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# Dorsal root ganglion progenitors differentiate to gamma-aminobutyric acid- and choline acetyltransferase-positive neurons<sup>☆</sup>

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## Abstract

This study examined the isolation and differentiation of dorsal root ganglion progenitor cells for therapeutic use in neurodegenerative diseases. Rat embryonic dorsal root ganglia progenitors were isolated and purified using the differential adhesion method combined with cytosine arabinoside treatment. After culture in serum-free medium supplemented with B27, basic fibroblast growth factor and epidermal growth factor, these cells remained viable and survived for more than 18 months *in vitro*. Most cells differentiated to neurons that were immunoreactive for gamma-aminobutyric acid and choline acetyltransferase as detected by immunohistochemical staining. In addition, nerve growth factor and neurotrophic tyrosine kinase receptor expression were also observed in dorsal root ganglion progenitors and differentiated cells. K252a, an inhibitor that blocks nerve growth factor-induced signaling, inhibited cell survival, suggesting the possible existence of a nerve growth factor autocrine loop in these proliferating cells.

**Key Words:** dorsal root ganglion; neural progenitor; differentiation characterization; nerve growth factor; tyrosine kinase receptor type 1

**Abbreviations:** DRG, dorsal root ganglion; MAP2, microtubule-associated protein 2; CNP, cyclic nucleotide phosphodiesterase; ChAT, choline acetyltransferase; GABA, gamma-aminobutyric acid; NGF, nerve growth factor

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## INTRODUCTION

Neural stem cells or progenitors exist in the nervous system<sup>[1]</sup> and various non-neural peripheral tissues such as gut<sup>[2]</sup>, skin<sup>[3]</sup>, connective tissue<sup>[4]</sup> and even heart<sup>[5]</sup> of embryonic or adult mammals. Studies to isolate and characterize such progenitors have attempted to purify particular phenotypes with which to generate pluripotent cell lines, with the objective of increasing the efficacy of therapy with grafts of neuronal precursors or mature neurons into injured neurological systems to replace degenerated neurons such as acetylcholine neurons in Alzheimer's disease and dopaminergic neurons in Parkinson's disease<sup>[6]</sup>. Many sensory organ diseases resulting from aging or wounding are usually accompanied by sensory nerve fiber loss or sensory neuron degeneration. Thus, it is important to isolate sensory neural progenitors or stem cells for cell replacement therapy of sensory organ diseases<sup>[7]</sup>. An early study demonstrated that allotransplant and xenotransplant of fetal

dorsal root ganglion (DRG) neurons could survive and reinnervate the denervated host peripheral targets in a variety of locations in a gangliectomized adult rat<sup>[8]</sup>. In addition, xenografts of human fetal DRGs could extend axons into the central nervous system to form functional connections in the deafferented rat spinal cord<sup>[8-9]</sup>. Although neural stem/progenitor cells have been isolated from postnatal and adult DRG<sup>[10-12]</sup>, adult otic placode-derived spiral ganglion<sup>[13]</sup> and inner ear<sup>[14]</sup>, the differentiation status and neurotransmitter phenotype have not been characterized. In addition, embryonic stem cell-derived neuronal precursors can survive and differentiate to glia and neurons after central nervous system transplantation<sup>[15-16]</sup>. In the present study, DRG progenitors were isolated and purified from embryonic day 17 rats to investigate their proliferation and differentiation capacities. We hope that our study will be helpful for understanding neurogenesis in the peripheral nervous system and for harnessing the potential application of peripheral neural stem cells for the treatment of sensory organ diseases.

## RESULTS

### Purification of neuronal progenitors from embryonic day 17 DRG

To obtain a high yield of neurons from DRG neuronal progenitors, cells from embryonic day 17 rat DRGs were purified using the differential adhesion method combined with 0.5  $\mu$ M cytosine arabioside treatment. Cultures comprising approximately 90% neurons were obtained after purification (Figure 1).

