

# Galatrox is a C-type lectin in *Bothrops atrox* snake venom that selectively binds LacNAc-terminated glycans and can induce acute inflammation

Marco A Sartim<sup>2</sup>, Thalita B Riul<sup>2</sup>, Camillo Del Cistia-Andrade<sup>2</sup>, Sean R Stowell<sup>3</sup>, Connie M Arthur<sup>3</sup>, Carlos A Sorgi<sup>2</sup>, Lucia H Faccioli<sup>2</sup>, Richard D Cummings<sup>4</sup>, Marcelo Dias-Baruffi<sup>1,2</sup>, and Suely V Sampaio<sup>1,2</sup>

<sup>2</sup>Department of Clinical Analyses, Toxicology and Food Sciences, School of Pharmaceutical Sciences of Ribeirão Preto, University of São Paulo, Ribeirão Preto 14040903 São Paulo, Brazil; <sup>3</sup>Department of Pathology and Laboratory Medicine, Center for Transfusion and Cellular Therapies, Emory University School of Medicine, Atlanta 30322, GA, USA; and <sup>4</sup>Department of Biochemistry and The Glycomics Center, Emory University School of Medicine, Atlanta 30322, GA, USA

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Previous studies indicate that snake venom contains glycan-binding proteins (GBPs), although the binding specificity and biological activities of many of these GBPs is unclear. Here we report our studies on the glycan binding specificity and activities of galatrox, a *Bothrops atrox* snake venom-derived GBP. Glycan microarray analysis indicates that galatrox binds most strongly to glycans expressing *N*-acetylglucosamine (LacNAc), with a significant preference for Gal $\beta$ 1-4GlcNAc $\beta$  over Gal $\beta$ 1-3GlcNAc $\beta$  compounds. Galatrox also bound immobilized laminin, a LacNAc-dense extracellular matrix component, suggesting that this GBP can bind LacNAc-bearing glycoproteins. As several endogenous mammalian GBPs utilize a similar binding LacNAc binding preference to regulate neutrophil and monocyte activity, we hypothesized that galatrox may mediate *B. atrox* toxicity through regulation of leukocyte activity. Indeed, galatrox bound neutrophils and promoted leukocyte chemotaxis in a carbohydrate-dependent manner. Similarly, galatrox administration into the mouse peritoneal cavity induced significant neutrophil migration and the release of pro-inflammatory cytokines IL-1 $\alpha$  and IL-6. Exposure of bone marrow-derived macrophages to galatrox induced generation of pro-inflammatory mediators IL-6, TNF- $\alpha$ , and keratinocyte-derived chemokine. This signaling by galatrox was mediated via its carbohydrate recognition domain by activation of the TLR4-mediated MyD88-dependent signaling pathway. These

results indicate that galatrox has pro-inflammatory activity through its interaction with LacNAc-bearing glycans on neutrophils, macrophages and extracellular matrix proteins and induce the release of pro-inflammatory mediators.

**Keywords:** *Bothrops atrox* / C-type lectin / Inflammatory response / Neutrophil migration / Snake venom

## Introduction

Inflammation is critical in the host response to tissue damage and infection, helping the immune system to preserve homeostasis and leading to complex physiological mechanisms characterized by the involvement of vascular and cellular events (Medzhitov and Horng 2009; Soehnlein and Lindbom 2010). Deficiencies in inflammatory responses can increase susceptibility to infectious disease and limit normal wound repair (Netea et al. 2012; White and Mantovani 2013). In contrast, exuberant inflammatory responses are associated with pathophysiology of several diseases, leading to tissue damage and organ injury (Buckley et al. 2013; Huang et al. 2013; Strydom and Rankin 2013). Given the critical balance of inflammation in facilitating host immunity while avoiding injury, many factors appear to have evolved unique roles in the regulation of both these mechanisms (Ariel and Timor 2013; Frey et al. 2013; Lu et al. 2013; Rogala and Glück 2013). Among these factors, glycan-binding proteins (GBPs) play a critical role in inflammation by regulating a wide variety of cells involved in the inflammatory process (Dias-Baruffi et al. 1995a,b; Rabinovich et al. 2000; Nishi et al. 2003; Nieminen et al. 2005; Stowell, Qian, et al. 2008; Sato et al. 2009).

