

# Gangliosides of myelosupportive stroma cells are transferred to myeloid progenitors and are required for their survival and proliferation

Ana L. ZIULKOSKI\*<sup>†</sup>, Cláudia M. B. ANDRADE\*, Pilar M. CRESPO<sup>‡</sup>, Elisa SISTI\*<sup>†</sup>, Vera M. T. TRINDADE\*, Jose L. DANIOTTI<sup>‡</sup>, Fátima C. R. GUMA\* and Radovan BOROJEVIĆ<sup>§</sup>

\*Laboratório de Bioquímica e Biologia Celular de Lipídios, Departamento de Bioquímica, Instituto de Ciências Básicas da Saúde, Universidade Federal do Rio Grande do Sul, Porto Alegre, Brazil, <sup>†</sup>Instituto de Ciências da Saúde, Centro Universitário FEEVALE, Novo Hamburgo, Rio Grande do Sul, Brazil, <sup>‡</sup>Centro de Investigaciones in Química Biológica de Córdoba, CIQUIBIC (UNC-CONICET), Departamento de Química Biológica, Facultad de Ciencias Químicas, Universidad Nacional de Córdoba, Córdoba, Argentina, and <sup>§</sup>Instituto de Ciências Biomédicas & Programa Avançado de Biologia Celular Aplicada à Medicina, Hospital Universitário Clementino Fraga Filho, Universidade Federal do Rio de Janeiro, Rio de Janeiro, Brazil

In previous studies, we have shown that the myelopoiesis dependent upon myelosupportive stroma required production of growth factors and heparan-sulphate proteoglycans, as well as generation of a negatively charged sialidase-sensitive intercellular environment between the stroma and the myeloid progenitors. In the present study, we have investigated the production, distribution and role of gangliosides in an experimental model of *in vitro* myelopoiesis dependent upon AFT-024 murine liver-derived stroma. We used the FDC-P1 cell line, which is dependent upon GM-CSF (granulocyte/macrophage colony-stimulating factor) for both survival and proliferation, as a reporter system to monitor bioavailability and local activity of GM-CSF.  $G_{M3}$  was the major ganglioside produced by stroma, but not by myeloid cells, and it was required for optimal stroma myelosupportive function. It was released into the supernatant and selectively incorporated into the myeloid progenitor cells, where it segregated into rafts

in which it co-localized with the GM-CSF-receptor  $\alpha$  chain. This ganglioside was also metabolized further by myeloid cells into gangliosides of the *a* and *b* series, similar to endogenous  $G_{M3}$ . In these cells,  $G_{M1}$  was the major ganglioside and it was segregated at the interface by stroma and myeloid cells, partially co-localizing with the GM-CSF-receptor  $\alpha$  chain. We conclude that myelosupportive stroma cells produce and secrete the required growth factors, the cofactors such as heparan sulphate proteoglycans, and also supply gangliosides that are transferred from stroma to target cells, generating on the latter ones specific membrane domains with molecular complexes that include growth factor receptors.

**Key words:** ganglioside, granulocyte/macrophage colony-stimulating factor (GM-CSF), intercellular environment, membrane raft, myelopoiesis.

## INTRODUCTION

Cells of the myeloid lineage can either dwell inside solid tissues in close contact with adjacent cells and the intercellular matrix or circulate free in liquid tissues, such as blood and lymph. The former situation is required for proliferation and commitment of myeloid cell progenitors, which occur mainly in the bone marrow. Myeloid cell populations can be amplified further in peripheral tissues. Circulation of leucocytes and myeloid cells in blood is only transitory, as the final site of their migration and activity occurs inside normal or injured tissues [1–3]. In both bone marrow and peripheral tissues, myeloid cell proliferation and activation are dependent upon the local stroma cells, the extracellular matrix and the cytokines or inflammatory mediators that are locally produced or brought into the tissues by blood [4–6].

Intercellular mediators and cytokines usually attain only relatively low concentrations in circulating biological fluids and their activity is dependent upon binding to the available membrane receptors on the target cells. Inside solid tissues, the intercellular space is commonly narrow and the mediators can attain high concentrations and steep gradients, interacting with components of the extracellular matrix or the cell glycocalyx. Their bioavailability and activity are thus largely dependent upon macromolecular complexes that surround the membrane receptors [3,7,8]. Moreover, while settling or moving inside the tissues, myeloid cells

and their progenitors become polarized and redistribute the specialized membrane regions, which can fully segregate the cytokine receptors and the associated membrane components by capping. The cell can thus be highly responsive to a given cytokine on one of its sides and insensitive on the other. Under such conditions, the interaction of myeloid cells with the adjacent stroma can be highly specific and spatially ordered [9].

GM-CSF (granulocyte/macrophage colony-stimulating factor), IL (interleukin)-3 and IL-5 are central haemopoietins that act through a common  $\beta$ -receptor chain. They control proliferation, differentiation and activation of myeloid cells [10]. Besides their function in the bone marrow myelopoiesis, GM-CSF and IL-5 also participate in tissue repair and regeneration [11,12]. Chronic activation of GM-CSF production in injured tissues leads to exacerbated reactions, such as proliferation and activation of myofibroblasts and formation of granulation tissue and/or fibrosis [13,14]. GM-CSF expression also occurs in many cancer cells, correlating with the tumour aggressiveness and the extension of pericancerous stroma reaction [15]. Local controls of GM-CSF activities are thus relevant for pathobiology and evolution of tissue reactions in pathological states.

Experimental *in vitro* models of myelopoiesis have been well established [16]. One of the major findings is the fact that relatively high concentrations of haemopoietins are required to drive proliferation of myeloid progenitors in liquid cell cultures,

Abbreviations used: CMF-BSS, calcium- and magnesium-free balanced salt solution; DMEM, Dulbecco's modified Eagle's medium; FBS, fetal bovine serum; GM-CSF, granulocyte/macrophage colony-stimulating factor; GM-CSF-R $\alpha$ , GM-CSF-receptor  $\alpha$  subunit; HPTLC, high-performance TLC; IL, interleukin; PDMP, D,L-threo-1-phenyl-2-decanoylamino-3-morpholino-1-propanol.