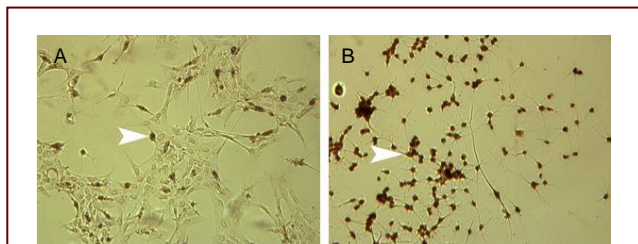
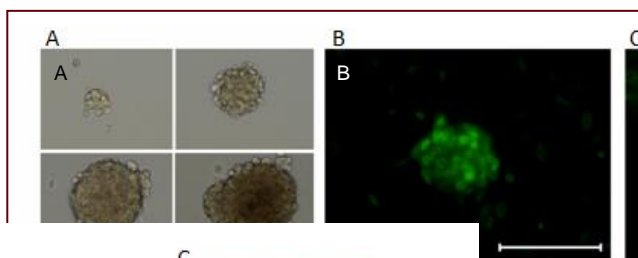


Figure 1 Dorsal root ganglion (DRG) neuronal progenitors before (A) and after (B) purification (magnification  $\times$  200).

Cells were plated on coverslips and purified by the differential adhesion method combined with 0.5  $\mu$ M cytosine arabioside treatment, after which the cells were fixed and immuno-labeled with a monoclonal anti-microtubule-associated protein 2 antibody.

Positive staining was revealed by the avidin-biotin-peroxidase complex method. Arrows show microtubule-associated protein 2 positive cells.

### Progenitors derived from embryonic day 17 DRG can be expanded and passaged long term (Figures 2 and 3)



(A) Formation of spheres from purified DRG neurons; (B) spheres stained for incorporated 5-bromodeoxyuridine (green); (C) spheres stained for stem/progenitor cell marker nestin and revealed by fluorescein isothiocyanate-conjugated secondary antibody (green).

Samples were visualized via fluorescent microscopy. Scale bars represent 25  $\mu$ m.

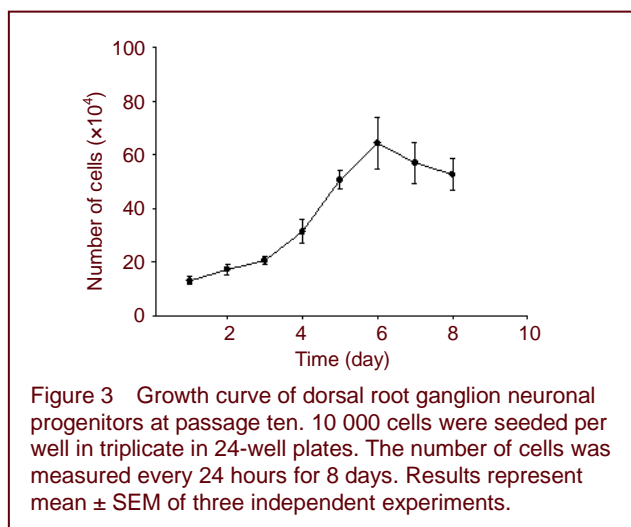


Figure 3 Growth curve of dorsal root ganglion neuronal progenitors at passage ten. 10 000 cells were seeded per well in triplicate in 24-well plates. The number of cells was measured every 24 hours for 8 days. Results represent mean  $\pm$  SEM of three independent experiments.

To assess the ability of progenitor cells to proliferate, primary spheres were dissociated to a single-cell suspension, and new or secondary spheres were obtained between 1 to 2 weeks (Figure 2). Bromodeoxyuridine (BrdU) incorporation and subsequent immunostaining for BrdU and nestin revealed that neuronal progenitors existed in the embryonic DRG neurons (Figure 2). Cells maintained in culture for periods of more than 1 year retained their dependency on epidermal growth factor and basic fibroblast growth factor. In the early stage of culture (1–5 passages), there was a moderate increase in cell numbers. After passage 5, the growth rate increased, followed by stabilization at passage 15. The growth curve was measured at passage 10, and demonstrated that a logarithmic growth phase occurred 4 to 6 days after sub-culturing, and that the saturated phase occurred on

### Is aged and died (Figure 3). G progenitors after differentiation microtubule-associated protein 2 nucleotide phosphodiesterase

type of cells derived from the DRG were stained for MAP2, glial fibrillary acidic protein (GFAP), and nestin. We observed that most cells were stained for MAP2, with a high proportion being GFAP positive cells, whereas no glial fibrillary acidic protein positive cells were observed. CNP is highly expressed in myelin-forming glial cells and is widely used as an immunohistological marker for oligodendrocytes<sup>[17]</sup>. However, in the present study, CNP was present in patches throughout the soma and dendrites of cells, and significantly co-localized with MAP2. This expression pattern suggested an important role for CNP in the neuronal progenitors (Figure 4). Western blot analysis confirmed the immunocytochemical findings (Figure 5).