While disruptions to normal host homeostasis can significantly alter the inflammatory balance, environmental factors also appear capable of compromising normal homeostatic processes (Ramos and Fernandez-Sesma 2012; Aziz et al. 2013). Several biological active molecules from *Bothrops* spp. snake venoms are responsible for stimulating an intense inflammation during envenomation, triggering immunological events, such as prominent leukocyte infiltration, and promoting local and systemic disorders (Bonavita et al. 2006; Carneiro et al. 2008; Teixeira et al. 2009). As an example of these factors within venom, the C-type  $\beta$ -galactoside-binding lectins have been reported to be involved in inflammatory response (Lomonte et al. 1990; Havt et al. 2005; Panunto et al. 2006; Elifio-Esposito et al.

<sup>1</sup>To whom correspondence should be addressed: Tel: +55-16-36024286; Fax: +55-16-3602-4719; e-mail: suvilela@usp.br (S V S); Tel: +55-16-36024256; Fax: +55-16-3602-4725; e-mail: mdbaruffi@fcf.usp.br (M D-B)

2011). Given the role of endogenous  $\beta$ -galactoside-binding lectins, such as galectins, within mammals in the regulation of inflammation and immunity (Stowell, Qian, et al. 2008; Sato et al. 2009), these results suggest that snake venom C-type lectins may involve related glycan ligands and also induce changes related to those previously shown for endogenous GBPs.

Galatrox, a C-type lectin isolated from *Bothrops atrox* snake venom, displays significant lactose-binding properties and appears to recognize mammalian glycans (Mendonça-Franqueiro et al. 2011), suggesting that galatrox may engage leukocyte ligands and induce significant inflammatory changes. However, the glycan-binding specificity of galatrox remains undefined. Furthermore, whether galatrox recognizes leukocyte glycan ligands or induces alterations in leukocyte function is not known. In this study, we examined the glycan-binding specificity and proinflammatory activity of galatrox. Our results indicate that galatrox displays similarity in glycan-binding characteristics to endogenous mammalian GBPs and induces significant changes in leukocyte migration and activation that are likely to contribute to host inflammation following *B. atrox* envenomation.

## Results

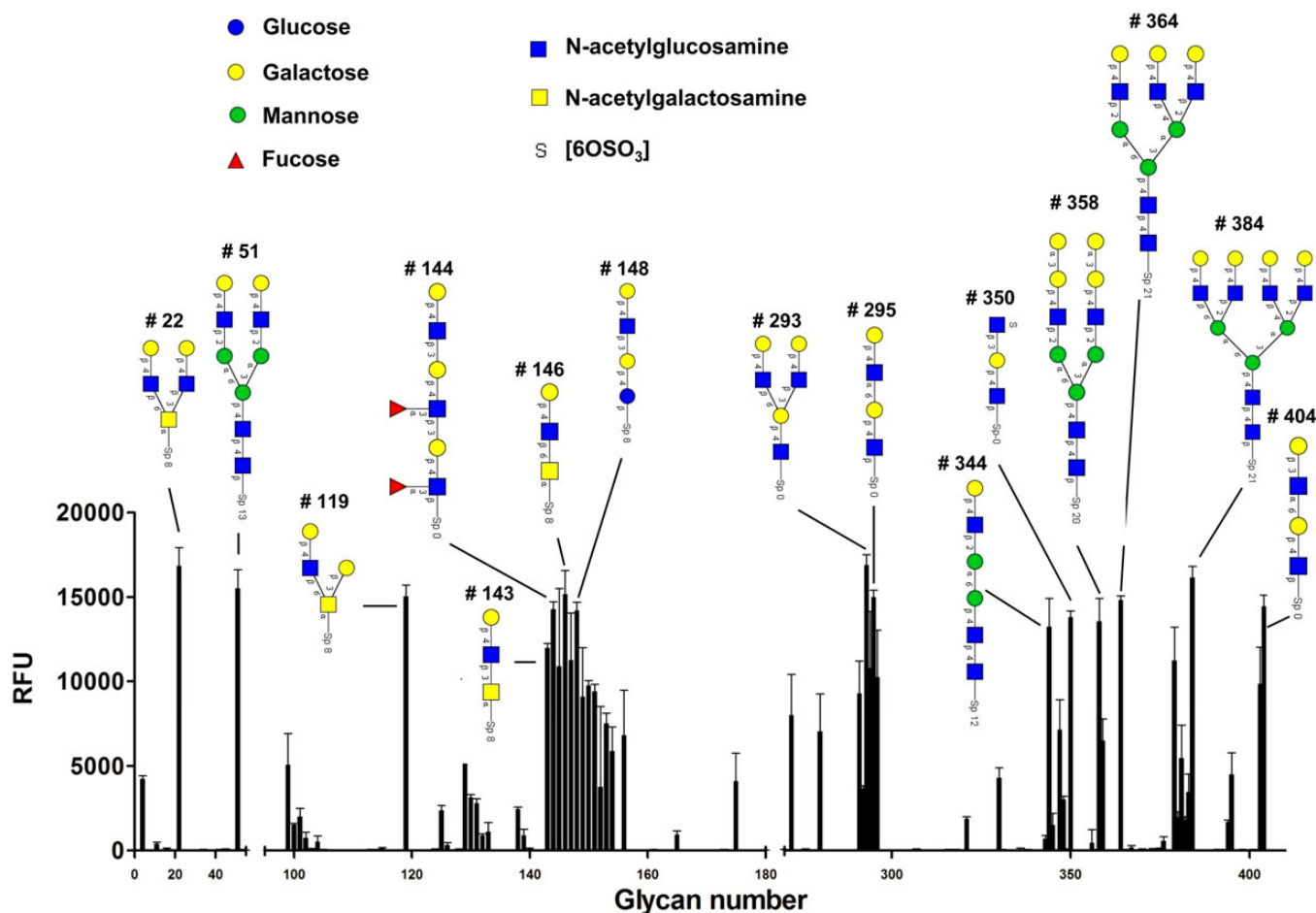
### *Galatrox preferentially binds to glycans terminating in N-acetyllactosamine*