<sup>1</sup> To whom correspondence should be addressed (email radovan@histo.ufrj.br).

whereas very low or barely detectable levels are sufficient in cultures provided with the myelosupportive stroma. Several studies have indicated that the mitogenic activity of GM-CSF can be modulated by the glycosaminoglycan moiety of heparan sulphate proteoglycans produced by stromal cells, which could either enhance or abrogate the biological availability or activity of the cytokine [17–19]. We have also shown [20] that the polarity of the intercellular micro-environment was determinant for optimal interaction between GM-CSF and glycosaminoglycans. This was sensitive to digestion with sialidase. In view of these findings, we proposed that clusters of negatively charged molecules, such as sialylated glycoproteins and glycosphingolipids (gangliosides), associated with membranes of stromal cells, haematopoietic progenitors or both cell types, could be determinants for the local physical and chemical properties leading to the decreased pH, which induces the conformational change in GM-CSF required for its optimal interaction with glycosaminoglycans [6,20,21]. In the present study, we have addressed the question of the production, distribution and role of gangliosides in the experimental model of *in vitro* myelopoiesis in the presence of a myelosupportive stroma. We have used the FDC-P1 cell line, which is dependent upon GM-CSF for both survival and proliferation, as a reporter system to monitor the bioavailability and the local activity of GM-CSF.

## EXPERIMENTAL

### Materials

DMEM (Dulbecco's modified Eagle's medium), RPMI 1640 medium, C<sub>6</sub>-NBD-ceramide {*N*-[7-(4-nitrobenzo-2-oxa-1,3-diazole)]-6-aminohexanoyl-D-erythro-sphingosine}, FITC-conjugated cholera toxin, rabbit anti-(cholera toxin) antibody and standard lipids were from Sigma–Aldrich; FBS (fetal bovine serum) was from Cultilab; D-[U-<sup>14</sup>C]galactose (300 mCi/mmol) was from Amersham BioSciences; plastic tissue-culture dishes were Nunc; Fluorsave<sup>®</sup> and PDMP (D,L-threo-1-phenyl-2-decanoylamino-3-morpholino-1-propanol) were from Calbiochem; silica-gel HPTLC (high-performance TLC) plates were from Merck; a rabbit polyclonal antibody to GM-CSF-R $\alpha$  (GM-CSF-receptor  $\alpha$  subunit) was from Santa Cruz Biotechnology; FITC-conjugated goat anti-mouse serum, CY3-conjugated goat anti-mouse serum and TRITC-conjugated donkey anti-rabbit serum were from Jackson ImmunoResearch. DH2, a monoclonal anti-G<sub>M3</sub> antibody, was generously supplied by Dr Sen-itiroh Hakomori (University of Washington, Seattle, WA, U.S.A.).

### Cells and cell cultures

The murine AFT-024, FDC-P1 and WeHi-3B cell lines were obtained from the Rio de Janeiro Cell Bank (PABCAM, Federal University, Rio de Janeiro, Brazil). AFT-024 (established originally from fetal liver) and WeHi-3B cells were maintained in RPMI 1640 and DMEM respectively, supplemented with 10% (v/v) FBS. The FDC-P1 cell line was maintained in RPMI 1640 medium containing 10% (v/v) FBS and 10% (v/v) supernatant of WeHi-3B cells, which secrete IL-3 and GM-CSF constitutively. Cultures were maintained under a humidified 5% CO<sub>2</sub> atmosphere at 37°C. For a typical co-culture experiment, FDC-P1 cells were washed with CMF-BSS (calcium- and magnesium-free balanced salt solution) in order to remove IL-3 and GM-CSF, inoculated on to the semi-confluent monolayers of AFT-024 cells in 24-well plates (1 × 10<sup>5</sup> cells/well) and maintained in culture for 48 h. Cell proliferation was monitored under a phase-contrast-equipped inverted microscope. For co-cultures under PDMP treatment, the AFT-024 culture was maintained

with 10  $\mu$ M PDMP for 48 h, FDC-P1 cells were then inoculated on to the semi-confluent monolayers, as described above, and maintained in culture for 24, 48 and 72 h in the presence of 10  $\mu$ M PDMP. The PDMP-containing medium was renewed every 24 h. Controls were done using FDC-P1 cultures maintained in standard AFT-024 supernatant supplemented at the beginning of the culture with 10  $\mu$ M PDMP.

### Determination of the FDC-P1 proliferation rate

FDC-P1 cells were inoculated on to 24-well culture plates at 1 × 10<sup>5</sup> cells/well and maintained in three different medium preparations: (i) RPMI supplemented with 10% (v/v) FBS and 50% (v/v) medium conditioned by AFT-024 cells; (ii) RPMI supplemented with 10% (v/v) FBS and 50% (v/v) medium conditioned by AFT-024 cells treated with 10  $\mu$ M PDMP; (iii) RPMI supplemented with 10% (v/v) FBS and 50% (v/v) medium conditioned by AFT-024 cells and addition of 10  $\mu$ M PDMP; and (iv) RPMI supplemented with 10% (v/v) FBS and 50% (v/v) medium conditioned by AFT-024 cells and 20% (v/v) DH2 hybridoma supernatant (producing monoclonal anti-G<sub>M3</sub> antibody). After 24 h, the viable cells were counted on a haemocytometer. RPMI supplemented with only 10% (v/v) FBS was used as the negative control and RPMI supplemented with 10% (v/v) FBS and 10% (v/v) WeHi-3B-cell-conditioned medium was used as the positive control. In order to monitor the potential non-specific effect of DH2 over FDC-P1 proliferation, cells were maintained in positive-control medium with 20% (v/v) DH2.

### Metabolic labelling and lipid extraction

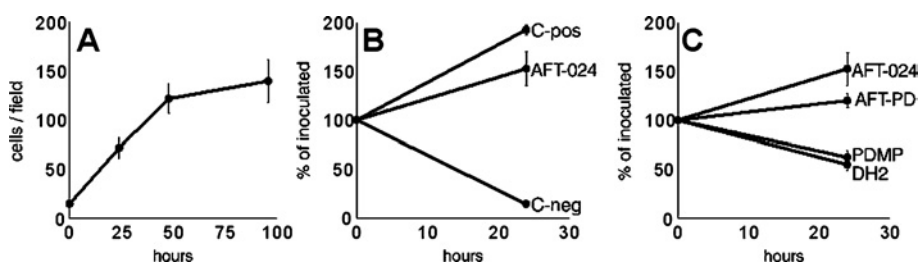
Cultures of stromal cells that reached confluence and cultures of FDC-P1 cells were incubated with 0.5  $\mu$ Ci/ml [<sup>14</sup>C]galactose for 12 h. Subsequently, cells were washed three times with ice-cold PBS, harvested from the plate and pelleted by a brief centrifugation [22]. Lipids were extracted from radiolabelled cell pellets with chloroform/methanol (2:1, v/v), and glycosphingolipids were purified using a Sep-Pack C<sub>18</sub> column [23]. Inhibition of PDMP was determined after incubation of stromal cells for 24 and 72 h with 10  $\mu$ M PDMP and incubation with 0.5  $\mu$ Ci/ml [<sup>14</sup>C]galactose for the last 12 h of incubation.

