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however, did not differ significantly between 4-OHT-treated and untreated rats. We roughly estimated the number of transduced cells in the striatum at 5×10^4 based on cell counts performed on the tissue sections. This efficiency of transduction is sufficient to parallel the functional effects on behavior observed in other studies [3,5,11]. Double-labeling with both anti-Cre and anti-TH antibodies, or with anti-Cre and anti-AADC antibodies, showed that more than 80% of Cre-immunoreactive cells were also positive for TH (164 of 200) and AADC (176 of 200) in three 4-OHT-untreated rats (Fig. 4C). Immunostaining for glial fibrillary acidic protein (GFAP) or AADC in parallel sections demonstrated transduction of striatal cells without obvious reactive astrocytosis (Fig. 4D). Dopamine content (Fig. 5A) and TH activity (Fig. 5B) within the lesioned striatum were significantly lower in 4-OHT-treated compared to untreated rats. Dopamine levels in the transduced striatum in the 4-OHT-treated and untreated rats were 0.66 and 4.3%, respectively, those of the unlesioned striatum. Unlike primary dopamine in the nigrostriatal system, which is stored in synaptic vesicles, genetically produced dopamine in the lesioned striatum might be readily metabolized without storage. Since the dopamine level in the lesioned side of AAV-LacZ-injected rats was much lower (0.3%), a 10-fold increase in dopamine level after triple transduction with TH, AADC, and GCH genes caused a remarkable therapeutic effect. Average TH activity, measured in terms of L-dopa production (pmol/min/mg protein), reached 51.6% that of the normal striatum (65.9 ± 11.0 versus 127.7 ± 3.7) in rats transduced with dopamine-synthesizing enzymes. In 4-OHT-treated rats, TH activity fell to 10.8% of normal (13.8 ± 4.1).

Discussion

Our results show efficient viral vector-mediated delivery of tamoxifen-dependent CreER^{T2} recombinase into rodent brains and that transgenic floxed sequences can be deleted in a temporally controlled manner. In a rat model of PD, recombinant AAV vector-mediated delivery of CreER^{T2} into the striatum enabled 4-OHT-induced excision of a floxed TH transgene, resulting in reduced virally mediated dopamine synthesis. We targeted TH, a rate-limiting enzyme for dopamine biosynthesis that converts dietary L-tyrosine to L-dopa. Using this strategy, AADC activity was retained so that L-dopa, a substrate for AADC capable of crossing the blood–brain barrier, could be converted to dopamine in the striatum. Thus, in clinical situations, the therapeutic effects of orally administered L-dopa would likely be preserved, even after 4-OHT treatment to reduce TH expression in cases of dopamine overproduction [11]. Although transduction with AADC alone, in combination with oral administration of L-dopa, might not achieve continuous delivery of L-dopa, in contrast to that which could potentially be achieved with triple transduction of TH, AADC, and GCH, dopamine production could be regulated by altering the dose of L-dopa, thereby providing a safer option for gene therapy. We previously demonstrated that dopamine synthesis was enhanced (greater than fivefold) after systemic administration of L-dopa in AAV-TH/-AADC/-GCH-injected striatum in the primate model of PD using *in vivo* dialysis [4]. A phase I clinical trial involving gene transfer of AADC alone is currently under way.

AAV vectors are powerful tools by which to deliver therapeutic genes into the mammalian brain. Many striatal neurons of rodents and nonhuman primates are transduced with AAV vectors via stereotaxic injection, and long-term gene expression has been achieved without substantial toxicity or immune response [2,12]. AAV vectors have safety advantages over other viral vectors when it comes to *in vivo* gene delivery, since they are derived from nonpathogenic wild-type viruses. Moreover, most recombinant AAVs are present in cells as episomes, thus reducing the probability of insertional activation of oncogenes, compared to retroviruses, which integrate into host chromosomes [13]. Although it is difficult to use a single AAV vector for multiple gene transfer due to its limited packaging capacity (<5 kb), a single cell can be simultaneously transduced with multiple AAV vectors. In the present study, dual immunofluorescence staining showed efficient cotransduction of cells with different AAV vectors in the rat striatum, a finding consistent with that which we observed in a previous study [5].

Gene therapy strategies for the treatment of PD using AAV vectors include gene delivery of dopamine-synthesizing enzymes into the striatum to restore dopamine production, as well as gene delivery of neuroprotective molecules, such as glial cell line-derived neurotrophic factor, to block or slow down further degeneration [14]. In addition, AAV vectors harboring genes encoding neurotrophic factors might be delivered by intramuscular administration in an attempt to protect spinal motoneurons in patients suffering from amyotrophic lateral sclerosis [15–17]. Although no adverse effects due to overexpression of transgenes have been reported in animals to date, it is necessary to develop vectors that allow for regulation of transgene expression, thus avoiding transgene overexpression. In PD, overproduction of dopamine has the potential to cause dyskinesia or hallucinations, and sustained exposure to high concentrations of neurotrophic factors could result in tumor formation.

Inducible Cre recombinases have been used to generate a number of conditional knockout mice. They are invaluable tools for investigators studying the role of gene function in development, as well as a number of physiological and pathological processes. The tamoxifen-dependent CreER^{T2} recombinase has proven particularly helpful [9,18,19]. It has been shown that 4-OHT does not alter dopamine content within the striatum in mice [20],

and we did not observe any adverse effects of 4-OHT treatment in the present experiment. In addition, tamoxifen, which is metabolized by the liver into 4-OHT, has neuroprotective effects [21,22]. Thus, the CreER^{T2} system might be useful for gene therapy in the treatment of neurological diseases.

We have expanded upon the use of inducible Cre recombinase technology to regulate transgene expression using an AAV vector-mediated gene delivery system. Recently, a viral vector-mediated RNA interference (RNAi) approach has been developed and localized gene knockdown achieved in the adult brain [23–25]. Although RNAi-mediated suppression of gene function has a wide variety of applications, more specific and inducible transgene silencing can be achieved with AAV-CreER^{T2}. Selective ablation of floxed transgenes reduces the possibility of down-regulation of normal cellular proteins. This system works as a molecular switch, increasing the safety of long-acting gene therapy by avoiding or minimizing side effects due to overproduction of the protein product and by providing the ability to shut down expression if toxicities are encountered or treatment is completed. The ability to restrict somatic recombination of transgenes spatially and temporally has a wide range of applications, in both gene therapy and biological study requiring somatic genetic manipulation.