As toxins often utilize host ligands to mediate their activity, we hypothesized that galatrox may induce inflammation through engagement of ligands typically employed by endogenous GBPs. Among mammalian GBPs with  $\beta$ -galactoside-binding activity, several studies demonstrate that members of the galectin family possess remarkable ability to regulate immunity, including the regulation of leukocyte activity (Dias-Baruffi et al. 2003; Nieminen et al. 2005; Garín et al. 2007; Toscano et al. 2007; Stowell, Qian, et al. 2008). However, whether galatrox and galectin family members share similarities in carbohydrate-binding activity remains unknown. To examine this in detail, we turned to the consortium for functional glycomic glycan microarray, which has been previously used to characterize the binding specificity of multiple GBPs. To accomplish this, we directly labeled galatrox with fluorescein isothiocyanate (FITC) and then examined glycan-binding activity on the glycan microarray (Version 3.2) (<http://www.functionalglycomics.org>). Of all the carbohydrates represented on the glycan microarray, FITC-galatrox displayed the highest affinity toward glycans expressing non-reducing *N*-acetyllactosamine (LacNAc) sequences, with a significant preference for type 2 (Gal $\beta$ 1-4GlcNAc $\beta$ ) over type 1 (Gal $\beta$ 1-3GlcNAc $\beta$ ) LacNAc glycans (Figure 1 and Table I). In addition, although FITC-galatrox appeared to display preference for LacNAc presented on extended glycan structures, it did not bind to LacNAc repeats within glycans, as no increases in binding to polyLacNAc could be detected, and internal modifications to LacNAc failed to alter polyLacNAc binding (Supplementary data, Table S1), similar to previous results with several members of the galectin family (Stowell et al. 2004; Stowell, Arthur, Mehta, et al. 2008). Furthermore, FITC-galatrox failed to bind sialylated LacNAc glycans (in both  $\alpha$ 2-3 and/or  $\alpha$ 2-6 linkages) (Figure 1 and Table I), strongly suggesting that sialylation inhibits galatrox LacNAc recognition (Supplementary data, Table S1), similar to previous results on galectin-2 (Stowell, Arthur, Mehta, et al. 2008). Similar inhibition