### Chromatography and analysis of gangliosides

The purified lipid extracts were evaporated under N<sub>2</sub> and run on HPTLC silica gel 60 plates with two successive solvent systems: chloroform/methanol (4:1, v/v) and chloroform/methanol/0.25% aqueous CaCl<sub>2</sub> (15:9:2, by vol.). Lipids were developed in a tank as described by Nores et al. [24]. Radiolabelled sphingolipids were visualized by exposing to a radiographic film at –70°C, and their relative content was determined by densitometric scanning of the X-ray film in a CS 930 Shimadzu UV/vis densitometer. The standards were visualized by exposure to resorcinol/HCl [25]. Ganglioside designations are based on TLC co-migration with standards. The identity of labelled phosphatidylcholine was confirmed by alkaline hydrolysis and co-migration with purified phospholipid standard as described by Rosales-Fritz et al. [26].

### Immunocytochemistry

Semi-confluent monolayers of AFT-024 and co-cultures (maintained for 12 h) grown on coverslips were fixed in 4% (v/v) paraformaldehyde for 30 min, washed in PBS and incubated in 3% (w/v) BSA-containing PBS for 1 h at 37°C to block non-specific binding sites. FDC-P1 cells were washed with PBS, and 5 × 10<sup>4</sup> cells were suspended in 100  $\mu$ l of PBS. They were



**Figure 1** Myelosupportive activity of AFT-24 cells for survival and proliferation of FDC-P1 cells

(A) FDC-P1 cell proliferation in co-culture with AFT-024 cells. (B) FDC-P1 cell proliferation in the presence of the supernatant of AFT-024 cells, of the supernatant of WeHi-3B cells as the positive control (C-pos), and in the RPMI 1640 culture medium supplemented with 10% (v/v) FBS as the negative control (C-neg). (C) Inhibition of the FDC-P1 cell proliferation in the presence of AFT-024 cell supernatant harvested from cultures in which the synthesis of gangliosides was inhibited by PDMP, standard AFT-024 cell supernatant in the presence of 10  $\mu$ M PDMP added at the beginning of culture (AFT-PD) or AFT-024 cell supernatant in the presence of DH2 (a monoclonal antibody to  $G_{M3}$ ) compared with the proliferation in the presence of the standard AFT-024 cell supernatants (AFT-024).

cytocentrifuged at 40  $g$  on to coverslips, fixed in 4% (v/v) paraformaldehyde for 30 min and the non-specific binding sites were blocked as described above. The immunocytochemistry procedures were done as described by Crespo et al. [25]. The primary antibodies were: mouse monoclonal anti- $G_{M3}$  (clone DH2) and rabbit polyclonal anti-(GM-CSF- $R\alpha$ ) (1:200) antibodies. The secondary antibodies were: CY3-conjugated goat anti-mouse Ig (1:1000), FITC-conjugated goat anti-mouse Ig (1:40) or TRITC-conjugated goat anti-(rabbit Ig) (1:500) antibodies. After a final washing with PBS, specimens were mounted with Fluor-save. Confocal images were obtained using a Carl Zeiss LSM5 Pascal laser scanning confocal microscope (Facultad de Ciencias Químicas, Universidad Nacional de Córdoba, Argentina) equipped with an argon/helium/neon laser and a 63 $\times$  (1.4 numerical aperture) oil-immersion objective (Zeiss Plan-Apochromat). For  $G_{M1}$  immunostaining, two methods were used: (i) fixed cells were incubated with 4  $\mu$ g/ml non-toxic cholera toxin B subunit, followed by mouse monoclonal anti-(cholera toxin) antibody; and (ii) fixed cells were incubated with FITC-conjugated non-toxic cholera toxin B subunit. Other immunocytochemistry procedures were the same as described above.

### Shedding of gangliosides

The AFT-024 and FDC-P1 cells were metabolically labelled with 0.5  $\mu$ Ci/ml [ $^{14}$ C]galactose for 12 h, washed and cultured in fresh medium for 48 h. Subsequently, the culture supernatant was collected, dialysed against distilled water for 24 h, lyophilized and the total lipids were extracted from the lyophilized powder with chloroform/methanol (1:1, v/v). The cells were harvested and lipids extracted with chloroform/methanol (2:1, v/v) as indicated above. The gangliosides present in the cells and in the supernatant were purified and analysed [27].

### Uptake of radioactive lipids from AFT-024-conditioned medium by FDC-P1 cells

AFT-024 cells were metabolically labelled with 0.5  $\mu$ Ci/ml [ $^{14}$ C]galactose for 12 h, washed and cultured in fresh medium for 48 h. The culture supernatant was collected and centrifuged at 1200  $g$  for 20 min to eliminate cells and debris. Subsequently, FDC-P1 cells were cultivated overnight in RPMI supplemented with 10% (v/v) FBS and 50% (v/v) AFT-024 cell-conditioned medium. The cells were pelleted by brief centrifugation and lipids were extracted, purified and analysed as described previously [25]. The same protocol was applied to AFT-024 cells treated previously with 10  $\mu$ M PDMP for 72 h and maintained in the presence of PDMP in order to obtain the conditioned medium.