### DRG progenitors differentiate to cells co-expressing choline acetyltransferase (ChAT) and gamma-aminobutyric acid (GABA)

To assess the progenitor cell neuronal subtypes, we screened for the presence of neurotransmitters using

antibodies specific for tyrosine hydroxylase (TH), GABA, ChAT and 5-hydroxytryptamine/serotonin (5-HT). We observed both ChAT and GABA positive reactivity in the

majority of the cells examined. However, 5-HT and TH were undetectable (Figure 4). Immuno-labeling analysis implied the co-localization of GABA and ChAT<sup>[4]</sup>.

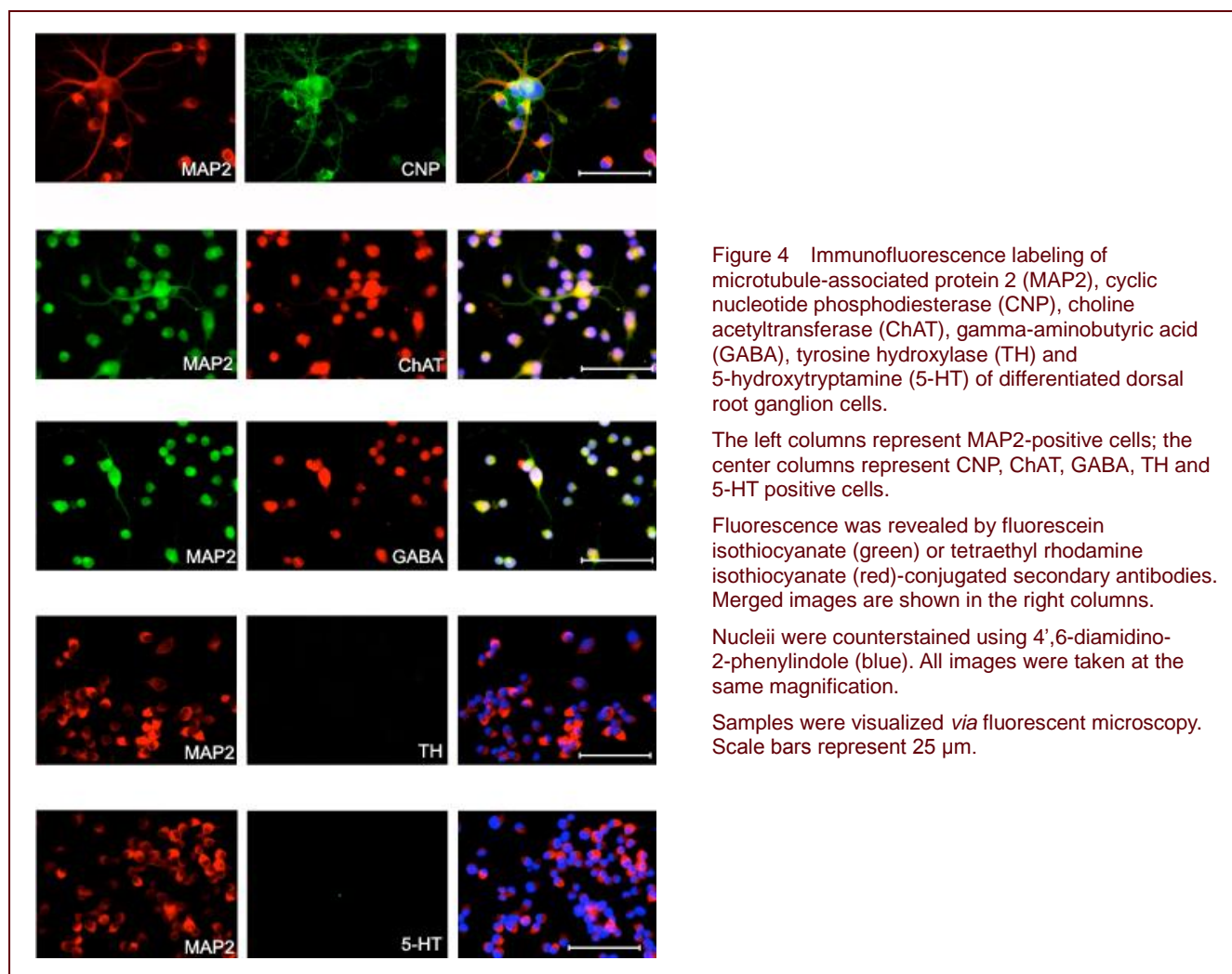


Figure 4 Immunofluorescence labeling of microtubule-associated protein 2 (MAP2), cyclic nucleotide phosphodiesterase (CNP), choline acetyltransferase (ChAT), gamma-aminobutyric acid (GABA), tyrosine hydroxylase (TH) and 5-hydroxytryptamine (5-HT) of differentiated dorsal root ganglion cells.

The left columns represent MAP2-positive cells; the center columns represent CNP, ChAT, GABA, TH and 5-HT positive cells.

Fluorescence was revealed by fluorescein isothiocyanate (green) or tetraethyl rhodamine isothiocyanate (red)-conjugated secondary antibodies. Merged images are shown in the right columns.

Nucleii were counterstained using 4',6-diamidino-2-phenylindole (blue). All images were taken at the same magnification.

Samples were visualized *via* fluorescent microscopy. Scale bars represent 25  $\mu$ m.

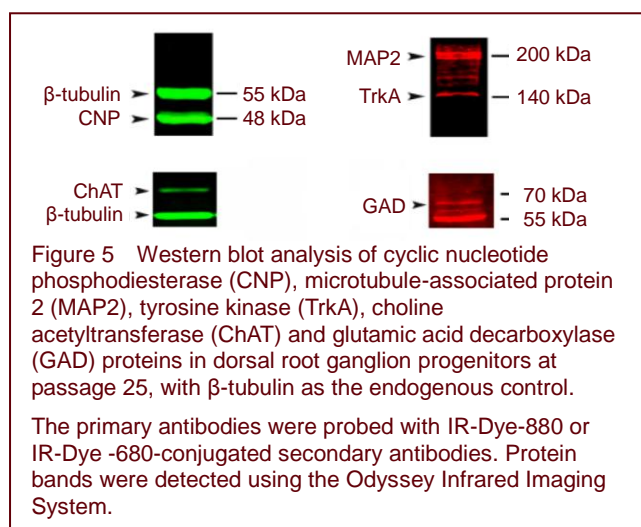


Figure 5 Western blot analysis of cyclic nucleotide phosphodiesterase (CNP), microtubule-associated protein 2 (MAP2), tyrosine kinase (TrkA), choline acetyltransferase (ChAT) and glutamic acid decarboxylase (GAD) proteins in dorsal root ganglion progenitors at passage 25, with  $\beta$ -tubulin as the endogenous control.