of LacNAc recognition appears to occur if the glycans are  $\alpha$ 1-2 fucosylated. This weak binding is also observed for either GalNAc $\alpha$ 1-3(Fuc $\alpha$ 1-2)LacNAc (Supplementary data, Table S1) or Gal $\alpha$ 1-3(Fuc $\alpha$ 1-2)-LacNAc (Supplementary data, Table S1) glycans, as present in blood group A and B antigens, respectively. In addition, LacNAc-containing glycans expressing Lewis<sup>x</sup>, Lewis<sup>a</sup> and sialyl-f<sup>x</sup> antigens [Gal $\beta$ 1-4(Fuc $\alpha$ 1-3)GlcNAc-R, Gal $\beta$ 1-3 (Fuc $\alpha$ 1-4)GlcNAc-R and NeuAc $\alpha$ 2-3Gal $\beta$ 1-4(Fuc $\alpha$ 1-3)GlcNAc-R, respectively] also displayed weak binding to galatrox, as well as LacNAc glycans modified to express 3-O, 4-O or 6-O sulfation (Supplementary data, Table S1), with exception ([6OSO3]GlcNAc $\beta$ 1-3Gal  $\beta$ 1-4GlcNAc- $\beta$ -Sp0) which appeared to be one of the highest bound glycans on the array (Figure 1; Table I). The binding specificity displayed by galatrox on the glycan microarray is very similar to the binding specificities of exhibited by many members of the galectin family, which also tend to display a higher preference of extended non-sialylated terminal LacNAc glycans, LacNAc or lactose in the context of blood group antigens and sulfated LacNAc or lactose containing glycans. Given the similarities in binding specificity, these results strongly suggest that galatrox may possess the ability to bind similar cell mammalian glycans as engaged by galectins and therefore directly impact the behavior of cells involved in inflammation.

### *Galatrox binds to laminin “in vitro”*

Although the glycan microarray provides an unprecedented platform to elucidate the binding specificity of GBPs toward defined glycans, we next sought to determine whether galatrox might display high binding toward a previously characterized mammalian glycoprotein ligand. While previous studies suggest that galectins likely mediate leukocyte regulation through engagement of cell surface LacNAc-containing glycans, they appear to do so through formation of complex galectin-mediated lattice formation that may result in the recruitment of many different glycoproteins (Brewer et al. 2002; Belardi et al. 2012). In contrast, early studies demonstrated that more defined glycoproteins, such as laminin, may also serve as glycoprotein ligands for galectins (van den Brùle et al. 2003; Camby et al. 2006; Friedrichs et al. 2007). As a result, we next sought to determine whether galatrox might also display high affinity for laminin, which exhibits a large number of *N*-glycans composed of poly-LacNAc<sub>*n*</sub> residues containing  $\beta$ -Gal and  $\alpha$ -Gal terminal residues, and forms the major component of the extracellular matrix (ECM; Fujiwara et al. 1988; Jin et al. 1995). FITC-galatrox displayed significant binding to murine laminin in a concentration-dependent manner (Figure 2). Alternatively, we have assessed the capacity of galatrox to interact to other ECM component that lacks of terminal LacNAc motif such as fibronectin. Human plasma fibronectin is a glycoprotein that contains  $\alpha$ (2,6)-linked sialic acid at the terminal galactose residues on all seven oligosaccharides (N- and O-linked) and only two have fucosylated glycans (Katnik-Prastowska et al. 2006; Katnik-Prastowska and Orczyk-Pawłowicz 2011). Interestingly, FITC-galatrox did not bind to fibronectin, presenting similar binding pattern as bovine serum albumin (BSA; Figure 2). These results suggest that galatrox interaction to laminin is specific and mediated by recognition of LacNAc-containing glycans on this component of the ECM.



**Fig. 1.** Galatrox glycan microarray analysis. FITC-galatrox, at concentration of 0.1  $\mu\text{g/mL}$ , was assessed for glycan recognition pattern using mammalian printed array version 3.2 which consists of 406 glycans in replicates of 6. The results are expressed by the mean RFU and SEM and top 15 ranked glycan structures are illustrated with its respective glycan number.