### Labelling of co-cultures

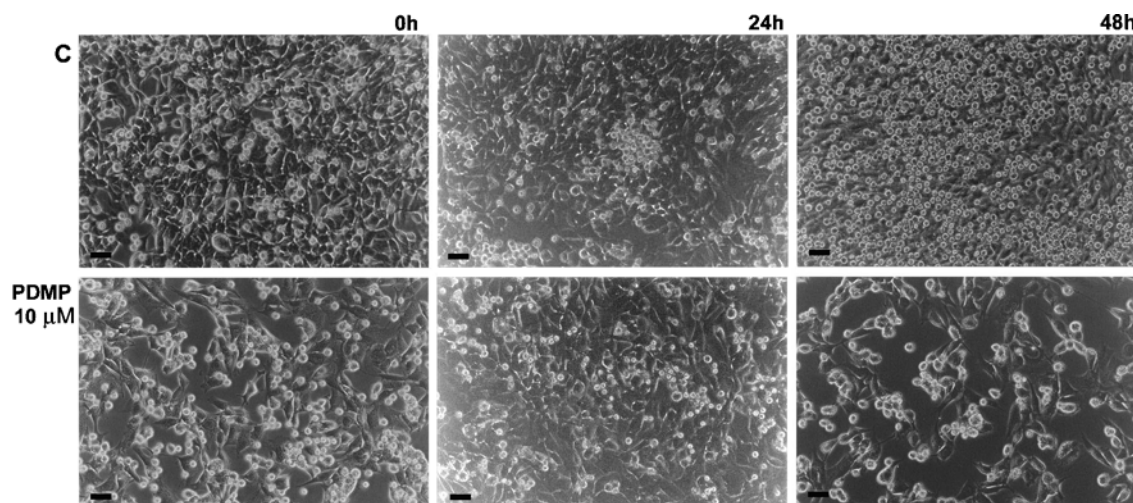
Co-cultures were submitted to different metabolic labelling protocols. (i) COC group: AFT-024 and FDC-P1 cells were incubated with 0.5  $\mu$ Ci/ml [ $^{14}$ C]galactose for the last 12 h of 48 h co-cultures. (ii) ST group: semi-confluent monolayers of AFT-024 cells were incubated with 0.5  $\mu$ Ci/ml [ $^{14}$ C]galactose for 12 h, washed three times with CMF-BSS and FDC-P1 cells were inoculated and co-cultured for the following 48 h. (iii) PDMP group: AFT-024 cells were incubated with 0.5  $\mu$ Ci/ml [ $^{14}$ C]galactose for 12 h, washed three times with CMF-BSS and FDC-P1 cells treated previously with 10  $\mu$ M PDMP for 6 h were inoculated on to the semi-confluent monolayers of AFT-024 cells and co-cultured for the following 48 h in the presence of PDMP. (iv) PREC group: FDC-P1 cells were incubated with 0.5  $\mu$ Ci/ml [ $^{14}$ C]galactose for 12 h, washed three times with CMF-BSS, inoculated on to the semi-confluent monolayers of AFT-024 cells and co-cultured for the following 48 h. For harvesting FDC-P1 cells from co-cultures, the medium was collected and the culture plates were washed three times with PBS at room temperature, under gentle stirring. FDC-P1 cells were harvested from washes. Under these conditions, a fraction of adherent FDC-P1 cells remained attached to the AFT-024 cell stroma.

### Exosome preparations

Exosomes were collected from the medium of the ST and PDMP groups described above. Supernatants were centrifuged at 300  $g$  for 5 min and at 1200  $g$  for 20 min in order to eliminate cells and debris, followed by a centrifugation for 1.5 h at 70 000  $g$ . Two fractions were obtained: a high-density (pellet) and a low-density (hypodense) fraction. The exosomes concentrated in the pellet were washed in a large volume of PBS and centrifuged at 70 000  $g$  for 1.5 h [28]. The lipids were extracted, purified and analysed as described previously [28].

## RESULTS

FDC-P1 cells proliferated both in co-culture with AFT-024 stroma (Figures 1A and 2) and in the presence of the AFT-024 cell-culture supernatant (Figure 1B). This is in accordance with our previous study [29] showing that AFT-024 cells, although lacking IL-3, produced and secreted high amounts GM-CSF, as well as heparan-sulphate-bearing proteoglycans that interacted with GM-CSF. These stroma cells could thus be used for studies on the GM-CSF-dependent myeloid cell proliferation. Inhibition of ganglioside synthesis in the stroma layer with PDMP significantly decreased the myelosupportive capacity in both experimental



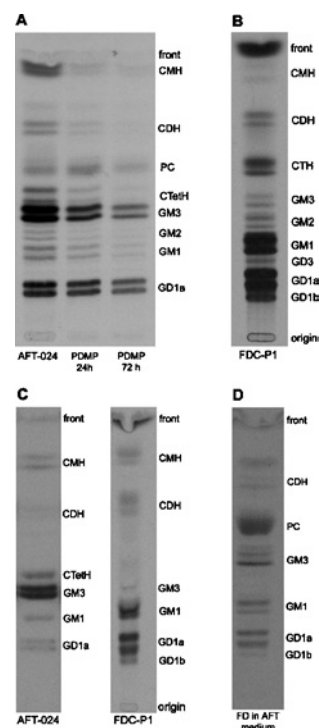
**Figure 2** Inhibition of FDC-P1 cell survival and proliferation in co-culture with AFT-024 cells in which ganglioside synthesis was inhibited by PDMP

FDC-P1 cells washed previously with CMF-BSS were inoculated on to the semi-confluent monolayers of AFT-024 cells at  $1 \times 10^5$  cells/well and maintained for 0, 24 and 48 h. C, co-cultures maintained in standard conditions. PDMP, co-cultures maintained with  $10 \mu\text{M}$  PDMP after inoculation of FDC-P1 cells upon AFT-024 semi-confluent cell monolayer treated previously with the inhibitor for 48 h. Phase-contrast microscopy of the cells is shown. Scale bar,  $20 \mu\text{m}$ .

conditions (Figures 1C and 2), indicating that the synthesis and release of gangliosides into the supernatants were required for FDC-P1 cell proliferation. To exclude a direct toxic effect of PDMP on FDC-P1 cells, supernatant was collected from standard AFT-024 cell cultures and PDMP was added prior to its addition to FDC-P1 cells. Results indicated a small decrease in FDC-P1 proliferation, but this direct effect of PDMP was much lower than the inhibition observed when gangliosides shed by stroma cells was reduced due to PDMP treatment (Figure 1C).

We had suspected previously that sialic-acid-containing molecules played an important role in modulating the cross-talk between stromal and myeloid cells, since capping of negatively charged molecules, sensitive to sialidase digestion, occurred in both cells at the site of interaction [21]. For this reason, we next analysed ganglioside synthesis in AFT-024 and FDC-P1 cells (Figures 3A and 3B, and Table 1). AFT-024 accumulated essentially  $\text{G}_{\text{M}3}$  and, to a lower extent,  $\text{G}_{\text{D}1\text{a}}$  and  $\text{G}_{\text{M}1}$ . This is similar to other myelosupportive stromata that we have studied: the S17 cell line derived from the bone marrow (F. C. R. Guma, unpublished work) and the liver connective tissue cells involved in inflammatory fibro-granulomatous reactions that have a tissue origin similar to AFT-024 cells (C. M. B. Andrade, unpublished work). Conversely, FDC-P1 cells accumulated only a low amount of  $\text{G}_{\text{M}3}$ , which was metabolized further to several gangliosides of the *a* and *b* series.  $\text{G}_{\text{M}1}$  was the major ganglioside accumulated by FDC-P1 cells. All major gangliosides ran as doublets, in accordance with the fact that sphingolipids have differences in ceramide structures [30,31].