The primary antibodies were probed with IR-Dye-880 or IR-Dye -680-conjugated secondary antibodies. Protein bands were detected using the Odyssey Infrared Imaging System.

**Tyrosine kinase (TrkA) and nerve growth factor (NGF) expression in DRG progenitors and differentiated neural cells**

NGF is essential for the survival, differentiation, and

maintenance of many sensory neurons. NGF induces biological effects through binding with receptor tyrosine kinase, TrkA, or the death-like domain containing receptor, p75<sup>NTR</sup>, or both depending on their presence on a given cell and their surface localization. NGF activates TrkA and/or p75<sup>NTR</sup>, triggering signal transduction cascades. Most DRG neurons require NGF for survival in early development<sup>[18]</sup>. We maintained and analyzed DRG progenitors *in vitro* for more than 1 year without adding exogenous NGF to the culture medium. As DRG cells can express NGF following injury *in vivo*<sup>[14]</sup>, NGF and TrkA may also be involved in the proliferation and differentiation of the progenitor cells. The expression of TrkA and NGF was evaluated in progenitors, and differentiated cells by immunofluorescence. We observed that the majority of the cells were stained positive for TrkA and NGF (Figure 5). Western blot analysis confirmed the expression of TrkA protein, but not NGF protein in those cells (Figure 6), possibly indicating that NGF was either expressed at a very low level or was being rapidly degraded.