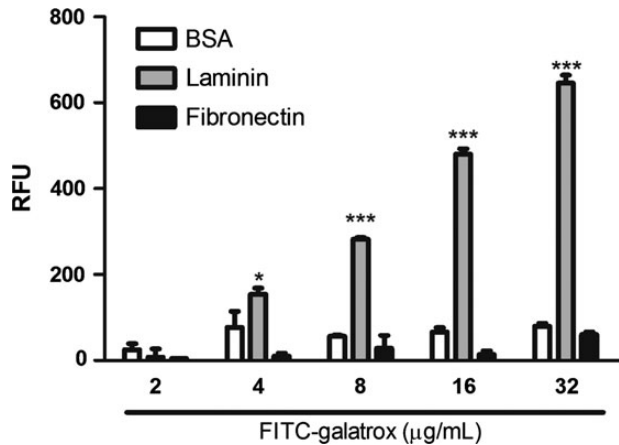
**Table I.** Top 15 rank of galatrox glycan array recognition

Glycan no.	Glycan structure	Rank <sup>a</sup>
293	Gal $\beta$ 1-4GlcNAc $\beta$ 1-3(Gal $\beta$ 1-4GlcNAc $\beta$ 1-6)Gal $\beta$ 1-4GlcNAc-Sp0	100
22	Gal $\beta$ 1-4GlcNAc $\beta$ 1-3(Gal $\beta$ 1-4GlcNAc $\beta$ 1-6)GalNAc $\alpha$ -Sp8	99.7
384	Gal $\beta$ 1-4GlcNAc $\beta$ 1-2(Gal $\beta$ 1-4GlcNAc $\beta$ 1-4)Man $\alpha$ 1-3(Gal $\beta$ 1-4GlcNAc $\beta$ 1-2)(Gal $\beta$ 1-4GlcNAc $\beta$ 1-6)Man $\alpha$ 1-6) Man $\beta$ 1-4GlcNAc $\beta$ 1-4GlcNAc $\beta$ -Sp21	95.6
51	Gal $\beta$ 1-4GlcNAc $\beta$ 1-2Man $\alpha$ 1-3(Gal $\beta$ 1-4GlcNAc $\beta$ 1-2Man $\alpha$ 1-6)Man $\beta$ 1-4GlcNAc $\beta$ 1-4GlcNAc $\beta$ -Sp13	91.8
146	Gal $\beta$ 1-4GlcNAc $\beta$ 1-3Gal $\beta$ 1-4GlcNAc $\beta$ -Sp0	89.7
119	Gal $\beta$ 1-3(Gal $\beta$ 1-4GlcNAc $\beta$ 1-6)GalNAc $\alpha$ -Sp8	89.0
295	Gal $\beta$ 1-4GlcNAc $\alpha$ 1-6Gal $\beta$ 1-4GlcNAc $\beta$ -Sp0	88.7
364	Gal $\beta$ 1-4GlcNAc $\beta$ 1-2(Gal $\beta$ 1-4GlcNAc $\beta$ 1-4)Man $\alpha$ 1-3(Gal $\beta$ 1-4GlcNAc $\beta$ 1-2Man $\alpha$ 1-6) Man $\beta$ 1-4GlcNAc $\beta$ 1-4GlcNAc-Sp21	87.7
404	Gal $\beta$ 1-3GlcNAc $\alpha$ 1-6Gal $\beta$ 1-4GlcNAc $\beta$ -Sp0	85.5
144	Gal $\beta$ 1-4GlcNAc $\beta$ 1-3Gal $\beta$ 1-4(Fuc $\alpha$ 1-3)GlcNAc $\beta$ 1-3Gal $\beta$ 1-4(Fuc $\alpha$ 1-3)GlcNAc $\beta$ -Sp0	84.5
148	Gal $\beta$ 1-4GlcNAc $\beta$ 1-3Gal $\beta$ 1-4Glc $\beta$ -Sp8	84.1
350	[6-OSO <sub>3</sub> ]GlcNAc $\beta$ 1-3Gal $\beta$ 1-4GlcNAc $\beta$ -Sp0	81.7
358	Gal $\alpha$ 1-3Gal $\beta$ 1-4GlcNAc $\beta$ 1-2Man $\alpha$ 1-3(Gal $\alpha$ 1-3Gal $\beta$ 1-4GlcNAc $\beta$ 1-2Man $\alpha$ 1-6)Man $\beta$ 1-4GlcNAc $\beta$ 1-4GlcNAc $\beta$ -Sp20	80.2
344	Gal $\beta$ 1-4GlcNAc $\beta$ 1-2Man $\alpha$ 1-6Man $\beta$ 1-4GlcNAc $\beta$ 1-4GlcNAc-Sp12	78.4
143	Gal $\beta$ 1-4GlcNAc $\beta$ 1-3GalNAc $\alpha$ -Sp8	70.9

<sup>a</sup>The rank values were obtained with the following calculation: rank = 100  $\times$  (RFU bound/highest RFU) (Heimburg-Molinario et al. 2011).