The required presence of gangliosides in the stroma-mediated support of myelopoiesis was shown by PDMP-dependent inhibition of their synthesis. Exposure of AFT-024 cells to  $10 \mu\text{M}$  PDMP for 3 days led to a noticeable decrease in ganglioside synthesis (57 and 82% in 24 and 72 h respectively) and in their precursors lactosylceramide [CDH (ceramide dihexoside)] and glucosylceramide [CMH (ceramide monohexoside)] (Figure 3A). We questioned whether the major ganglioside of the stroma,  $\text{G}_{\text{M}3}$ , was specifically involved in this interaction. In the presence of DH2 (a monoclonal antibody against  $\text{G}_{\text{M}3}$ ), inhibition of FDC-P1 cell proliferation was equivalent to that elicited by



**Figure 3** Analysis of glycosphingolipids in AFT-024 and FDC-P1 cells

(A) Synthesis of glycosphingolipids in AFT-024 cells. Cell cultures were incubated for 12 h with  $[^{14}\text{C}]$ galactose and for the last 12 h of treatment with  $10 \mu\text{M}$  PDMP for 24 or 72 h. (B) Synthesis of glycosphingolipids in FDC-P1 cells. (C) Glycosphingolipids shed by AFT-024 and FDC-P1 cells. Cell cultures were incubated 12 h with  $[^{14}\text{C}]$ galactose and the lipids were shed into the medium for 48 h. (D) FDC-P1 uptake of glycosphingolipids from AFT-024 cell-conditioned medium. AFT-024 cultures were incubated for 12 h with  $[^{14}\text{C}]$ galactose and the lipids were shed into the medium for 48 h. FDC-P1 cells washed previously with CMF-BSS were then inoculated in RPMI supplemented with 10% (v/v) FBS and 50% (v/v) of medium conditioned by AFT-024 cells. Lipids were extracted, purified, analysed by HPTLC and visualized by fluorography. The radioactive bands correspond to the lipids indicated. CTetH: ceramide tetrahexoside; CTH, ceramide trihexoside; CDH, ceramide dihexoside; CMH, ceramide monohexoside; PC, phosphatidylcholine. Gangliosides are named according to Svennerholm [47].

**Table 1** Densitometric analysis of metabolically labelled lipids from AFT-024 and FDCP-1 cells

Cell cultures were incubated for 12 h with [ $^{14}\text{C}$ ]galactose and lipids were extracted, purified, analysed by HPTLC and visualized by fluorography. Results are expressed as a percentage of the total radioactivity incorporated.  $\text{G}_{\text{D1b}}$ ,  $\text{G}_{\text{D1a}}$ ,  $\text{G}_{\text{M1}}$ ,  $\text{G}_{\text{M2}}$ ,  $\text{G}_{\text{D3}}$  and  $\text{G}_{\text{M3}}$  are gangliosides named according to Svennerholm [47]; CtetH, ceramide tetrahexoside; CTH, ceramide trihexoside; CDH, ceramide dihexoside; CMH, ceramide monohexoside; PC, phosphatidylcholine; n.d., not detected.

Lipids	Proportion of glycolipids (% of total radioactivity)	
	AFT-024	FDC-P1
$\text{G}_{\text{D1b}}$	n.d.	6.9
$\text{G}_{\text{D1a}}$	20.6	26.7
$\text{G}_{\text{D3}}$	n.d.	2.4
$\text{G}_{\text{M1}}$	3.6	24.2
$\text{G}_{\text{M2}}$	0.9	3.8
$\text{G}_{\text{M3}}$	52.8	2.6
CtetH	7.0	n.d.
PC	2.5	n.d.
CTH	n.d.	9.7
CDH	3.0	4.0
CMH	8.1	n.d.

the overall inhibition of ganglioside synthesis, indicating the crucial role of  $\text{G}_{\text{M3}}$  in the myelosupportive function of the stroma (Figure 1C). Additionally, we performed experiments to discard a direct inhibition of cell proliferation by DH2. FDC-P1 cells were incubated in the standard culture medium containing DH2 at the same concentration as used in the experiment in Figure 1(C). No effect of DH2 on FDC-P1 cell proliferation was found, clearly excluding a direct effect of DH2 (results not shown).

Since the response to GM-CSF stimulation requires its interaction with the FDC-P1 membrane through binding to the GM-CSF-R, we investigated the spatial distribution of gangliosides and GM-CSF-R $\alpha$  on FDC-P1 cells (Figure 4). In accordance with the TLC analysis of metabolic incorporation of [ $^{14}\text{C}$ ]galactose (Figures 3A and 3B, and Table 1), AFT-024 cells displayed a strong labelling with DH2 (monoclonal anti- $\text{G}_{\text{M3}}$  antibody) in the perinuclear region and a faint labelling of cell membranes (Figure 4A). Conversely, the  $\text{G}_{\text{M3}}$  labelling of FDC-P1 cells was barely detectable (Figure 4B). However, when co-cultures were studied, an intense labelling of  $\text{G}_{\text{M3}}$  was observed on the membranes of FDC-P1 cells, but not in the perinuclear region. Instead, labelling disclosed a full capping on one side of the cell only (Figure 4C). A similar labelling is shown in Figures 4(D) and 4(G) and it can be compared with the labelling of the GM-CSF-R $\alpha$  (Figures 4E and 4H). The merged images disclosed that GM-CSF-R $\alpha$ -stained areas co-distributed almost completely with  $\text{G}_{\text{M3}}$  staining, despite extensive areas of  $\text{G}_{\text{M3}}$  staining outside of areas of GM-CSF-R $\alpha$  staining (Figures 4F and 4I). In many cases, the capping was observed at the contact regions both between two FDC-P1 cells and between stroma and FDC-P1 cells (Figures 4Ia', 4Ib' and 4J–4M).

