The Fetal Hypothalamus Has the Potential to Generate Cells with a Gonadotropin Releasing Hormone (GnRH) Phenotype

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Abstract

**Background:** Neurospheres (NS) are colonies of neural stem and precursor cells capable of differentiating into the central nervous system (CNS) cell lineages upon appropriate culture conditions: neurons, and glial cells. NS were originally derived from the embryonic and adult mouse striatum subventricular zone. More recently, experimental evidence substantiated the isolation of NS from almost any region of the CNS, including the hypothalamus.

**Methodology/Findings:** Here we report a protocol that enables to generate large quantities of NS from both fetal and adult rat hypothalami. We found that either FGF-2 or EGF were capable of inducing NS formation from fetal hypothalamic cultures, but that only FGF-2 is effective in the adult cultures. The hypothalamic-derived NS are capable of differentiating into neurons and glial cells and most notably, as demonstrated by immunocytochemical detection with a specific anti-GnRH antibody, the fetal cultures contain cells that exhibit a GnRH phenotype upon differentiation.

**Conclusions/Significance:** This in vitro model should be useful to study the molecular mechanisms involved in GnRH neuronal differentiation.

Introduction

NS were first discovered by Reynolds and Weiss [1,2], as colonies of cells containing neural stem and precursor cells in embryonic and adult mouse striatum subventricular zone. These primary NS could be expanded in vitro by mechanical dissociation to single cells that generated new (secondary) NS, in a process that can be repeated many times. Furthermore, cells in each NS could be induced to differentiate into neurons and glial cells, thus demonstrating the two cardinal features of stem cells: self-renewal and multipotentiality. This initial report fueled many other investigations that further confirmed and substantiated the idea that neurospheres can be isolated from many other regions of the CNS of rodents and humans [3–7]. Overall, these results suggest that the adult CNS has strong neurogenic potential, presumably due to the presence of putative stem and progenitor cells. The precise nature of these cells is still a matter of investigation and controversy [8–10]. It is assumed that under normal conditions, these cells are kept in a quiescent state by inhibitory signals, but can be induced to proliferate upon exposure to adequate growth factors, most notably EGF and FGF-2 [11–13]. The recent finding by Markakis et al. [14] that GnRH-immunoreactive cells can be derived from in vitro expanded adult progenitor cells prompted us to set up a NS assay using either fetal or adult rat hypothalamic tissue and verify if we could detect cells with GnRH-phenotype in these cultures. The advantage of the NS assay is that stem/progenitor cells can be 1) isolated and propagated in a serum free medium, 2) studied at clonal density, and 3) stimulated to induce the differentiation of the whole population of the growing cells. This method should also help to determine whether fetal CNS progenitors have the potential to generate GnRH neurons. Here we report that both the fetal and the adult rat hypothalami are a rich source of NS that can be expanded and passaged for a long time, with the capability to give rise to neurons and glial cells under differentiating conditions. In addition, we report the detection of GnRH-immunoreactive cells among differentiating NS derived from fetal cultures.

Results

Hypothalamic tissue is a rich source of neurospheres with self-renewing capacity

Hypothalami were recovered from E18 embryos (Wistar rats) and after tissue dissociation, cells were seeded into 6-well plates at
hypothalamic NS after 7DIV. C, Adult hypothalamic NS after 7DIV. Scale bars: 50 μm.

Figure 1. Morphology of hypothalamic-derived NS, phase-contrast images. A, Dividing cells after 2 days in vitro (2DIV). B, Fetal hypothalamic NS after 7DIV. C, Adult hypothalamic NS after 7DIV. Scale bars: 50 μm.

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Hypothalamic NS express Neuronal, Glial and Neuroepithelial markers, and can differentiate into Neuron, Astrocyte and Oligodendrocyte cells