Materials and Methods

AAV vector production. The AAV vector plasmids contained an expression cassette with a human cytomegalovirus immediate-early promoter (CMV promoter), followed by the first intron of human growth hormone, target cDNA, and a simian virus 40 polyadenylation signal sequence (SV40 poly(A)), between the inverted terminal repeats of the AAV-2 genome. The plasmids pAAV-LacZ, pEGFP, pAAV-AADC, pAAV-GCH, pCre, and pCreER^{T2} contained the cDNAs of LacZ, EGFP, human AADC, human GCH, Cre recombinase with a nuclear localization signal [26], and Cre recombinase fused to a mutated form of the ligand-binding domain of estrogen receptor α (CreER^{T2})[27], respectively. The plasmid pAAV-floxed TH contained human TH1 cDNA flanked by two loxP sequences between the CMV promoter and the SV40 poly(A). To generate pAAV-EGFP/Red, a DNA fragment containing d2EGFP (BD Biosciences, San Jose, CA, USA) and the SV40 poly(A) was flanked by loxP sequences and inserted between the CMV promoter and DsRed-Express-DR (BD Biosciences). The two helper plasmids, pHLP19 and pladeno1 (Avigen, Alameda, CA, USA), harbored the AAV *rep* and *cap* genes, as well as the *E2A*, *E4*, and *VA RNA* genes of the adenovirus genome, respectively. HEK293 cells were cotransfected by the calcium phosphate coprecipitation method with the vector plasmid, pHLP19, and pladeno1. The AAV vectors were then harvested and purified by two rounds of continuous iodoxale ultracentrifugations. Vector titers were determined by quantitative DNA dot-blot hybridization or by quantitative PCR of DNase I-treated vector stocks. We routinely obtained 10^{12} to 10^{13} vector genome copies (vg).

In vitro transduction. HEK293 cells were seeded at 3×10^5 cells/well in six-well plates. After 24 h, cells were infected with appropriate combinations of AAV vectors (5×10^9 vg per vector). 4-OHT was added to the culture medium at a concentration of 10 or 100 nM at 5 h after infection. Culture medium was collected for dopamine assay 48 h after infection and the cells were fixed for immunostaining.

AAV injections. All animal experiments were performed in accordance with the institutional guidelines. Three B6,129-*Gt(ROSA))26Sor*^{tm2Sho}/J mice (The Jackson Laboratory, Bar Harbor, ME, USA) [10] were stereotaxically injected into the caudoputaminal unit or cerebral cortex with 1×10^9 vg (1 μl) of AAV-CreER^{T2}. After 1 week, 4-OHT (1 mg) was administered intraperitoneally every day for 5 days, after which 2 of the mice were killed. One mouse that did not receive 4-OHT treatment was killed as a control. Creation of PD model rats and stereotaxic injections of AAV vectors were carried out as previously described [5]. Briefly, 60 male albino Wistar rats (weighing 200–250 g) were unilaterally lesioned at the left medial forebrain bundle (coordinates AP − 4.3 mm and ML 1.6 mm, relative to the bregma, and DV − 7.8 mm relative to the dura, with the incisor bar set 3.3 mm below the interaural line) with 4 μl of 4.5 mg/ml 6-OHDA HBr (Sigma, St. Louis, MO, USA) in 0.02% ascorbate saline prior to intrastriatal transduction. These rats were stereotaxically injected with AAV vectors (5×10^7 vg per site for each vector) at three sites in the lesioned striatum (coordinates relative to the bregma and dura, AP +1.5, +1.0, and +0.5 mm; ML 2.6, 3.0, and 3.2 mm; DV − 5.2 mm). Forty-eight rats were injected with a 1:1:1 mixture of AAV-floxed TH, AAV-AADC, and AAV-GCH plus AAV-Cre (n = 12) or AAV-CreER^{T2} (n = 36). Twelve rats received AAV-LacZ alone as a control. Among the AAV-CreER^{T2}-treated rats, 24 were intraperitoneally injected with 4-OHT (4 mg/kg) for 5 consecutive days, starting either at the same time as or 4 weeks after vector injection.

Behavioral testing. The rats were tested weekly for rotational behavior and spontaneous limb use, as described previously [28]. The total number of complete body turns was counted during an observation period of ≥ 60 min following intraperitoneal injection of apomorphine HCl (0.1 mg/kg; Sigma). Only those animals exhibiting seven or more contralateral rotations/min in a 60-min period at 4 weeks after the 6-OHDA injection were included in further analysis. Spontaneous limb use was scored according to the cylinder test method [29]. Rats were placed in a clear glass cylinder large enough to ensure free movement. After they had performed 10 rears during which they were observed to place at least one paw on the cylinder wall, the number of times both forepaws contacted the wall of the cylinder was counted until at least 20 contacts were made. Data indicating the number of times a contralateral forepaw made contact with the wall are expressed as a percentage of the total. We also evaluated rotational behavior in response to a low dose of L-dopa methyl ester (5 mg/kg; Sigma) coadministered with 2.5 mg/kg of a peripheral decarboxylase inhibitor (benserazide hydrochloride; Sigma) 10 weeks after AAV injection.

Biochemical assays. Levels of dopamine in the cell culture medium (n=4 for each group) and within the brain samples (n=4 for each group) were determined by high-performance liquid chromatography (HPLC), as previously described [5]. Rats were killed by decapitation under sodium pentobarbital anesthesia 12 weeks after vector injection, after which their brains were immediately dissected and placed on dry ice. The striatum was punched out bilaterally using a sharp-edged, stainless steel tube. Wet tissue samples were weighed and stored at −80°C until subsequent analysis. Tissues were homogenized in 20 volumes of homogenization buffer and then mixed immediately with 0.76 M perchloric acid prior to centrifugation at 15,000*g* for 10 min. After the supernatant was neutralized with sodium acetate, the samples were analyzed by HPLC analysis. Determination of TH activity was based on the formation of L-dopa from L-tyrosine, as demonstrated by HPLC electrochemical detection. The reaction mixture contained 200 mM sodium acetate buffer (pH 6.0), 100 mM 2-mercaptoethanol, 0.2 mg/ml catalase, 0.2 mM L-tyrosine, and 1 mM tetrahydrobiopterin. The mixture was incubated for 10 min at 37°C. The reaction was stopped by adding perchloric acid, and L-dopa was extracted using an alumina column [30].