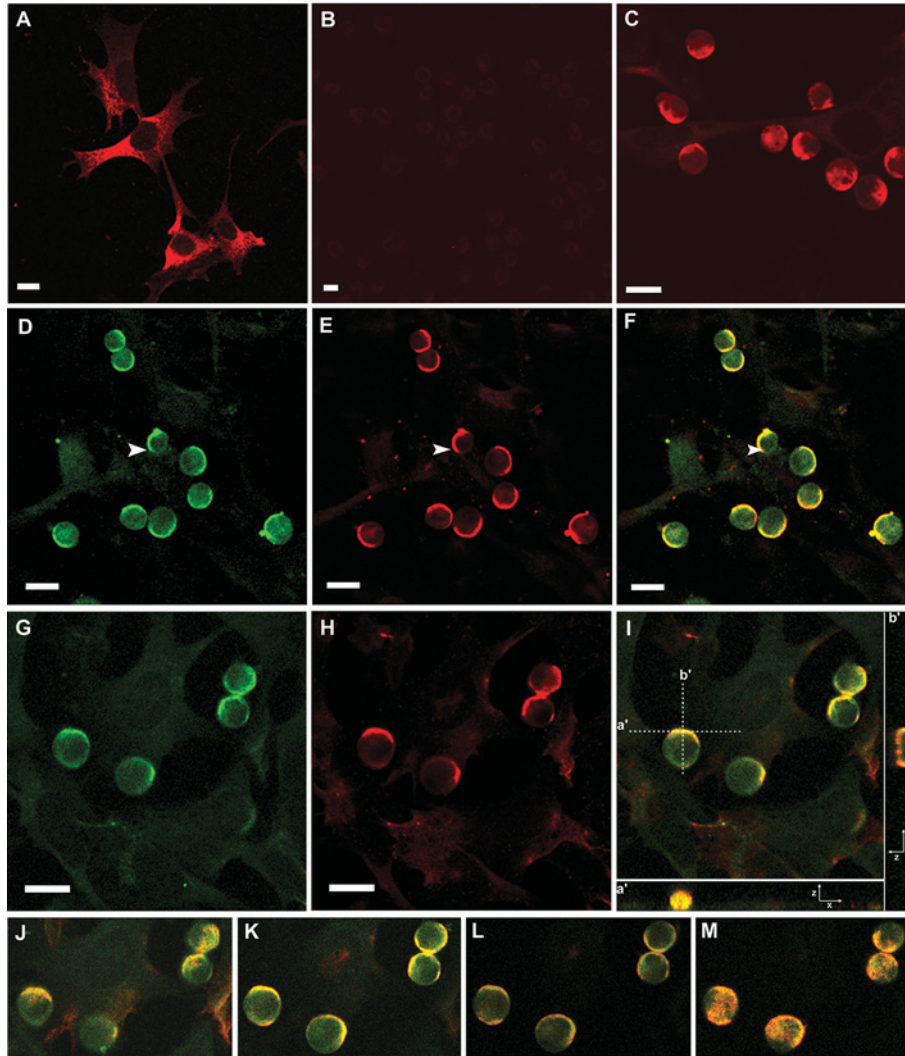
Taking into consideration that  $\text{G}_{\text{M1}}$  is the major ganglioside produced by FDC-P1 cells and to a lesser extent in AFT-024 cells (Figures 3A and 3B, and Table 1), the subcellular distribution of this lipid was analysed and compared with the distribution of  $\text{G}_{\text{M3}}$ .  $\text{G}_{\text{M1}}$  was present homogeneously on the membranes of both the stroma and myeloid cells (Figures 5A and 5B respectively). When co-cultures were studied, the pattern of  $\text{G}_{\text{M1}}$  distribution was similar to  $\text{G}_{\text{M3}}$ . A significant capping of  $\text{G}_{\text{M1}}$  was observed at the contact region between AFT-024 and FDC-P1 cells (Figures 5C and 5D). However, the analysis of GM-CSF-R $\alpha$

distribution disclosed only a partial co-distribution of the receptor staining with  $\text{G}_{\text{M1}}$  staining, mainly at the contact region (Figures 5E–5G).

The increased presence of  $\text{G}_{\text{M3}}$  on FDC-P1 cells in co-culture with AFT-024 raised the question of the mechanism of this apparent contribution of the stroma to the ganglioside composition of the myeloid cell membrane. In order to provide direct evidence of the functional intercellular transfer of sphingolipids in the model of myelosupportive stroma, shedding of radiolabelled gangliosides from AFT-024 and FDC-P1 cells was monitored (Figure 3C). The major ganglioside shed by the stroma cells was  $\text{G}_{\text{M3}}$ , whereas FDC-P1 cells shed essentially  $\text{G}_{\text{M1}}$  and  $\text{G}_{\text{D1a}}$ . The gangliosides harvested from the culture medium represented essentially the pattern of their synthesis in the two cell types, with a relatively low presence of  $\text{G}_{\text{D1a}}$  in the supernatant compared with its presence in AFT-024 cells. When unlabelled FDC-P1 cells were incubated with the supernatant harvested from the AFT-024 cell culture treated with [ $^{14}\text{C}$ ]galactose in which  $\text{G}_{\text{M3}}$  was the major component (Figure 3B), a number of labelled compounds were detected (Figure 3D). The pattern suggests that the  $\text{G}_{\text{M3}}$  captured from the medium was metabolized further in a similar manner to the metabolism of endogenous  $\text{G}_{\text{M3}}$  in FDC-P1 cells (Figure 3B). When the same experiment was done with conditioned medium obtained from AFT-024 cells metabolically labelled and treated with PDMP, the presence of the same ganglioside species in FDC-P1 cells was observed, although the incorporated radioactivity was lower than in control cells (results not shown).