To characterize the hypothalamic NS obtained and establish whether cells forming the NS exhibit multipotentiality, the expression of a panel of cell-type specific marker genes such as...
NSE (neuron specific enolase neuronal marker), GFAP (astrocyte marker) and Nestin (neuroepithelial/undifferentiated cell marker) was studied by RT-PCR. The NS generated within each separate culture were pooled and processed in order to prepare RNA which was then used for RT-PCR analysis. For these experiments, we only used NS derived from passaged primary fetal and adult NS (secondary and tertiary NS), to avoid any possibility of contamination with cells derived from primary culture. Fig. 4A shows the results of a representative experiment from adult NS (n=4 samples from independent cultures analyzed), confirming the expression of all the cell markers assessed. These are compared to the expression of the same markers in RNA extracted from adult rat hypothalami (HT). Beta-2 microglobulin gene (β-2m) was chosen as housekeeping gene to verify sample quality and PCR conditions. NS isolated from other parts of the CNS could differentiate into neurons and glial cells when cultured under appropriate differentiating conditions, usually in media supplemented with 1–2% FBS but without the presence of the mitogens EGF/FGF-2. Therefore, we next verified whether the hypothalamic-derived NS could differentiate into both neurons and glial...
cells. Either secondary or tertiary NS were mechanically dissociated, and the resulting single-cells were cultured in a media devoid of growth factors but supplemented with 1% FBS. Cells kept under this condition attached to the surface of vessels and started to assume differentiated morphologies [see Fig. 4B]. A representative picture from immunocytochemical analysis of fetal cultures at 7DIV is displayed in Figure 4C and demonstrates that the major CNS phenotypes can be detected, using specific antibodies for neurons (beta Tubulin III, red staining), and astrocytes (GFAP, green staining). Cells immunoreactive for the neuronal marker beta Tubulin III represented 5 ± 1.5% of the total population derived from fetal NS (n = 3 cultures analyzed, from independent differentiating secondary and tertiary NS cultures). Parallel analysis for differentiated NS derived from adult NS gave similar results (data not shown). In addition, we also were able to detect, in fetal differentiating cultures, cells expressing the oligodendrocyte marker O4 (see Fig. 4D).

Hypothalamic-specific transcription factors are expressed in NS derived from adult and fetal rat hypothalami

The transcription factors Otp, Brn2, Arnt2 and Ttf1 have been shown to play a pivotal role in the development of different neuroendocrine cell lineages within the hypothalamus [21–23]. To assess whether hypothalamic-derived NS retained characteristics related to the region from which they were isolated, the expression of these transcription factors was assessed by RT-PCR in adult and fetal secondary/tertiary NS. As shown in Fig. 5, these factors were all expressed in both adult and fetal hypothalamic RNA extracts, as expected, as well as in RNA extracts from the adult NS cultures. In contrast, in RNA extracts from fetal NS only Ttf1 and Arnt2 were detected (right side of Fig. 5). Of note, none of these factors was expressed in the adult anterior pituitary gland, used as a control, which instead expressed the pituitary-specific transcription factor Pit1 [24]. Pit1 was neither detected in the adult hypothalamus nor in the adult/fetal NS. Overall, these results indicate that our hypothalamic-derived NS cultures retain some specificities typical of the region from which they are derived.

GnRH-Immunoreactive cells are present among differentiating NS of Fetal origin

We next wondered if cells with a GnRH phenotype could be present in these hypothalamic-derived NS cultures after differentiation. To detect GnRH expressing cells, we performed immunocytochemistry on cells fixed in 4% paraformaldehyde using either a polyclonal antibody directed against GnRH [25] or a commercially available anti-GnRH antibody. Differentiated cultures were first studied after 7DIV in standard NS medium containing 1% FBS but without growth factors, and on poly-D-Ornithine treated coverslips. GnRH immunoreactive cells were indeed detected at the end of this incubation and represented an average of 10 ± 0.5% of all cells (n = 3 cultures analyzed, from independent differentiating tertiary NS cultures). We next tried several other incubation media and finally were able to enrich the GnRH-immunoreactive cells population by using a differentiating...
protocol which consisted in incubating dissociated tertiary fetal NS for 5 days in a DMEM/F12 plus FGF-2 (20ng/mL) media followed by a further 5 days incubation in a DMEM/F12 plus B27 (0.5%) media(cells grown on poly-D-Ornithine treated coverslips). Under these conditions, the GnRH-immunoreactive cells (Fig. 6, green fluorescent cells) represented 21 ± 2% of all cells analyzed (n = 3 cultures analyzed, from independent differentiating tertiary NS cultures), thus basically doubling the yield obtained from the previous protocol. In contrast, no GnRH-immunoreactive cells could be detected in adult derived NS (data not shown).

