was demonstrated to be significantly increased by this treatment. This result confirms and extends the published data by showing that these morphogens can increase TH expression by progenitors obtained from adult humans [53]. Furthermore, following a 4 day treatment of RA1FN5-Shh, the dopamine level of the spent conditioned medium was significantly enhanced, indicating that the morphogens promoted the release of dopamine, which is important for future studies transplanting lineage restricted hONPs into PD animal models. Among all 4 lineage restricted hONP lines, those cells transfected with pIRES-Pitx3-Nurr1 produced and released the highest levels of dopamine in the presence of Shh, RA and FN. This result is consistent with the analysis of the lineage restricted cells in the absence of treatment with the morphogens. This data further supports the conclusion that hONPs transfected with pIRES-Pitx3-Nurr1 are the most efficient line in dopamine production studies to date, and therefore are likely candidates for engraftment into an animal model of PD. Shh is secreted by the notochord and floor plate at early stage of development [99], RA is detectable in the mid-brain of chick and mice embryos [100], and FN is highly concentrated in the rat substantia nigra [101]. The local distribution of these morphogens *in situ* should influence the engrafted hONPs and may further support their survival and dopamine release following transplantation. The higher level of dopamine released following Shh, RA and FN treatment suggests their potential utility for cell-replacement therapy for PD. Previous studies on the non-human primate PD models, demonstrated that the transplanted responsive human embryonic progenitor cells were still capable of differentiation to DA phenotype within the micro-environment around the lesioned adult host SN, an unexpected finding was that the engraftment also up-regulated an endogenous progenitor population [12]. The results of our studies utilizing a paradigm that combines transfection and morphogen induced lineage modulation highlight the potential therapeutic utility of olfactory epithelial-derived neural progenitors

as an autologous cell source for cell-based replacement and regenerative strategies for patients with Parkinson's disease.

### 4.3. Lineage Restricted hONPs Retain Their Capability to Produce Neurotrophic Factors

It's been reported that neurotrophins such as BDNF, CNTF and NT-3 are crucial in the recovery of primate and rodent models of Parkinson's disease [12,102]. BDNF is a member of the neurotrophin family which supports the maturation and survival of dopaminergic neurons in substantia nigra [44,103]. In the presence of BDNF, more TH positive cells can be found in cultures of ventral mesencephalic tissue than in the absence of the neurotrophin [103,104]. NT-3 belongs to the same family of neurotrophins as BDNF, and has been shown to play a protective role in the degeneration of adult central noradrenergic neurons *in vivo* [105,106]. CNTF has been reported to rescue the degenerating striatal neurons in primate and rodent models [45,107]. Furthermore, the absence of CNTF leads to the apoptosis of motor neurons in adult mice [46,108]. Collectively these studies strongly suggest an important role for these neurotrophins in future therapeutic strategies for neurodegenerative diseases, including PD, Alzheimer's disease and Huntington disease. Therefore, a cell population that can produce neurotrophins could be an ideal for therapy for these diseases. They can provide protective micro-environments *in vivo* and prevent, rescue and or replace neuronal degeneration. The pre-transfected hONPs were found to produce several neurotrophins including BDNF, NT-3, and even nerve growth factor (NGF) when in a serum enriched medium [10]. The stably transfected lines were examined to determine if lineage restriction to dopaminergic neurons or absence of serum alters the synthesis of these neurotrophins since they play a role in neuronal survival, differentiation and maturation. As shown in the results, the transfection of hONPs did not alter neurotrophin production. The post-transfected hONPs produce BDNF, NT-3 and CNTF at equivalent levels with the pre-transfected progenitors. Therefore, genetically modified hONPs can not only serve as replacements of the dead or dysfunctional dopaminergic neurons but also can provide protective micro-environments to help rescue dying or damaged neurons from further degeneration and to enhance the endogenous progenitor populations. The stably lineage restricted hONPs are unique populations with high potential for cell transplantation for animal models of Parkinson's disease.

The long term goal of this study is to develop restricted hONP lines that will have therapeutic utility in cell replacement strategies for patients with PD. The *in vivo* viability and stability are important variables, especially

considering the likelihood that with time the engrafted population may die and require replacement. Therefore, experiments were undertaken to determine the stability and viability of frozen stocks of transfected cells. hONPs survived under the pressure of selection after removal from cryostorage and retained their ability to express TH, as well as produce and release dopamine and neurotrophins, which further demonstrates the unique potential of these progenitors to perhaps serve as an autologous cell source for cell-based strategies for the long-term treatment of Parkinson's disease.

Human adult olfactory epithelial-derived progenitors may provide a unique autologous cell population for cell-based therapy of Parkinson's disease, because of their potential to become dopaminergic neurons which produce and release dopamine and their capability to provide neurotrophic factor enriched micro-environments which support cell survival, protect cells from degeneration and activate endogenous stem cell populations. *In vivo* studies are in progress to determine the ability of hONPs to diminish Parkinson like locomotory deficits in a rodent model.

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