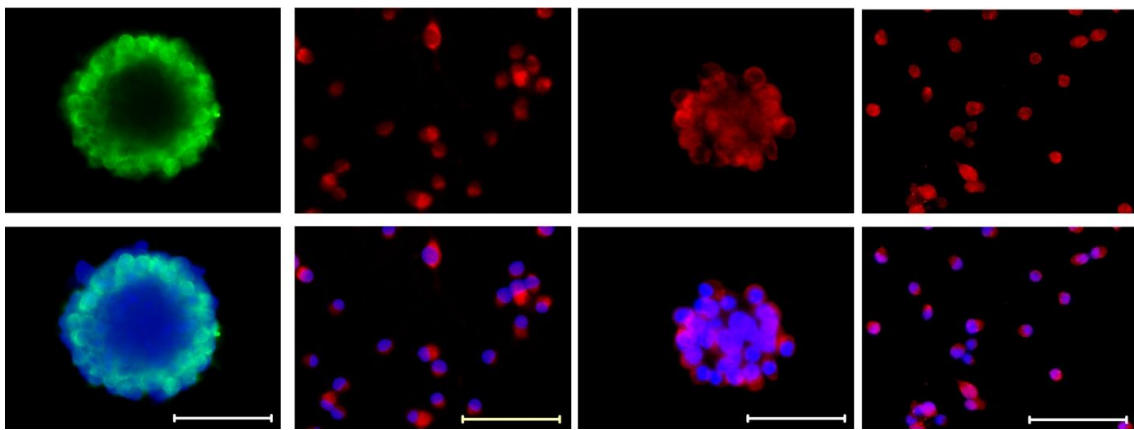


Figure 6 Immunofluorescence labeling for tyrosine kinase (TrkA) and nerve growth factor (NGF) in proliferating and differentiated dorsal region ganglion (DRG) cells.

The left columns represent TrkA-positive cells; the second columns represent TrkA-positive cells in differentiated DRG cells; the third columns represent NGF-positive cells in proliferating DRG cells; and the fourth columns represent NGF-positive cells in differentiated DRG cells.

The fluorescence was revealed by fluorescein isothiocyanate (green) or tetraethyl rhodamine isothiocyanate (red)-conjugated secondary antibodies. Nuclei were counterstained with 4',6-diamidino-2-phenylindole (blue). All images were taken at the same magnification. Samples were visualized via fluorescent microscopy. Scale bars represent 50  $\mu\text{m}$ .

As abundant TrkA and NGF were detected in differentiated DRG neurons by immunofluorescence, we investigated whether endogenous NGF produced by the cells was required for their maintenance. K252a, an inhibitor which blocks NGF-induced signaling in PC12 cells<sup>[19-21]</sup> was added to the culture medium. Survival rates of the cells were diminished with increasing concentrations of K252a (Figure 7). This result implied that NGF might have a key role in supporting the survival and function of DRG progenitors.

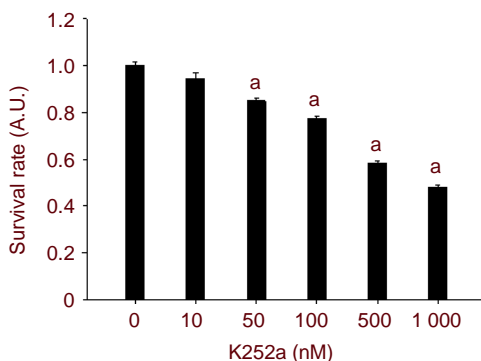


Figure 7 K252a, an inhibitor that blocks nerve growth factor-induced signaling, attenuates the survival of dorsal region ganglion (DRG) progenitors. DRG progenitors were maintained in differentiation medium at 24 hours before incubation with K252a at increasing concentrations.