Discussion

We report here that primary cultures obtained from either fetal (E18) or adult rat hypothalamic tissue are an abundant source of neuronal precursor cells that proliferate in culture as free-floating spheres. By supplementing standard hypothalamic primary cultures [26] with growth factors such as EGF and FGF-2, we were able to reproducibly obtain large amounts of NS that can be maintained for a long period of time while retaining the capability of differentiating into the principal CNS cell phenotypes: neurons and glial cells. These data are in agreement with a recent report demonstrating the generation of NS from explants derived from small, anatomically-defined, subregions of adult rat CNS, including parts of the hypothalamus [6]. Interestingly, these authors also found that hypothalamic explants were among the richest sources of NS compared with other regions of the CNS such as the hippocampus.

We found here that the different growth factors used in the cultures influenced differently the development of NS in fetal compared to adult hypothalami. In fetal cultures, EGF and FGF-2 were almost equally efficient in promoting the generation of primary NS, but without exhibiting any additive effect. This observation implies that the same population of progenitor/stem cells is responding to stimulation with either growth factor, suggesting the expression of their two specific receptors in the same cells. In cultures from adult hypothalami, generation of NS was obtained effectively and reliably only with FGF-2, despite our
concomitant finding that the EGFR is expressed in adult hypothalami. This observation is nevertheless consistent with previous data [20], but its significance remains to be elucidated.

We were also able to demonstrate that these NS are derived from the hypothalamic parenchyma, at least in the adult rat cultures, since the entire third ventricle lining including its immediately surrounding tissue was removed before culturing. This point appears important because the ependymal and subependymal layers of the lateral ventricles were once believed to be the only regions where putative neural progenitors/neuronal stem cells reside [27]. Consistently, more recent data show that the ependymal layer of the third ventricle also contains neuronal precursor cells capable of proliferation and migration within the hypothalamic parenchyma [20]. However, it is now relatively well established that parenchymal tissue from different CNS regions such as the cortex, septum, striatum and the optic nerve contain quiescent neuronal progenitors that can be induced to proliferate in vitro [4,7,11]. In our system, the hypothalamic-derived NS cultures were able to retain some phenotypic specificity of their area of origin. We could notably demonstrate the expression of transcription factors essential for the differentiation of many neuroendocrine cell lineages of the hypothalamus [21–23] in NS cultures from adult and fetal hypothalami. This result is entirely in line with the demonstration that neural stem cells and their progeny are regionally specified in the developing brain [28].

The postnatal hypothalamus has long been considered a neurogenically silent region with no proliferative capacity, but recent in vivo studies are challenging this hypothesis: in rats, the icv infusion of BDNF results in the accumulation of BrdU-positive cells in several CNS regions, including the hypothalamus [19]. These proliferating cells are scattered in a large hypothalamic parenchymal area around the third ventricle, and many of them express neuronal markers. Similarly, the icv infusion of CNTF in adult mice is exerting a strong anorexigenic effect leading to long-lasting weight loss that is dependent upon the de novo generation of neuronal cells [18]. In fact, CNTF-treated adult mice showed accumulation of BrdU-positive cells in the hypothalamus, especially in the ventromedial and arcuate nuclei, two areas involved in the control of feeding. The present data are therefore consistent with these studies [18,19] and support the hypothesis that putative neural precursor cells/stem cells residing in the adult hypothalamic parenchyma are normally in a quiescent state, but capable of responding to stimulation by opportune exogenous factors such as BDNF and CNTF.

We therefore reasoned that CNTF or BDNF should have a similar proliferative effect in vitro. However, despite many attempts and the fact that the receptors for these molecules are expressed in our system, we never succeeded in generating primary NS when the two growth factors EGF or FGF-2 were substituted by either CNTF, BDNF or a combination of the two. It therefore appears that the putative CNTF/BDNF-responsive cells residing within the hypothalamic parenchyma are either not present or not capable of responding to stimulation in the primary neural cell cultures. One possible explanation would be that even though CNTF/BDNF is able to induce neurogenesis in vivo, it may not be sufficient to support all the physiological processes required for the in vitro generation of NS and that other factors may interact with CNTF/BDNF to promote proliferation. Another possibility could be that EGF and FGF-2 responsive cells are already committed for proliferation in vivo [29,30], and that they maintain this characteristic in vitro, resulting in a prompt response to stimulation. Other cell types may be less robust, or loose their commitment toward proliferation once away from their natural microenvironment.