Quantification of the densitometric analysis of  $\text{G}_{\text{M3}}$  and  $\text{G}_{\text{D1a}}$  (Figure 6) in FDC-P1 cells co-cultured with AFT-024 cells showed an increased quantity of labelled  $\text{G}_{\text{M3}}$  (COC group = 20%) when compared with FDC-P1 cells cultivated alone (2.6%). The increased amount of  $\text{G}_{\text{M3}}$  labelled in FDC-P1 cells isolated from co-cultures was clearly detectable when only stroma cells were labelled (ST group = 32%). These results indicate the transfer of  $\text{G}_{\text{M3}}$  from AFT-024 cells to FDC-P1 cells. To exclude the possibility of the  $\text{G}_{\text{M3}}$  synthesis by FDC-P1 cells from metabolically labelled ganglioside precursors captured from AFT-024 cells, the *de novo* synthesis of FDC-P1 glycosphingolipids was inhibited by PDMP before inoculation on to the AFT-024 cellular monolayer (PDMP group) labelled previously. Under this condition, an increased fraction of  $\text{G}_{\text{M3}}$  (22%) was found in FDC-P1 cells isolated from co-culture, indicating again the transfer of  $\text{G}_{\text{M3}}$ . Moreover, when only FDC-P1 cells were metabolically labelled before inoculation on to stroma cells (PREC group),  $\text{G}_{\text{M3}}$  was not detected in FDC-P1 cells isolated from co-cultures. The presence of  $\text{G}_{\text{D1a}}$  in ST and PDMP groups confirms the further metabolism of the captured  $\text{G}_{\text{M3}}$ . Taken together, our results stand in contrast with the data reported for fibroblasts, neurons and neuroblastoma cells that incorporated 1- $^3\text{H}$ ]sphingosine and synthesized labelled sphingolipids [32]. In this study, the released sphingolipids were taken up by cells, but they segregated to lysosomes and were entirely catabolized, indicating that, in this case, trafficking of sphingolipids to and from the extracellular environment does not allow the modification of cell membrane composition [32].

The results shown in Figures 4 and 5 are in agreement with our previous observation of extensive capping of raft-like structures on membranes of the stroma in contact with FDC-P1 cells [21]. Taken together, these data suggest formation of specific regions of the membranes enriched in proteoglycans, glycosphingolipids and growth factor receptors at the interface between the stroma and target cells. This model is compatible with a non-random transfer of membrane compounds between the adjacent stroma and the myeloid cells that represents a new model of co-operation between the myelosupportive stromata and the target cells. We



**Figure 4** Subcellular location of  $G_{M3}$  and GM-CSF-R $\alpha$  in cell cultures and co-culture systems

Cells were immunostained for  $G_{M3}$  with DH2 and immunostained for GM-CSF-R $\alpha$  with a polyclonal anti-(GM-CSF-R $\alpha$ ) antibody. (A) AFT-024 cells immunostained for  $G_{M3}$ . (B) FDC-P1 cells immunostained for  $G_{M3}$ . (C) Co-culture immunostained for  $G_{M3}$ . (D and G) Co-cultures immunostained for  $G_{M3}$  with monoclonal antibody DH2. (E and H) Co-cultures immunostained for GM-CSF-R $\alpha$ . (F) Merged image from (D) and (E). (I) Merged image of (G) and (H). Single confocal sections ( $x$ ,  $y$ ) of  $0.7 \mu\text{m}$  were taken parallel to the coverslip, except for (I) in which a series of  $z$  sections were collected and displayed using the ortho-mode in LSM5 Pascal software provided with the microscope. A single  $x$ ,  $z$  section ( $a'$ ) is shown at the bottom of (I) and a  $y$ ,  $z$  section ( $b'$ ) at the right of (I). (J–M) are single  $x$ ,  $y$  sections from the same  $z$  stack of (I) (from bottom to top). The arrows indicate capping of  $G_{M3}$  (D), GM-CSF-R $\alpha$  (E) and co-capping of  $G_{M3}$  and GM-CSF-R $\alpha$  (F). Scale bars,  $10 \mu\text{m}$ .

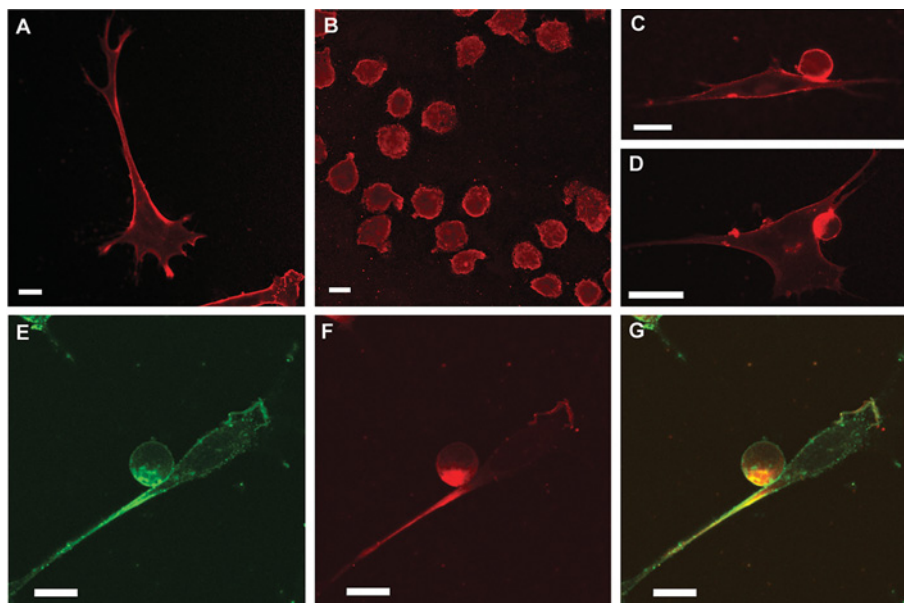
questioned further whether the intercellular transfer of membrane regions with sorted compounds, which potentially represent operational molecular units, could be achieved by the release and subsequent uptake of exosomes. The exosomal fraction was isolated by ultracentrifugation, as described previously by Skokos et al. [28]. We found that 40% of radiolabelled gangliosides released into the supernatant in a co-culture containing AFT-024 and FDC-P1 cells were found in the exosome fraction.

## DISCUSSION

The concept of tissue stromata has gained increasing interest in view of the extending use of cell therapies, based upon introduction of exogenous stem or progenitor cells into the tissues that require repair or regeneration. It is expected that the introduced cells can receive sufficient information from the stroma in order to

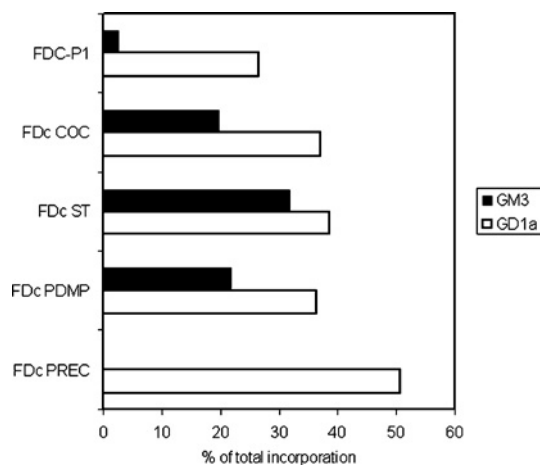
proliferate within the physiological limits and to differentiate into the required cell types in the injured or degenerated tissues. Cells have also to re-establish appropriate spatial tissue organization. Tissue stromata have been studied extensively in the context of providing the required cytokines and chemokines, as well as mediators of homeostasis or inflammatory reactions and repair. They are also known to provide extracellular matrix that provides spatial tissue support. The present study indicates that, in addition, stroma cells may also supply membrane elements to the target cells, which can considerably modulate their ability to respond to given cytokines by segregating the receptors and enhancing or inhibiting intracellular signalling cascades.