Immunostaining of cultured cells and brain sections. Cultured cells were fixed in 4% paraformaldehyde (PFA) in PBS. Brains were perfused with 4% PFA, soaked in 30% sucrose, and dissected into coronal sections (30 μm). The following primary antibodies were used: TH monoclonal (1:800 or 1:8000; DiaSorin, Stillwater, MN, USA) or polyclonal (1:10,000; provided by Ikuko Nagatsu, Fujita Health University, Japan), AADC polyclonal (1:10,000; I. Nagatsu), Cre recombinase monoclonal (1:500; Covance, Princeton, NJ, USA) or polyclonal (1:500; Covance), GFP polyclonal (1:200; BD Biosciences or Chemicon, Temecula, CA, USA), DsRed

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polyclonal (1:1000; BD Biosciences), and GFAP (1:1000; Chemicon). Appropriate fluorescence-tagged (Invitrogen, Carlsbad, CA, USA) or biotinylated (Vector Laboratories, Burlingame, CA, USA) secondary antibodies were used for visualization. Immunoreactivity was assessed under microscopy (Axioplan, Zeiss, Germany) or confocal laser scanning microscopy (TCS NT; Leica Microsystems, Germany). To analyze quantitatively the numbers of TH-positive neurons and AADC-positive neurons, every 10th 30-μm section (total of 11 sections) covering a 3-mm thickness from each animal ($n = 3$ per group) was examined. Coexpression efficacy was analyzed by dual immunofluorescence staining.

Statistical analysis. One-way analysis of variance (ANOVA) was performed to determine differences in dopamine levels, as well as TH activity, followed by Tukey's test (StatView 5.0 software; Abacus). Behavioral changes were analyzed by a repeated measure ANOVA, followed by Tukey's test, with $P < 0.05$ considered statistically significant. Results are presented as means ± SEM.

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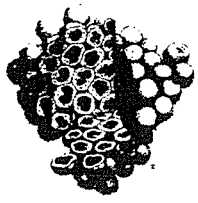

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Improved Safety of Hematopoietic Transplantation with Monkey Embryonic Stem Cells in the Allogeneic Setting

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Improved Safety of Hematopoietic Transplantation with Monkey Embryonic Stem Cells in the Allogeneic Setting

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Key Words. Cynomolgus monkey • Hematopoiesis • Embryonic stem cell • In utero transplantation • Teratoma • Purging Tumor prevention

ABSTRACT

Cynomolgus monkey embryonic stem cell (cyESC)-derived in vivo hematopoiesis was examined in an allogeneic transplantation model. cyESCs were induced to differentiate into the putative hematopoietic precursors in vitro, and the cells were transplanted into the fetal cynomolgus liver at approximately the end of the first trimester (n = 3). Although cyESC-derived hematopoietic colony-forming cells were detected in the newborns (4.1%–4.7%), a teratoma developed in all newborns. The risk of tumor formation was high in this allogeneic transplantation model, given that tumors were hardly observed in immunodeficient mice or fetal sheep that had been xeno-transplanted with the same cyESC derivatives. It turned out that the cyESC-derived donor cells included a residual undifferentiated fraction positive for stage-specific embryonic antigen (SSEA)-4 (38.2% ± 10.3%) despite the rigorous differentiation culture. When an SSEA-4-negative fraction was transplanted (n = 6), the teratoma was no longer observed, whereas the cyESC-derived hematopoietic engraftment was unperturbed (2.3%–5.0%). SSEA-4 is therefore a clinically relevant pluripotency marker of primate embryonic stem cells (ESCs). Purging pluripotent cells with this surface marker would be a promising method of producing clinical progenitor cell preparations using human ESCs. STEM CELLS *2006;24:1450–1457*

INTRODUCTION

Human embryonic stem cells (hESCs) hold great potential in the treatment of a variety of diseases and injuries because embryonic stem cells (ESCs) have the ability to proliferate indefinitely in culture and to differentiate into any cell type [1, 2]. Because ESCs are able to form teratomas when transplanted into immunodeficient mice, safety concerns would be raised against the clinical application of hESCs [3, 4]. It will be necessary to test the safety of these cells in animal transplantation models before clinical application. Nonhuman primate transplantation models would be desirable for this purpose; however, there have been only a few reports on these models [5–7]. The successful engraftment of transplanted cells in primates will not be achieved unless the immune rejection of transplanted cells is circumvented (e.g., through immunosuppressive treatment) [6]. The early gestational fetus may be a good recipient with which to circumvent immune rejection because the immune system is premature [8]. In addition, in the animal fetus, transplanted cells would engraft without conditioning of recipients such as irradiation or immunosuppressive treatment [9–12]. We have previously established a system for allogeneic transplantation of cynomolgus ESCs (cyESCs) using preimmune fetal monkeys as recipients [5].

We have also reported a novel method for hematopoietic engraftment from cyESCs in sheep [13]. The method is a combination of three steps: (a) differentiation in vitro to generate the putative hematopoietic precursors [14]; (b) transplantation of the cells in utero [15]; and (c) development into hematopoietic cells in vivo using the hematopoietic microenvironment of the fetal liver [16]. In the present study,

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we have examined the safety as well as the efficacy of hematopoietic engraftment of cells derived from cyESCs in the allogeneic transplantation model.

MATERIALS AND METHODS

Animals

Pregnant cynomolgus monkeys (16–22 years old) were obtained by mating and were reared at the Tsukuba Primate Research Center in accordance with Rules for Animals Care and Management set forth by the Research Center and Guiding Principles for Animal Experiments Using Nonhuman Primates formulated by the Primate Society of Japan. Experimental procedures were approved by the Animal Welfare and Animal Care Committee of the National Institute of Infectious Diseases. The animals were free of intestinal parasites and were seronegative for herpes virus B, varicella-zoster-like virus, measles virus, and simian immunodeficiency virus.