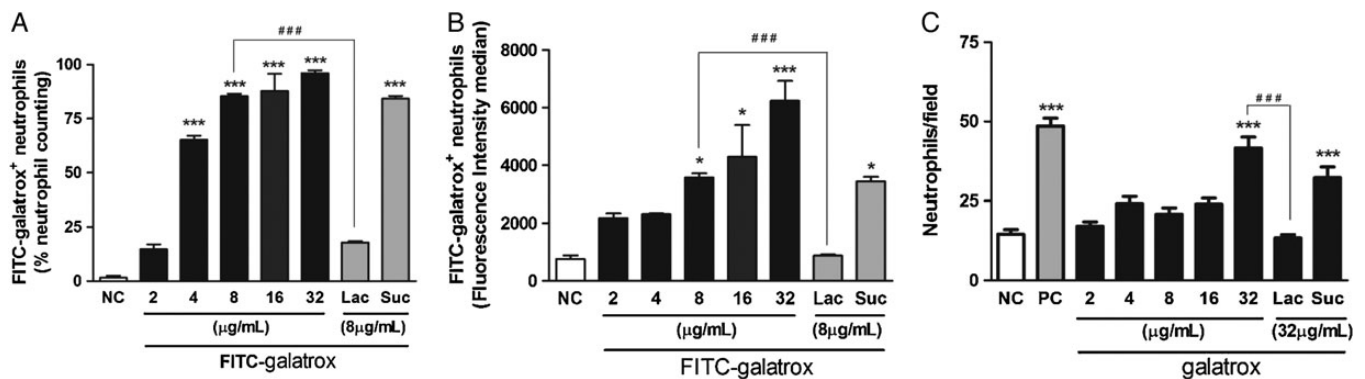
*Galatrox interacts with human neutrophil surfaces in a glycan-dependent manner and induces neutrophil chemotaxis*

Given the remarkable similarities in carbohydrate-binding specificity between galatrox and previously studied members of the galectin family, we next sought to determine whether galatrox might similarly engage leukocytes (Allavena et al. 2004; Camby et al. 2006; Fulcher et al. 2006). We evaluated the ability of



**Fig. 2.** Analysis of galatrox binding to ECM components. Polystyrene microplate wells were coated with ECM components laminin and fibronectin, and BSA as random binding ( $n = 3$  for each component and BSA). After each well was incubated for 1 h with several concentrations of FITC-galatrox (2–32 µg/mL). The analysis was performed by fluorescence reading using a Power Wave X reader on an emission wavelength of 585/20 nm. The results obtained were expressed as mean of RFU  $\pm$  SEM and are representative of two independent experiments. \* $P < 0.05$  and \*\*\* $P < 0.001$  compared with BSA binding.

galatrox to bind to human neutrophils and examined the involvement of the carbohydrate-recognition domain (CRD) in this interaction, as neutrophils represent a primary cellular mediator of inflammation. Similar to laminin binding, FITC-galatrox displayed significant binding to neutrophils over a range of concentrations (Figure 3A and B). Interestingly, preincubation of FITC-galatrox with lactose (20 mM) inhibited binding, while preincubation with sucrose (20 mM), a hapten sugar which is not recognized by galatrox, failed to alter neutrophil recognition, strongly suggesting that galatrox recognizes neutrophils through a glycan-dependent interaction (Figure 3A and B). The ability of mammalian GBPs to bind ECM and cell surface glycans places them in a unique position to regulate cellular migration and trafficking. As galectins represent the only known soluble mammalian GBPs with LacNAc-binding activity, we next sought to determine whether incubation of neutrophils with galatrox may induce similar alterations in neutrophil function as previously observed following galectin engagement. To accomplish, we first examined the potential impact of galatrox on neutrophil chemotaxis, as several previous studies suggested that galectins might regulate this key feature of neutrophil function (Dias-Baruffi et al. 1995a; de Toledo et al. 2007; Elifio-Esposito et al. 2011; Auvynet et al. 2013). To determine whether galatrox promotes neutrophil migration, we examined potential galatrox-induced chemotaxis through polycarbonate membrane pores using a 48-well microchamber. Galatrox induced significant migration of human neutrophils, with the most significant alterations in leukocyte trafficking occurring following the addition of 32 µg/mL galatrox. Similarly to neutrophil recognition (Figure 3A and B), preincubation of cells with lactose (20 mM) reduced their migration to similar migration levels as observed in the negative control (NC; Figure 3C). However, preincubation with sucrose (20 mM) failed to alter galatrox-induced neutrophil migration (Figure 3C).