A cell counting kit-8 assay was used to evaluate cell viability. Data are presented as the percent viability of vehicle-treated control cells. Values are expressed as the mean  $\pm$  SEM of three independent experiments.

Fisher's least significant difference test was used to analyze the statistical significance of the differences. <sup>a</sup> $P < 0.01$ , vs. untreated cells (0 nM) K252a.

## DISCUSSION

Previously, several *in vitro* systems have been described for the derivation of neural stem/progenitor cells from the central nervous system or peripheral nervous system<sup>[22]</sup>. The entire peripheral nervous system is derived from a migratory cell population termed neural crest cells. These cells generate a wide variety of cell and tissue types during embryonic and adult development including cartilage and bone, connective tissue, pigment and endocrine cells as well as neurons and glia amongst many others. Due to these specific properties they have been studied for their potential application in cell-based tissue and disease-specific repair<sup>[23]</sup>. DRGs are derived from precursors in the neural crest, suggesting that early postnatal DRGs may contain a population of neuronal precursors that retain their capacity for neurogenesis. In the present study, we report the purification of a DRG neuronal stem/progenitor cell, and the further characterization of proliferation and differentiation of these cells.

### Progenitors derived from embryonic DRGs can be expanded long term

DRG cells from embryonic day 17 rats were purified using the differential adhesion method followed by treatment with cytosine arabinoside that causes the selective removal of glial cells. After purification, cells were cultured in serum-free medium DMEM/F12 (1:1) supplemented with B27, basic fibroblast growth factor and epidermal growth factor. Cells proliferated slowly in the first 2 to 3 weeks. After this time point, neurospheres were observed and new spheres were generated after each passage. These cells were routinely passaged once every 1 to 2 weeks depending on the density

seeded. The growth curve of the 15<sup>th</sup> passage cells demonstrated that progenitors from embryonic DRGs could proliferate efficiently. Cells were maintained in culture for more than 1 year and retained their potential for proliferation and differentiation as specialized subtypes. Such long-term proliferation was unexpected and to our knowledge has not been reported previously. The incorporation of BrdU, together with positive nestin immunofluorescence, suggested that the purified cells from embryonic DRG were proliferating<sup>[24]</sup>. Thus, we termed these cells DRG progenitors.

#### **DRG progenitors exhibit characteristics similar to neural precursors**

To investigate the differentiation characteristics of embryonic DRG progenitors, cells were incubated in culture medium with serum and without exogenous basic fibroblast growth factor and epidermal growth factor. Most DRG progenitors expressed MAP2, 40% of which were positive for both MAP2 and CNP. CNP is also present in various cell types in addition to myelinating cells, such as lymphocytes, retinal, liver, muscle, and Purkinje cells and hippocampal neurons<sup>[25-27]</sup>, indicating that CNP is also expressed in some subpopulations of neuronal cells. CNP is a regulator of tubulin polymerization, where it associates with the cytoskeleton and has microtubule-associated protein-like characteristics<sup>[28]</sup>. Taken together and combined with our findings, these results suggest that CNP may be important in the modulation of the cytoskeleton in the differentiating DRG progenitors.

In addition to glial cells, mature DRGs are composed of many neurons with different morphologies and distinct biochemical properties. How distinct cell fates are generated from an initially homogeneous cell population in the embryonic DRG is a compelling question in developmental biology. Moreover, once DRG precursors aggregate to their final positions, there are still a number of "fate choices" that can occur<sup>[29]</sup>. The sensory neurons present in mature DRG receive sensory information including pain, temperature, touch and proprioception. Cells in DRG produce multiple neurotransmitters, such as GABA, acetylcholine, and glutamate catecholamine. Tyrosine hydroxylase is expressed in a subpopulation of small DRG neurons in the adult mouse<sup>[30]</sup>.

Since therapeutic applications may require considerable *in vitro* expansion of neural precursors, we next investigated whether the expanded, multi-passage DRG progenitors retained key neural precursor properties. The purified DRG cells formed a subpopulation of neuronal progenitors rather than glial precursors. These cells could proliferate, but also expressed GABA and ChAT simultaneously while undergoing differentiation. Cholinergic neurons have been reported to be immunoreactive for either GABA or its synthesizing enzyme<sup>[31]</sup>. Furthermore, in the rat cerebral cortex, 88% co-localization of ChAT with GABA was observed in interneurons<sup>[32]</sup>. Thus, we assumed that the subpopulations of DRG progenitors mainly differentiated

to neurons coexpressing acetylcholine and GABA. However, no TH and 5-HT containing cells were detected.