Cell Preparation

A cyESC line (CMK6G) stably expressing green fluorescent protein (GFP) was established after transfection of the parental cyESC line (CMK6) with the enhanced GFP gene (Clontech, Palo Alto, CA, http://www.clontech.com) [17]. cyESCs were maintained on a feeder layer of mitomycin C (Kyowa, Tokyo, http://www.kyowa.co.jp)-treated mouse (ICR or BALB/c; Clea Japan, Tokyo, http://www.clea-japan.com) embryonic fibroblasts as previously described [18]. The mouse bone marrow stromal cell line OP9 was maintained in α-minimum essential medium (Invitrogen, Carlsbad, CA, http://www.invitrogen.com) supplemented with 20% fetal calf serum (FCS; Invitrogen) [19].

cyESCs were induced to differentiate into the putative hematopoietic precursors as previously described [13]. Briefly, undifferentiated cyESCs were transferred onto mitomycin C-treated confluent OP9 cells and cultured for 6 days in Iscove's modified Dulbecco's medium (Invitrogen) supplemented with 8% FCS, 8% horse serum (Invitrogen), 5×10^{-6} M hydrocortisone (Sigma, St. Louis, http://www.sigmaaldrich.com), and multiple cytokines, including 20 ng/ml recombinant human (rh) bone morphogenetic protein-4 (R&D Systems, Minneapolis, http://www.rndsystems.com), 20 ng/ml rh stem cell factor (Biosource, Camarillo, CA, http://www.biosource.com), 20 ng/ml rh vascular endothelial growth factor (VEGF; R&D Systems), 20 ng/ml rh Flt-3 ligand (PeproTech, Rocky Hill, NJ, http://www.peprotech.com), 20 ng/ml rh interleukin-3 (PeproTech), 10 ng/ml rh interleukin-6 (PeproTech), 20 ng/ml rh granulocyte colony-stimulating factor (PeproTech), and 2 IU/ml rh erythropoietin (Roche, Basel, Switzerland, http://www.roche.com). The cells were resuspended in 0.1% human serum albumin (Sigma)/Hanks' balanced saline solution (Sigma) for transplantation.

Flow Cytometry

Primary antibodies (Abs) used in the present study were anti-human CD34 monoclonal Ab (mAb; BD Pharmingen, San Diego, http://www.bdbiosciences.com/pharmingen), anti-human CD31 mAb (Pharmingen), anti-human CD45 mAb (Pharmingen), anti-human vascular endothelial (VE) cadherin mAb (Pharmingen), rabbit anti-human VEGF receptor (VEGFR)-2 Ab (Santa Cruz Biotechnology, Santa Cruz, CA, http://www.scbt.com), and anti-stage-specific embryonic antigen (SSEA)-4 mAb (Chemicon, Temecula, CA, http://www.chemicon.com). All of them cross-reacted to cynomolgus counterparts as previously demonstrated [18, 20–22]. Secondary Abs were phycoerythrin (PE)-conjugated rabbit anti-mouse immunoglobulins (Ig) Ab (DakoCytomation, Glostrup, Denmark, http://www.dako.com) and Alexa Fluor 647-conjugated goat anti-mouse IgG Ab (Molecular Probes, Eugene, OR, http://probes.invitrogen.com). Cells stained with unlabeled primary Abs were incubated with fluorescence-labeled secondary Abs. Cells were incubated with either primary or secondary Ab for 20–60 minutes at 4°C. Regarding staining with the anti-VEGFR-2 Ab, the cells were incubated with biotin-conjugated goat anti-rabbit IgG Ab (Beckman Coulter, Miami, http://www.beckmancoulter.com), followed by PE-conjugated streptavidin (Beckman Coulter). Fluorescence-labeled cells were analyzed with a FACS Calibur flow cytometer (Becton, Dickinson and Company, Franklin Lakes, NJ, http://www.bd.com). Data analysis was performed using the CellQuest software (Becton, Dickinson and Company). Isotype-matched, irrelevant mAbs (DakoCytomation or Beckman Coulter) served as negative controls. Nonviable cells were excluded from analysis by propidium iodide (Sigma) costaining.

Cell Sorting

Cell sorting was performed to purge SSEA-4^+ cells from among the cultured cyESCs in vitro. Cells were incubated with the anti-SSEA-4 mAb for 1 hour at 4°C and washed twice with Dulbecco's modified Eagle's medium supplemented with 10% FCS. The cells were then incubated with the PE-conjugated anti-mouse Ig Ab for 1 hour at 4°C and washed twice again. GFP-positive and SSEA-4-negative cells were sorted using an Epics Elite cell sorter (Beckman Coulter). Data acquisition was performed using the Expo2 software (Beckman Coulter).

Transplantation and Delivery

Transplant procedures were previously described [5]. Briefly, animals were anesthetized via an intramuscular administration of ketamine hydrochloride (Ketalar, 10 mg/kg; Sankyo, Tokyo, http://www.sankyo.co.jp) and received 0.5%–1.0% isoflurane by inhalation by means of an endotracheal tube. Cells ($0.16–46 \times 10^6$ cells per fetus; Table 1) were injected into the fetal liver through a 23-gauge needle using an ultrasound-guided technique at approximately the end of the first trimester. The fetuses were delivered by cesarean section at 2–3 months after transplant (gestation 120–157 days, full term 165 days).

Colony Polymerase Chain Reaction

Cynomolgus clonogenic hematopoietic colonies were produced as previously described [20]. After cells were cultured in methylcellulose medium for 10–14 days, well-separated individual colonies were plucked into 50 μl of distilled water and digested with 20 μg/ml proteinase K (Takara, Shiga, Japan, http://www.takara-bio.com) at 55°C for 1 hour, followed by 99°C for 10 minutes. Each sample (5 μl) was used for a nested polymerase chain reaction (PCR) to detect the *GFP* gene sequence. The outer primer set was 5′-AAGGACGACGGCAACTACAA-3′ and 5′-ACTGGGTGCTCAGGTAGTGG-3′, and the inner primer set was 5′-GCATCGACTTCAAGGAGGAC-3′ and 5′-GTTGTGGCGGATCTTGAAGT-3′. Amplification conditions for both the outer and inner PCR were 30 cycles of 95°C for 30 seconds, 65°C for 30 seconds, and 72°C for 30 seconds. The