**Fig. 3.** Galatrox human neutrophil surface interaction and chemotaxis. Human neutrophils ( $1 \times 10^6$  cells) were incubated with different concentrations of galatrox-FITC (2–32 µg/mL) for 30 min at 4°C. The CRD involvement was assessed by previous incubation of FITC-galatrox samples (8 µg/mL) with  $\alpha$ -lactose or  $\alpha$ -sucrose (20 mM). Cells treated with HBSS only were used as NC. After incubation, the suspension was washed with PBS and cells analyzed by flow cytometry. The treatment was performed in triplicate for each condition and the results were reported as (A) percentage  $\pm$  SEM and (B) fluorescence intensity median  $\pm$  SEM of galatrox-FITC<sup>+</sup> neutrophil population from two independent experiments. \* $P < 0.05$  and \*\*\* $P < 0.001$  compared with NC and #### $P < 0.001$  when compared with galatrox 8 µg/mL preincubated with  $\alpha$ -lactose 20 mM. (C) Human neutrophil chemotactic activity in vitro was performed using a 48-well Boyden microchamber with galatrox concentration ranging from 2 to 32 µg/mL. Also, galatrox (32 µg/mL), preincubated with 20 mM of  $\alpha$ -lactose (Lac) or  $\alpha$ -sucrose (Suc), was assessed for CRD involvement. NC, represented by the random migration, and fMLP ( $10^{-7}$  M) were used as the reference chemoattractant for PC. The cells that migrated from the upper chamber to the lower through the membrane were counted and recorded as the mean  $\pm$  SEM of total neutrophil per field (C). The assay was performed in triplicate for each condition, in two independent experiments. \*\*\* $P < 0.001$  compared with NC. #### $P < 0.001$  when comparing Galatrox 32 µg/mL which was pre-incubated or not with  $\alpha$ -lactose 20 mM.

*Galatrox induces in vivo neutrophil migration and the release of inflammatory cytokines*