#### **DRG cells express NGF and its receptor TrkA**

*In vivo*, NGF signaling is required for both survival and the differentiation of DRG neural phenotypes<sup>[33]</sup>. Embryonic DRG neurons initially require NGF for survival *in vitro*. Withdrawal of NGF from DRG neurons isolated at embryonic day 15 within the first 10 days of culture, resulted in apoptotic death. However, by 21 days of culture, the majority of these neurons survived for long periods without exogenous NGF support<sup>[34-35]</sup>. Recent studies reported that NGF mRNA is expressed in a variety of cell types in the injured spinal cord. NGF mRNA is also up-regulated in DRG neurons after spinal cord injury and the percentage of sensory neurons expressing NGF mRNA correlates with proximity to the lesion epicenter. This suggests that NGF expression in DRG may be up-regulated by damage to the central processes of sensory neurons<sup>[36]</sup>. In addition, it is likely that DRG neurons will express NGF when stimulated or under special conditions. In this study, the progenitors from embryonic day 17 DRG that were either proliferating or had differentiated were independent of exogenous NGF *in vitro*. We observed NGF and TrkA expression in both proliferating and differentiated cells. However, p75<sup>NTR</sup> positive cells were not detected (data not shown). In addition, incubation with K252a attenuated cell survival and implied that DRG progenitors expressed NGF for their survival and function. To conclude, we isolated a subpopulation of DRG neural progenitors, which could be induced to differentiate to neurons co-expressing ChAT and GABA. These cells also expressed TrkA and NGF. These novel DRG progenitor cells represent a useful tool for studying the mechanisms of cell proliferation and differentiation, and have potential application as peripheral neural stem cells for the treatment of sensory organ diseases.

## **MATERIALS AND METHODS**

### **Design**

A parallel controlled *in vitro* experiment.

### **Time and setting**

The experiments were performed at the Laboratory of Molecular and Cellular Neurophysiology, East China Normal University from September 2007 to April 2011.

### **Materials**

Healthy adult Sprague-Dawley rats, weighing 150–200 g, were provided by the Experimental Animal Research Center of Shanghai, China. All rats were maintained in air-conditioned quarters under regulated light and dark periods. For mating, two to three females and two vigorous males were caged together. The vaginal canal was examined in the morning for the presence of sperm or vaginal plug on subsequent days. The day sperm or vaginal plug was found in the vagina was considered as

embryonic day 0.5. At embryonic day 17.5, the mother was sacrificed, and the embryos were removed for dissection. All experimental procedures were carried out in accordance with the guidelines of the National Institutes of Health on animal care.

## Methods

### **Cell culture and growth curve analysis**

DRG neurons were isolated and purified using the differential adhesion method combined with treatment with 0.5  $\mu$ M cytosine arabinoside. Briefly, DRG from all spinal levels were removed, and incubated in pre-warmed Hank's buffered salt solution containing 0.25% trypsin (Invitrogen, Grand Island, NY, USA) for 20 minutes at 37°C. After enzymatic treatment, a single cell suspension was prepared by passing cells through a fire polished Pasteur pipette approximately 15 times, and then cells were plated in non-coated 6-well plates for 50 minutes to remove non-neuronal cells. The unattached cells were collected and cultured in serum-free Dulbecco's modified eagle's medium/F12, supplemented with 2% B27 (Invitrogen), and 0.5  $\mu$ M cytosine arabinoside (Sigma, St. Louis, MO, USA) was added after 12 hours. Cells were then cultured in proliferation medium Dulbecco's modified Eagle's medium/F12 supplemented with 2% B27, 20 ng/mL epidermal growth factor, 10 ng/mL basic fibroblast growth factor-2 (all from Gibco, Grand Island, NY, USA). Half of the medium volume was replaced every 3 days. The primary spheres were dissociated into a single-cell suspension, and new or secondary spheres could be obtained between 1 to 2 weeks. Expanded cultures were passaged routinely when new spheres formed and neared confluence (70–80% confluence). To evaluate the proliferation capacity, 10 000 cells were seeded per well in triplicate in 24-well plates. The number of the cells was counted every 24 hours for 8 days. To induce cell differentiation, spheres were mechanically dissociated into single cell suspensions before being placed on poly-L-lysine and laminin coated coverslips in 24-well plates, and differentiation medium supplemented with 2% fetal calf serum was added and refreshed every 3 days.