Table 1. ESC-derived hematopoiesis and tumor formation

Animals	Animal no.	Transplanted cells	Purging SSEA-4$^+$ cells	Cell number per fetus ($\times 10^6$)	Donor-derived CFU in recipients[a] at birth (donor/total colony number)	Tumor formation	Observation period (months)
Monkeys	0031	Undifferentiated	−	3.90	n.d.	+	3
	2311	ESCs	−	0.16	n.d., Dead	+	2
	0321		−	0.21	n.d., Dead	+	2
	0841	Day-6 ESC-	−	10	4.1% (2/49)	+	3
	1551	derived cells	−	46	n.d., Dead	+	2.5
	0021		−	46	4.7% (4/85)	+	3
	0691	Day-6 ESC-	+	0.16	3.2% (2/62)	−	3
	0381	derived cells	+	1.40	5.0% (4/80)	−	3
	0022		+	0.17	2.3% (2/86)	−	3
	0981		+	0.31	4.1% (3/73)	−	3
	0051		+	0.31	n.d., Dead[b]	−	3
	1552		+	0.75	4.4% (2/45)	−	4
Sheep[c]	57	Day-6 ESC-	−	50	1.1% (1/91)	−	18
	55	derived cells	−	50	1.1% (1/91)	−	26
	141		−	78	1.1% (1/91)	−	26
	182		−	14	1.6% (1/63)	−	21

[a]Percentage of donor-derived CFU was calculated by dividing the number of CFU positive for the green fluorescent protein gene sequence by the number of CFU positive for the β-actin gene sequence. Donor-derived CFU were analyzed at delivery.
[b]Death due to ablation of placentae. Other deaths were presumably tumor-related.
[c]As published by Sasaki et al. [13].
Abbreviations: CFU, colony-forming units; ESC, embryonic stem cell; n.d., not done; SSEA, stage-specific embryonic antigen.

outer PCR products were purified using a QIA quick PCR purification kit (Qiagen, Valencia, CA, http://www.qiagen.com). Simultaneous PCR for the β-actin sequence was also performed to ensure DNA amplification of the sample in each colony. The primer set for β-actin was 5′-CATTGTCATGGACTCTGGCGACGG-3′ and 5′-CATCTCCTGCTCGAAGTCTAGGGC-3′. Amplification conditions for β-actin PCR were 40 cycles of 95°C for 30 seconds, 65°C for 30 seconds, and 72°C for 30 seconds. Amplified GFP (131 bp) and β-actin (234 bp) products were resolved on 2% agarose gel (Sigma) and visualized by ethidium bromide (Invitrogen) staining.

RNA PCR

Total RNA was extracted from cells of interest using the EZ1 RNA universal tissue kit (Qiagen). RNA was reverse-transcribed at 50°C for 30 minutes using the RNA LA PCR kit (Takara) with oligo dT primer. The resulting cDNA was then subjected to PCR. Regarding PCR for Oct-4, the primer set was 5′-GGACACCTGGCTTCGGATT-3′ and 5′-TTCGCTTTCTCTTTCGGGC-3′. The PCR conditions were 35 cycles of 95°C for 30 seconds, 67°C for 45 seconds, and 68°C for 1.5 minutes. Regarding PCR for Scl, the primer set was 5′-GGGCGGAAAGCTGTTTGCGATT-3′ and 5′-TCGCTGAGAGGCCTGCAGTT-3′. The PCR conditions were 35 cycles of 95°C for 30 seconds, 63°C for 1 minute, and 72°C for 1 minute. A simultaneous PCR for β-actin was also conducted on each cDNA sample as an internal control as described above. Amplified Oct-4 (697 bp), Scl (201 bp), and β-actin (234 bp) products were resolved on 2% agarose gel and visualized by ethidium bromide staining.

RESULTS

In Utero Transplantation and Delivery

cyESCs stably expressing GFP were used in this study [17]. In the setting of allogeneic transplantation, GFP was used as a genetic tag to track transplanted cell progeny. We employed the OP9 stromal cell coculture method instead of the embryoid body formation method to facilitate the hematopoietic differentiation [19, 23, 24] (Fig. 1A, 1B). According to the flow cytometric analysis, CD34, CD31 (platelet/endothelial cell adhesion molecule-1 [PECAM-1]), CD144 (VE-cadherin), and VEGFR-2 (Flk-1) were all upregulated on day 6 but decreased thereafter (Fig. 1C–1E, 1G). Among the markers examined, CD34 is a widely used surface marker of hematopoietic stem cells in both human and monkey subjects [25–27]. The others are key markers of hemangioblasts (which generate endothelial and hematopoietic lineages) in both mice and humans [14, 28]. Cells positive for both VEGFR-2 and VE-cadherin emerged on day 6 (Fig. 1H). CD45, however, was not detected until day 12 (Fig. 1F). Despite the hemangioblast marker expression on day 6, the hematopoietic *Scl* gene was upregulated at this time point as assessed by RNA PCR (Fig. 1I), implying that the hematopoietic commitment might have already occurred on day 6 [29, 30]. We therefore designated the day 6 cyESC-derived progenitor cells as putative hematopoietic precursors. The time course profiles presented here were similar to those of hESCs [14, 24]. The GFP expression was stable during the 6-day culture (Fig. 1A, 1B) and afterward (data not shown).