We were also able to demonstrate the presence of GnRH-expressing cells among NS derived from fetal (E18) cultures when they were kept in differentiating conditions. This is somewhat surprising considering the embryonic origin of GnRH neurons, which are thought to derive from migratory cells originating from the olfactory placode [31]. However, more recent data suggest that GnRH-expressing cells may have multiple embryonic origins, a hypothesis that would be consistent with the present results [32]. Moreover, data derived from transgenic mice expressing the LacZ reporter gene under the control of the GnRH promoter indicate that GnRH-expressing cells can be transiently detected in the forebrain during embryogenesis [33]. Finally, GnRH-immunoreactive cells derived from in vitro expanded progenitor cells were also reported from either adult hypothalamic or hippocampal cells [14]. For the latter studies, the authors used a density gradient centrifugation technique to isolate progenitor cells from either the

| Gene   | Adult HT | Adult NS | Adult Pit | Adult no-RT | Fetal HT | Fetal NS | Fetal Pit |
|--------|----------|----------|-----------|-------------|----------|----------|----------|
| Otp    |          |          |           |             | 35 cycles| 35 cycles| 35 cycles|
| Brn2   |          |          |           |             | 35 cycles| 35 cycles| 35 cycles|
| Ttf1   |          |          |           |             | 40 cycles| 40 cycles| 40 cycles|
| Arnt2  |          |          |           |             | 40 cycles| 40 cycles| 40 cycles|
| pit1   |          |          |           |             | 26 cycles| 26 cycles| 26 cycles|
| β-2m   |          |          |           |             |          |          |          |

Figure 5. Hypothalamic-derived adult and fetal NS express some region-specific transcription factors. Gene expression analysis was determined by RT-PCR. RNA was isolated from adult and fetal tertiary NS cultures, adult and fetal hypothalami (HT) and adult anterior pituitary glands (Pit). Primers were designed to detect Otp (454 bp), Brn2 (390 bp), Ttf1 (128 bp), Arnt2 (162 bp), Pit1 (505 bp), and the housekeeping gene β-2m (142bp) used as control. The number of cycles performed for the PCR reactions were indicated for each gene on the right side. doi:10.1371/journal.pone.0004392.g005
hypothalamus or the hippocampus, and they were able to detect many GnRH-immunoreactive cells in the cultures originated from both CNS regions. They suggest that when put in vitro, progenitor cells are freed of the constraints of their embryonic origin and can therefore generate different subtypes of neurons than they would do in vivo. This hypothesis could possibly explain the present results as well, but our study differs from the previously published experiments [14] in the following ways: the techniques used for the isolation of precursor cells were different, and so were the conditions of differentiation applied. Another important difference between our study and that of Markakis et al. [14] is that we could detect GnRH-immunoreactive cells only in fetal but not in adult hypothalamic cultures. Markakis et al. used 7 week-old Fisher rats to derive progenitor cells and we used Wistar rats, between 8 to 10 weeks of age, for the adult hypothalamic cultures. Therefore, we cannot rule out the possibility that either a difference in age or/and in the genetic background could explain these discrepancies. Alternatively, GnRH progenitors may reside in the adult SVZ, which was not isolated in our culture system. Nevertheless, our data and these previous studies [33,14] together suggest that GnRH-expressing neurons could arise from central nervous system progenitor cells, but further work will be necessary to assess the functional significance of this observation.

In conclusion, we report here that using a simple and robust culture protocol, both the fetal and adult hypothalami are a rich source of NS that can give rise to cells of all three lineages of the central nervous system. This capacity of the adult hypothalamic parenchyma to generate NS does not depend upon the presence of the ependymal layer of the third ventricle, and the neuronal cells thus obtained still express specific hypothalamic molecular markers. Finally, we were able to detect GnRH-expressing neurons among cells derived from these fetal NS, and we therefore suggest that this culture system represents a useful model to study the molecular mechanisms of GnRH neuronal cell differentiation, a central process for the neuroendocrine control of reproduction in mammals.