### **Immunohistochemistry for MAP2, ChAT, 5-HT, GABA, glial fibrillary acidic protein, CNP, TH, NGF, TrkA, and BrdU**

Cells were fixed in 4% paraformaldehyde for 15 minutes at room temperature, and then incubated with antibodies specific for MAP2 (Shanghai Branch of Sigma, Shanghai, China), ChAT (Shanghai Branch of Chemicon, Shanghai, China), 5-HT (Shanghai Branch of Chemicon), GABA (Shanghai Branch of Chemicon), glial fibrillary acidic protein (Shanghai Branch of Sigma), CNP (Shanghai Branch of Sigma), TH (Shanghai Branch of Sigma), NGF (Shanghai Branch of Santa Cruz Biotechnology, Shanghai, China), TrkA (Shanghai Branch of Chemicon) and BrdU (Shanghai Branch of Sigma) overnight at 4°C, followed by incubation with an appropriate fluorescein isothiocyanate or tetraethyl rhodamine

isothiocyanate-conjugated secondary antibody (Shanghai Branch of Chemicon) for 1 hour at room temperature. Cell nuclei were stained with 4',6-diamidino-2-phenylindole (Shanghai Branch of Chemicon). The staining for nestin (Shanghai Branch of Millipore, Shanghai, China) was performed using the streptavidin-biotin-alkaline phosphatase (ABC) method using the ABC kit (Shanghai Branch of Vectastain, Shanghai, China) according to the manufacturer's protocol. Stained samples were visualized *via* fluorescent microscopy (Leica DMI4000B). Images were analyzed using Leica Imager, and brightness and contrast were adjusted using ImageJ software (version 1.38x, China).

### **Western blot method for MAP2, ChAT, GABA, CNP, and TrkA expression**

Briefly, protein samples extracted from DRG cells were separated by 7.5–12% sodium dodecyl sulfate polyacrylamide gel electrophoresis, electrotransferred onto nitrocellulose membranes (Millipore, China), membranes incubated with 5% bovine serum albumin in Tris-buffer saline at room temperature for 1 hour to inhibit non-specific binding, and then incubated with antibodies specific for MAP2 (Sigma, China), ChAT (Chemicon, China), GABA (Chemicon, China), CNP (Sigma, China), TrkA (Chemicon, China),  $\beta$ -tubulin (Santa Cruz Biotechnology) overnight at 4°C. The primary antibodies were probed with IR-Dye-880 or IR-Dye -680-conjugated secondary antibodies (LI-COR Biosciences, USA). Specific protein bands were detected and measured by Odyssey Infrared Imaging System (LI-COR Biosciences).

### **Cell counting kit-8 assay**

DRG cell viability was detected using a cell counting kit-8 assay (Dojido Laboratories, Japan) according to the manufacturer's protocol. Cells were seeded at 5 000 cells/well in 96-well plates. K252a at various concentrations (0, 10, 50, 100, 500, 1 000 nM; Shanghai Branch of Sigma) was added to the culture for 24 hours before cell counting kit-8 detection.

### **Statistical analysis**

Results were expressed as mean  $\pm$  SEM. Analyses were performed using SigmaPlot version 10.0 (Systat Software, Chicago, IL, USA). Fisher's least significant difference was used to analyze the statistical significance of the differences. Results were considered statistically significant when  $P < 0.05$ .

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**Author contributions:** The work presented here was carried out in collaboration between all authors. Lingli Yu and Yindi Ding defined the research theme, designed methods and experiments, performed the laboratory experiments, analyzed the data, interpreted the results and wrote the paper. All authors have contributed to and approved the manuscript.

**Conflicts of interest:** None declared.

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