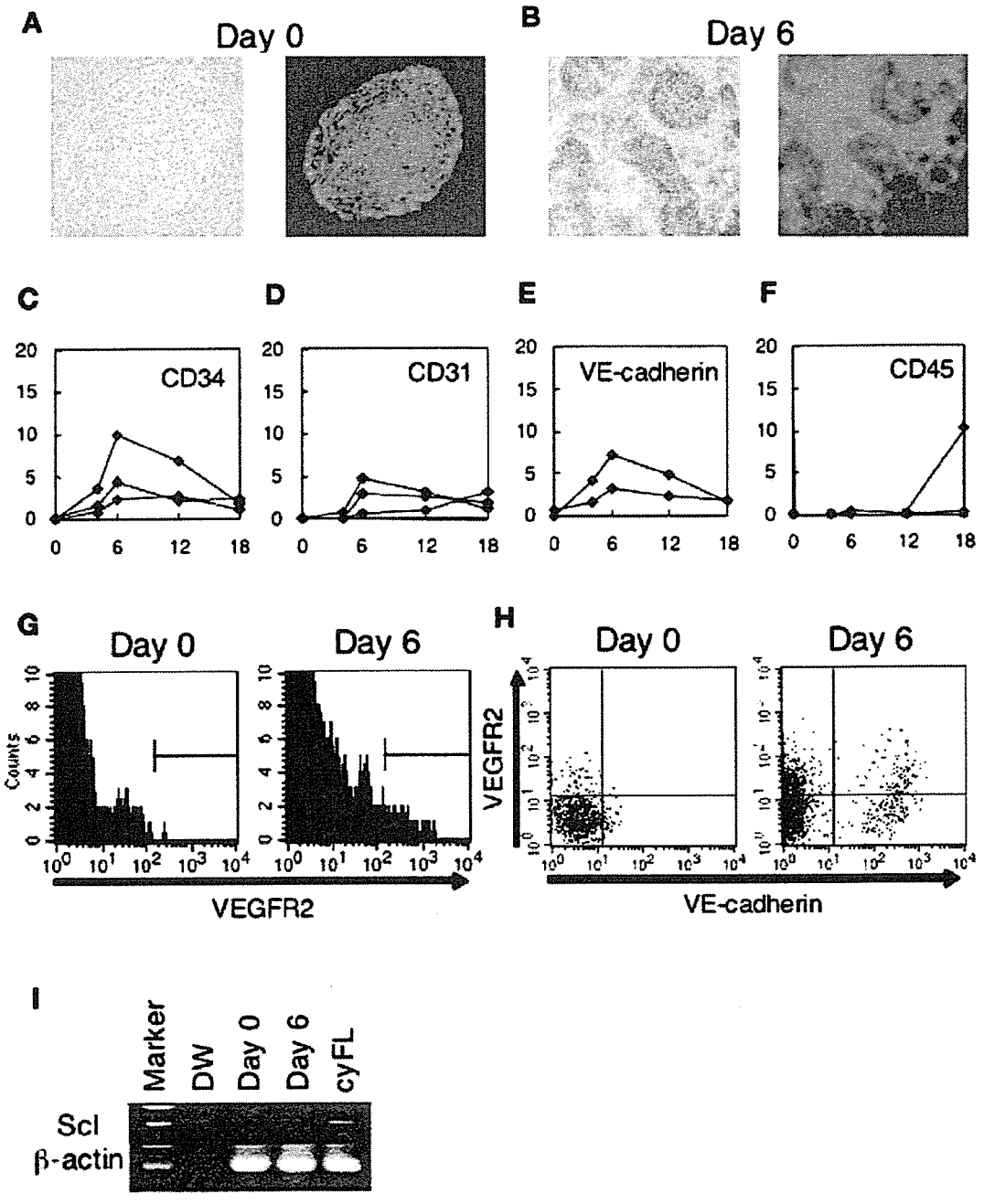

Figure 1. Flow cytometric analysis during the in vitro differentiation of cyESCs. Undifferentiated cyESCs expressing green fluorescent protein were cultured on OP9 cells with multiple cytokines (see Materials and Methods). **(A):** Cells on day 0 are shown in bright (left) and dark (right) fields. **(B):** Cells on day 6 are shown in bright (left) and dark (right) fields. **(C):** Cells on days 0, 4, 6, 12, and 18 were stained for CD34. **(D):** Cells on days 0, 4, 6, 12, and 18 were stained for CD31. **(E):** Cells on days 0, 4, 6, 12, and 18 were stained for VE-cadherin. **(F):** Cells on days 0, 4, 6, 12, and 18 were stained for CD45. The vertical axis shows the fraction (percentage) of cells that were stained positive. **(C–F):** Results of two or three independent experiments are shown. **(G):** Although cells on day 0 already express low levels of VEGFR-2, a VEGFR-2^{high} population did not emerge until day 6. **(H):** Dot-plot profiles for VEGFR-2 and VE-cadherin expression indicate that cells positive for both VEGFR-2 and VE-cadherin emerged until day 6. **(G, H):** Representative results from three independent experiments are shown. **(I):** The *Scl* gene expression was upregulated on day 6 to a level similar to that in the cynomolgus fetal liver as assessed by RNA polymerase chain reaction. Day-6 cells (putative hematopoietic precursors) were used for transplantation. Abbreviations: cyESC, cynomolgus embryonic stem cell; cyFL, cynomolgus fetal liver; DW, distilled water; VE, vascular endothelial; VEGFR, vascular endothelial growth factor receptor.

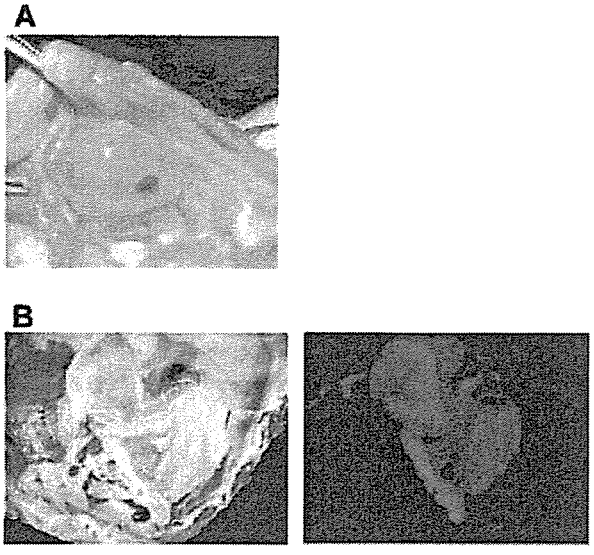

Figure 2. Tumor formation after the transplantation of cynomolgus embryonic stem cell (cyESC)-derived progenitor cells. Tumors formed in all three monkey fetuses transplanted with the day-6 cyESC-derived progenitor cells (putative hematopoietic precursors). **(A):** A representative tumor in the thoracic cavity at 3 months after transplantation (monkey no. 0841). **(B):** The tumor was observed in bright (left) and dark (right) fields under a fluorescence microscope.