We have demonstrated in the present and in parallel studies (C. M. B. Andrade and F. C. R. Guma, unpublished work) that  $G_{M3}$  is the major ganglioside in different myelosupportive stroma cells, such as AFT-024, S17 and GR cells, but it is only a minor ganglioside in the myeloid precursor FDC-P1 cells, although it



**Figure 5** Subcellular location of  $G_{M1}$  and GM-CSF-R $\alpha$  in cell cultures and co-culture systems

$G_{M1}$  was examined by incubation of cholera toxin, followed by a monoclonal anti-(cholera toxin) antibody or by the fluorescence of FITC-conjugated cholera toxin. Cells were immunostained for GM-CSF-R $\alpha$  with a polyclonal anti-(GM-CSF-R $\alpha$ ) antibody. (A) AFT-024 cells immunostained for  $G_{M1}$ . (B) FDC-P1 cells immunostained for  $G_{M1}$ . (C–E) Co-cultures immunostained for  $G_{M1}$ . (F) Co-cultures immunostained for GM-CSF-R $\alpha$ . (G) Merged images from (E) and (F). Single confocal sections of 0.7  $\mu$ m were taken parallel to the coverslip. Scale bars, 10  $\mu$ m.



**Figure 6** Analysis of FDC-P1 uptake of gangliosides  $G_{M3}$  and  $G_{D1a}$  from AFT-024-cell-conditioned medium

Results are expressed as the percentage of total radioactivity incorporated. FDC corresponds to myeloid progenitors isolated after co-culture submitted to different metabolic labelling protocols as described in the Experimental section. COC, metabolic labelling during the last 12 h of 48 h of co-culture; ST, metabolic labelling only of stromal cells prior to co-culture for 48 h; PDMP, metabolic labelling only of stromal cells and treatment of FDC-P1 cells with 10  $\mu$ M PDMP prior to co-culture for 48 h in the presence of 10  $\mu$ M PDMP; PREC, metabolic labelling only of myeloid precursor cells prior to co-culture for 48 h.

is required for their optimal response to GM-CSF. In the present study, we describe the transfer of  $G_{M3}$  from stroma to FDC-P1 cells, the capping of  $G_{M3}$  at the contact regions between the two cell types and the co-localization of GM-CSF-R $\alpha$  with  $G_{M3}$  on the myeloid cells. The GM-CSF-R $\alpha$  co-localizes only partially with ganglioside  $G_{M1}$ , a minor constituent of stroma cells, although the capping of  $G_{M1}$  by both cells at the contact regions was detectable, suggesting that  $G_{M3}$  and  $G_{M1}$  are located in different

membrane regions or in different rafts, as shown previously in other studies [33].

The biological significance of gangliosides shedding from normal and tumour cells and their effect on other cells has been well documented [27,34–36]. Gangliosides are shed from the cell plasma membrane as monomers, micelles or membrane vesicles enriched in caveolin-1 [37,38]. It has been also reported that exosomes, small vesicles formed from the fusion of multivesicular endosomes with the cell surface, could be transferred from professional to non-professional antigen-presenting cells for effective T-cell stimulation [39,40]. In this model, exosomes carry a number of immunoregulatory molecules and may be considered as immunologically sophisticated vectors for various antigens [41]. In the present study, we have shown that AFT-024 cells transfer  $G_{M3}$  to FDC-P1 cells. This transfer can be done by a direct contact between the cells or by uptake of  $G_{M3}$  shed to the extracellular environment. A random shedding and uptake of gangliosides is less probable, in view of their specific localization and metabolism on the target cells. The proliferation of FDC-P1 cells in the supernatant of AFT-024 cells was dependent on  $G_{M3}$  and we found that gangliosides were shed in the exosomal fractions. Therefore the exosomes might be an alternative way for secretion and cell uptake of  $G_{M3}$  now described in our experimental model.

Interaction of gangliosides with growth factor receptors has been studied extensively [42,43]. In many cases, glycosphingolipids were reported to have an inhibitory activity on signalling, decreasing the response to cytokines such as EGF (epidermal growth factor), PDGF (platelet-derived growth factor) or FGF (fibroblast growth factor). In contrast, in our model of stroma-dependent myelopoiesis, gangliosides and, in particular,  $G_{M3}$  are required for optimal growth factor signalling. We have already proposed that this is due to generation of a particular negatively charged intercellular space in which the interaction among the growth factor, heparan sulphate proteoglycans and the corresponding receptor are most active [20,21]. We can now propose

the following scenario. The physical contact between stroma and myeloid progenitors elicits extensive capping on both cells and the concentration of selected sets of membrane molecules at the cell interfaces. Stroma cells provide growth factors, which are required for survival and proliferation of myeloid cells. They also supply gangliosides, which are transferred from stroma to target cells, generating on the latter specific membrane domains with molecular complexes that include growth factor receptors surrounded by the associated molecules. It is known that such a formation of lipid rafts containing membrane receptors is often required, particularly in cases where the functional receptor is composed of many interacting units as, for example, the T-lymphocyte cell receptor [44].

Three mechanistic models for modulation of growth factor and the corresponding receptor activity by gangliosides have been proposed: (i) ganglioside–ligand interaction, (ii) ganglioside regulation of receptor dimerization, and (iii) ganglioside modulation of receptor activity and its subcellular localization [43]. The modulation of receptor activity involves regulation of kinase activity, such as PKC (protein kinase C) [45] and Src kinases [46]. In this sense, our results suggest that modulation of GM-CSF-R interaction with the ligand, through formation of macromolecular complexes involving  $G_{M3}$ , may be determinant for signal transduction through phosphorylation cascades. Further studies will be required to determine whether  $G_{M3}$  directly activates the GM-CSF interaction with its receptor or indirectly involves the activity of co-activators or inhibitors segregated in the same membrane microdomains.

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