Figure 6. GnRH-immunoreactive cells as detected by a specific anti-GnRH antibody in differentiating fetal hypothalamic NS cultures. Representative fluorescence micrographs showing some GnRH-labeled immunofluorescent cells (green fluorescence) as detected in differentiating cultures kept for 5 days in DMEM/F12, FGF-2 (20ng/mL) followed by 5 days in DMEM/F12, B27 (0.5%). Duplicate pictures show the nuclei stained in blue by the dye DAPI. Scale bars: 20 μm (A panel) and 10 μm (B panel).
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**Materials and Methods**

**Hypothalamic dissection and NS cell cultures**

Adult (8 to 10 weeks old) and fetal (E18) Wistar rats were used for this study (Charles River, Les Oncins, France). Animals were killed by decapitation and hypothalami were dissected out of brains and immediately placed into sterile trituration solution (0.1% Glucose, 1% Penicillin-Streptomycin in PBS). Under a tissue culture hood, the hypothalamic tissue was washed 3 times in sterile trituration solution to eliminate residual blood. Mechanical dissociation of hypothalamic tissue was done by gentle pipetting up and down with a sterile glass Pasteur pipette. This procedure was repeated 3 times, each time recovering the dissociated tissue away from tissue debris. Pooled dissociated tissue fractions were then centrifuged (5 minutes at 700–800 rpm) and resulting cell pellets from tissue debris. Pooled dissociated tissue fractions were then centrifuged (5 minutes at 700–800 rpm) and resulting cell pellets were suspended in 2–4 mL (volume depending on the amount of starting tissue) of culture media (DMEM/F12 supplemented with 0.5% B-27) supplemented with 1% BSA and 10% DMSO. Trypan Blue staining was used to count viable cells and proliferation NS were kept in liquid nitrogen, in their standard media (DMEM:F12, 0.5% B-27) supplemented with 1% BSA and 10% DMSO.

**RT-PCR**

Total RNA was extracted from either pooled neurospheres or differentiating cultures by using the TRIzol reagent (Roche). Extracted RNA was quantitated by spectrophotometry and quality evaluated by running samples on denaturing agarose gels. cDNA was prepared from 100–250 ng of total RNA using SuperScriptII (Invitrogen) enzyme and random primers, according to manufacturer’s instructions. The primer pairs used for the PCR reactions are shown on Table 1.

**Removal of third ventricle and surrounding tissue**

In order to remove third ventricle tissue, adult rats were anesthetized with chloral hydrate (400 mg/Kg bw i.p.) and stereotactically injected with either a solution of physiological saline or of an intracerebroventricular (icv) solution of 20 mM (0.1% Glucose, 1% Penicillin-Streptomycin in PBS). Under a tissue culture hood, the hypothalamic tissue was washed 3 times in sterile trituration solution to eliminate residual blood. Mechanical dissociation of hypothalamic tissue was done by gentle pipetting up and down with a sterile glass Pasteur pipette. This procedure was repeated 3 times, each time recovering the dissociated tissue away from tissue debris. Pooled dissociated tissue fractions were then centrifuged (5 minutes at 700–800 rpm) and resulting cell pellets were suspended in 2–4 mL (volume depending on the amount of starting tissue) of culture media (DMEM/F12 supplemented with 0.5% B-27 and 20 ng/mL of either FGF-2 or EGF growth factors). Fresh media to compensate for evaporation, cells were seeded into either 6-well or 12-well plates at a density of 10,000 cells/mL into new vessel plates. For long term storage cells with the help of yellow pipette tips and reseeding at a density of 10–20,000 cells/mL. Fresh media to compensate for evaporation, cells were seeded into either 6-well or 12-well plates. For long-term culture cells were kept in liquid nitrogen, in their standard media (DMEM:F12, 0.5% B-27) supplemented with 1% BSA and 10% DMSO.

**Immunocytochemistry**

To induce differentiation, proliferating NS were mechanically dissociated into single cells and seeded on 12-well plates containing glass coverslips treated with poly-L-Ornithine and cultured in DMEM:F12, 0.5% B27, 1% FBS, without growth factors. After 7 days in culture, cells were fixed in 4% Paraformaldehyde with 20 min incubation and then kept refrigerated in PBS till use. All primary antibodies used in the study were incubated overnight at 4°C. All secondary antibodies used in this study were incubated for 1 h at room temperature. Counterstaining was performed by incubating coverslips with DAPI solution (Molecular Probes, Eugene, OR) for 5 minutes. All coverslips were mounted on glass slides with the Mowiol solution (Calbiochem, Darmstadt, Germany).