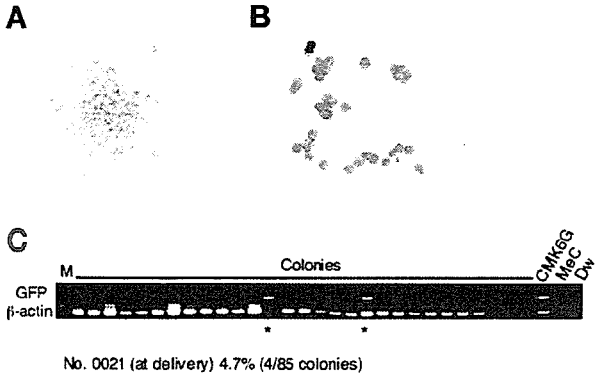

Figure 3. cyESC-derived hematopoiesis in vivo. **(A):** Bone marrow, cord blood, and liver cells were harvested from newborn monkeys and placed in methylcellulose medium to produce clonogenic hematopoietic colonies. **(B):** A cytospin specimen (stained with the May-Giemsa method) of plucked colonies reveals mature neutrophils. To identify cyESC-derived colonies, well-separated individual colonies were plucked and examined for the GFP sequence by PCR. Plucked MeC alone (not containing colonies) served as a negative control. PCR of the β-actin sequence in the same colonies was simultaneously performed as an internal control. Colony PCR was repeated at least twice. **(C):** Representative colony PCR results for monkey no. 0021. Asterisk indicates bands positive for the GFP sequence. Abbreviations: CMK6G, positive control green fluorescent protein-expressing cynomolgus cells; cyESC, cynomolgus embryonic stem cell; DW, distilled water; GFP, green fluorescent protein; M, molecular weight marker; MeC, methylcellulose; PCR, polymerase chain reaction.

Teratoma Formation

The undifferentiated cyESCs ($n = 3$) or cyESC-derived putative hematopoietic precursors ($n = 3$) were transplanted in utero into allogeneic fetuses in the liver under ultrasound guidance at approximately the end of the first trimester (49–66 days, full term 165 days) (Table 1). Regardless of whether the undifferentiated cyESCs or putative hematopoietic precursors were transplanted, tumors were found in the thoracic or abdominal cavities in all the six animals at 2–3 months after transplant (Table 1; Fig. 2A). The tumors fluoresced (Fig. 2B) and consisted of three germ layer cells. Thus, they were teratomas derived from transplanted cells. However, tumors were hardly observed in fetal sheep (1/10; [13] and our unpublished data) (Table 1) and immunodeficient (nonobese diabetic/severe combined immunodeficient) mice (3/10; our unpublished data) after the same putative hematopoietic precursors were transplanted.

In Vivo cyESC-Derived Hematopoiesis

Regarding the newborn monkeys that had been transplanted with the putative hematopoietic precursors, we harvested cells from the femur, cord blood, and liver and plated the cells in methylcellulose medium to produce clonogenic hematopoietic colonies (colony-forming units [CFU]) (Fig. 3A). The monkey cells generated colonies of clear hematopoietic morphology in this assay (Fig. 3B). To detect transplanted cell-derived, GFP-positive colonies, we tried to observe GFP fluorescence of colonies under a fluorescent microscope but were hampered by the high autofluorescence. We then conducted PCR for the *GFP* gene sequence in DNA isolated from each colony (colony PCR) (Fig. 3C). The transplanted cell-derived CFU were clearly detected in the animals (4.1% and 4.7%; Table 1). We repeated the colony PCR and confirmed that the results were reproducible.

We detected both granulocytic and erythroid cynomolgus CFU. In the peripheral blood, however, we were not able to detect cells expressing GFP by flow cytometry. It turned out that, as assessed by quantitative PCR, the fractions of GFP-positive cells in the peripheral blood were very small (<0.1%). Low peripheral "chimerism" has been reported more than once in other in utero transplantations of ESCs or hematopoietic stem cells such as in mice, sheep, and pigs [13, 31–33].

Purging SSEA-4$^+$ Cells of the Putative Hematopoietic Precursors

We examined the expression of an undifferentiated primate ESC marker, SSEA-4, in the undifferentiated cyESCs (day 0) and putative hematopoietic precursors (day 6). The proportion of SSEA-4$^+$ cells was 93.4% ± 8.1% and 38.2% ± 10.3% among the day-0 and -6 cells, respectively (Fig. 4A). A substantial number of cells were still positive for SSEA-4 after the rigorous differentiation culture. In addition, a considerable number of cells expressing another undifferentiated marker, Oct-4, remained among the day-6 population as assessed by RNA-PCR (Fig. 4B). Those residual undifferentiated cells might be responsible for the formation of teratomas in the recipients.

To prevent teratomas from forming in recipients, we purged SSEA-4$^+$ cells of the putative hematopoietic precursors and transplanted the SSEA-4$^-$ population into the fetal monkey liver ($n = 6$) (Fig. 4C). At delivery, tumors were no longer observed in the six animals that had been transplanted with the sorted SSEA-4$^-$ cells (Fig. 4D). The transplanted cell-derived CFU were clearly detected in the newborns, and

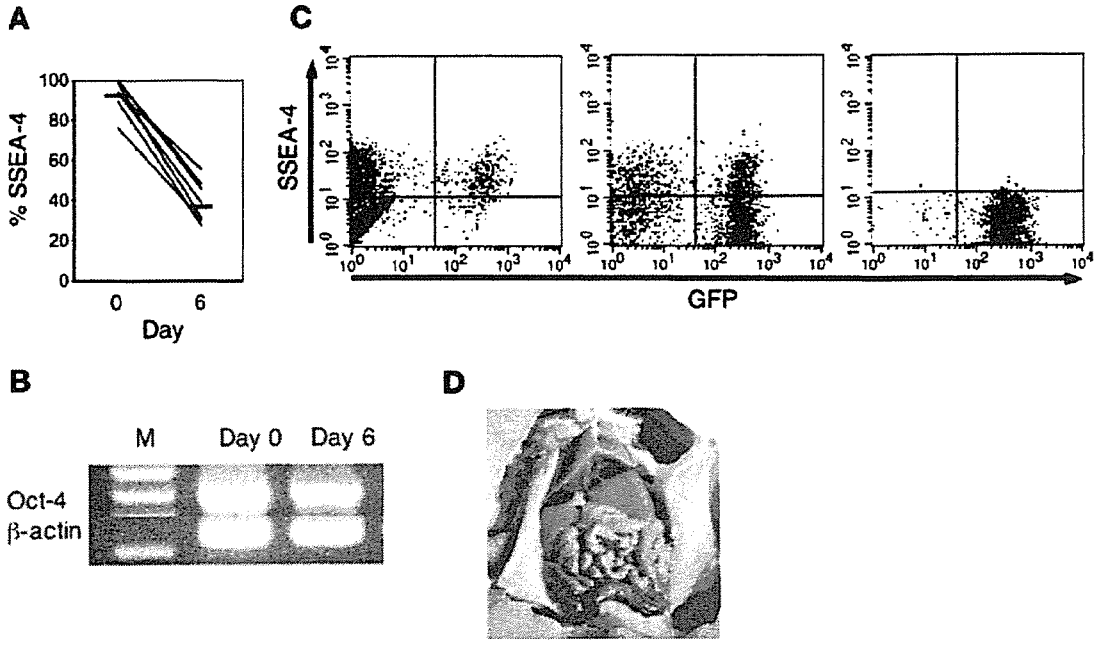

Figure 4. Purging SSEA-4$^+$ cells from among cyESC-derived progenitor cells. **(A):** Undifferentiated cyESCs (day 0) and cyESC-derivatives (day 6) were stained with anti-SSEA-4. The SSEA-4 expression (percentage of total) at day 0 and day 6 is shown (n = 8). **(B):** The Oct-4 expression at days 0 and 6 was also examined by RNA polymerase chain reaction. **(C):** Flow cytometric dot-plot profiles are shown for the SSEA-4 versus GFP expression at day 0 (left), at day 6 before the purge (middle), and at day 6 after the purge (right). Six independent experiments were conducted, and similar results were obtained. **(D):** No tumors were detected in any monkey after the transplantation of SSEA-4-negative day-6 cyESC derivatives (a representative monkey, no. 0981). Abbreviations: cyESC, cynomolgus embryonic stem cell; GFP, green fluorescent protein; M, molecular weight marker; SSEA, stage-specific embryonic antigen.

the fraction was not spoiled (2.3%–5.0%; Table 1), although the removed SSEA-4$^+$ fraction included some CD34$^+$ cells (data not shown).

DISCUSSION

We have previously described a method for hematopoietic engraftment from cyESCs [13]. cyESCs were first cultured for 6 days in vitro, and the day-6 cyESC-derived putative hematopoietic precursors were transplanted in vivo into fetal sheep liver after the first trimester, generating sheep with cynomolgus hematopoiesis. We transplanted the day-6 cells because the CD34 expression level was highest at this time point (Fig. 1C). We transplanted the cells into the liver because the liver is the major hematopoietic organ at this stage of gestation in sheep [34]. In the present study, we tested this method in a cynomolgus monkey allogeneic transplantation model and successfully detected cyESC-derived hematopoietic cells in cynomolgus recipients, albeit at low levels. cyESC-derived chimerism was, however, higher in the primate allogeneic transplantation model (2.3%–5.0%) than in our recently reported sheep xeno-transplantation model (1.1%–1.6%; [13]) (Table 1). To enhance ESC-derived hematopoiesis, further consideration is required of the in vitro culture conditions (i.e., the cytokine milieu, coculture- or embryoid body-associated cellular microenvironment, culture period, and genetic manipulation) and the in utero transplantation conditions (i.e., the preconditioning, route, and timing).

Teratomas developed in all animals, even after the transplantation of ESC-derived progenitor cells that had been cultured for 6 days in the differentiation medium. The risk of tumor formation was high, given that we could hardly detect tumors in immunodeficient mice or fetal sheep that had been transplanted with the same day-6 cyESC derivatives ([13] and our unpublished data). Innate immune responses against cynomolgus-derived tumors might be more rigorous in xeno-transplanted mice and sheep than in allo-transplanted monkeys, resulting in a failure to detect tumorigenesis in the xeno-transplantation models. Similarly, Erdo et al. reported that tumors developed after ESC-derived progenitor cell transplantation in the mouse-to-mouse setting, but not in the mouse-to-rat setting [35]. Our monkey allogeneic transplantation setting would therefore allow the strict evaluation of the in vivo safety of transplantation therapies using ESCs. However, given that teratomas indeed form when undifferentiated cyESCs alone are xeno-transplanted into immunodeficient mice, it is unclear why residual undifferentiated cells included among the day-6 cyESC derivatives did not form teratomas in immunodeficient mice or fetal sheep.

SSEAs that are developmentally regulated during early embryogenesis are widely used as markers to monitor the differentiation of both mouse and human embryos and ESCs [36–38]. Undifferentiated ESCs of both human and cynomolgus origin are characterized by the expression of SSEA-4 and by a lack of SSEA-1 [1, 2, 18]. We have therefore used SSEA-4 as a marker for the negative selection of an undifferentiated fraction. As a result of this negative selection, tumors were no longer detected in the monkeys after transplantation. On the other hand, Bieberich et al. recently developed a method for selective apoptosis of residual pluripotent stem cells using the transcription

factor Oct-4 as a pluripotency marker to prevent teratoma formation [39]. They found that the expression of Oct-4 is colocalized with that of prostate apoptosis response-4, a protein mediating ceramide-induced apoptosis. Treatment of ESC-derived neural precursors with ceramide resulted in selective elimination of residual Oct-4-positive pluripotent cells. Our method, however, uses a cell surface marker to purge pluripotent cells. With this method, one can see the purging efficiency in real-time. This would be meritorious for clinical applications. Although we used a cell sorter to obtain the SSEA-4$^-$ fraction in the present study, selection with beads would be easier and more appropriate for clinical applications.

To generalize the use of SSEA-4 for eliminating undifferentiated cells from among donor cells, we differentiated cyESCs into neural stem cells. After the culture, approximately 10% of cells were still positive for SSEA-4. When all the cells were transplanted into the striatum of Parkinson's cynomolgus monkeys, teratomas developed. We then transplanted cyESC-derived neural stem cells without an SSEA-4$^+$ fraction into the cynomolgus striatum and successfully detected the engraftment without tumor formation (our unpublished data). The removal of SSEA-4$^+$ cells is useful at least for hematopoietic and neural lineages.

CONCLUSION

We are now able to prevent the formation of tumors in nonhuman primate recipients by purging SSEA-4$^+$ cells from among ESC-derived progenitor cells without spoiling the engraftment. SSEA-4 is therefore a clinically relevant pluripotency marker of primate ESCs. Purging pluripotent cells with this marker would be a promising method for producing clinical progenitor cell preparations using hESCs to improve safety in vivo.

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DISCLOSURES

The authors indicate no potential conflicts of interest.

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