Primary antibodies used in this study were the following: mouse IgG monoclonal anti-ß-tubuline III (Sigma, Buchs, Switzerland) used at 1:1000 dilution, rabbit anti-GFAP antisera (Dako, HighWycombe, UK) used at 1:2000 dilution, rabbit anti-GnRH antibody (Chemicon, Basel, Switzerland) used at 1:5000 dilution, rabbit polyclonal anti-GnRH antisera [25] used at a 1:200 dilution and mouse IgM monoclonal anti-O4 (Chemicon, Basel, Switzerland) used at 1:1000 dilution, rabbit anti-GFAP antiserum (Dako, HighWycombe, UK) used at 1:2000 dilution, rabbit anti-GnRH antibody (Chemicon, Basel, Switzerland) used at 1:5000 dilution, rabbit polyclonal anti-GnRH antisera [25] used at a 1:200 dilution and mouse IgM monoclonal anti-O4 (Chemicon, Basel, Switzerland) used at 1:5000 dilution. Secondary antibodies (Jackson Immuno-Research, West grove, PA) used in this study were the following: FITC-conjugated goat antibody to rabbit IgG used at

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### Table 1. Primers used for the PCR analysis.

| Gene   | sense                                                     | antisense                                                   |
|--------|-----------------------------------------------------------|-------------------------------------------------------------|
| NSE    | 5’-AACCTGATGCTGGAGTTGATG-3’                               | 5’-CCTTGCAATGCTGTCCTCAC-3’                                  |
| GFAP   | 5’-AGCTCAATAGCGCTGCTGAG-3’                               | 5’-GACTCAACCTCTCCTCGAGTCA-3’                               |
| Nestin | 5’-AGTGTCGCTAGAGTGGAACAGACG-3’                           | 5’-CTCAAGAGGCTGTCACAGAGCTG-3’                               |
| Rtn2   | 5’-GGCCCTAAGCCGTCACC-3’                                  | 5’-GAAGATGGTGTCCTCAC-3’                                    |
| EphB   | 5’-CATCGGGAACCTCAGAA-3’                                  | 5’-TCCAATGCTGCTTCCA-3’                                     |
| FgfR1  | 5’-CATCCTCTGCATGTTGACC-3’                                | 5’-TCAGGGTCCTCACC-3’                                       |
| Cntfr  | 5’-CGATCCCTCAGAAGATGG-3’                                 | 5’-TCAGGGATGTTGGAGCTG-3’                                   |
| NK1    | 5’-CCTCTGAACTGAGCACCAG-3’                                | 5’-ATGGCTGGGCTGCTT-3’                                      |
| Gp130  | 5’-TCAACTTTGGGAACTCATTGG-3’                              | 5’-TCCAAGTACAGACTTGC-3’                                    |
| LIFR   | 5’-GCCTGGGATGCTAAAGGATCC-3’                              | 5’-AACAAATGTCCTTTCTGAGGCG-3’                               |
| Otp    | 5’-ATGGCTGCTCAGCCGCCAC-3’                                | 5’-ACCAAGATCGACTCCGCAGGAT-3’                               |
| Amt2   | 5’-CTGTTGCTGCTACTGGTG-3’                                 | 5’-CCTAGCTGGAATCAGGAA-3’                                   |
| Bnn2   | 5’-CTAGGCCACGCCGCACT-3’                                  | 5’-TGGATGATGGGCTGACTG-3’                                   |
| Ttf1   | 5’-AAATTTGGGGCCTTTCTTG-3’                                | 5’-AGAGTATCCACAGGAAGG-3’                                   |
| Pit1   | 5’-CGTGGAATGCGCCCTTTGATAC-3’                             | 5’-ACCTTCCGGCTGGACTC-3’                                    |

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Author Contributions

Conceived and designed the experiments: RS. Performed the experiments: RS mG JPR MJV. Analyzed the data: RS YA RCG PYR FPP. Contributed reagents/materials/analysis tools: PYR. Wrote the paper: RS YA FPP.