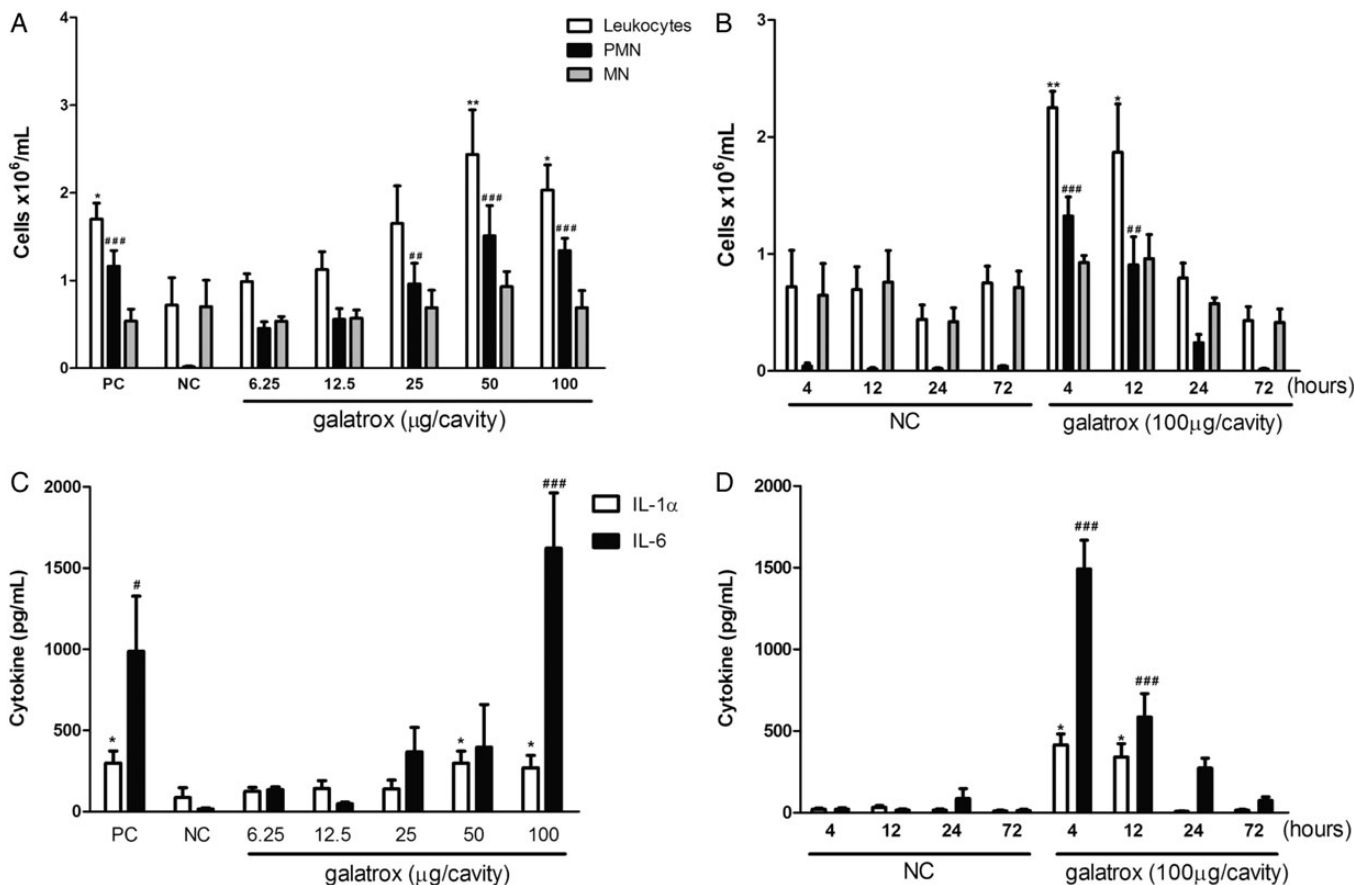
Since galatrox induced neutrophil migration in vitro and is released into tissues following envenomation, we explored whether galatrox could induce neutrophil migration in vivo. Injection of galatrox induced significant leukocyte migration into the peritoneal cavity 4 h postadministration in a concentration-dependent manner with neutrophils representing the dominant leukocyte population (Figure 4A and B). In contrast, mononuclear cell migration profile did not change during all treatments, with recruitment values remaining statistically equivalent to the NC (Figure 4A). A progressive reduction of cell infiltrate was observed following 12 h. Significant levels of leukocytes/neutrophils continued to be observed up to 12 h following administration, following which a decline in neutrophil numbers occurred, eventually reaching similar values as the NC from 24 up to 72 h following injection (Figure 4B).

In addition to examining the cellular infiltrate following galatrox injection, we also determine whether alterations in cytokines levels might accompany leukocyte recruitment. Galatrox

injection induced significant levels of interleukin (IL)-1 $\alpha$  and IL-6 in a concentration- and time-dependent manner (Figure 4C and D). In contrast, no different in the levels of tumor necrosis factor (TNF)- $\alpha$  and IL-10 were observed in comparison with NC (data not shown). Taken together, these results suggest that galatrox not only possesses the capacity to engage similar ligands as endogenous mammalian galectin GBPs but also appears to induce similar changes in neutrophil function an overall cytokine secretion of proinflammatory cytokines. Thus, galatrox may in part mediate the proinflammatory activity of *B. atrox* envenomation by engaging similar pathways normally utilized by endogenous GBPs (Escocard Rde et al. 2006; Stowell, Qian, et al. 2008; Moreira et al. 2012).

*Galatrox binds to glycoconjugates on BMDM surface and induces proinflammatory cytokine production mediated by Toll-like receptor 4*

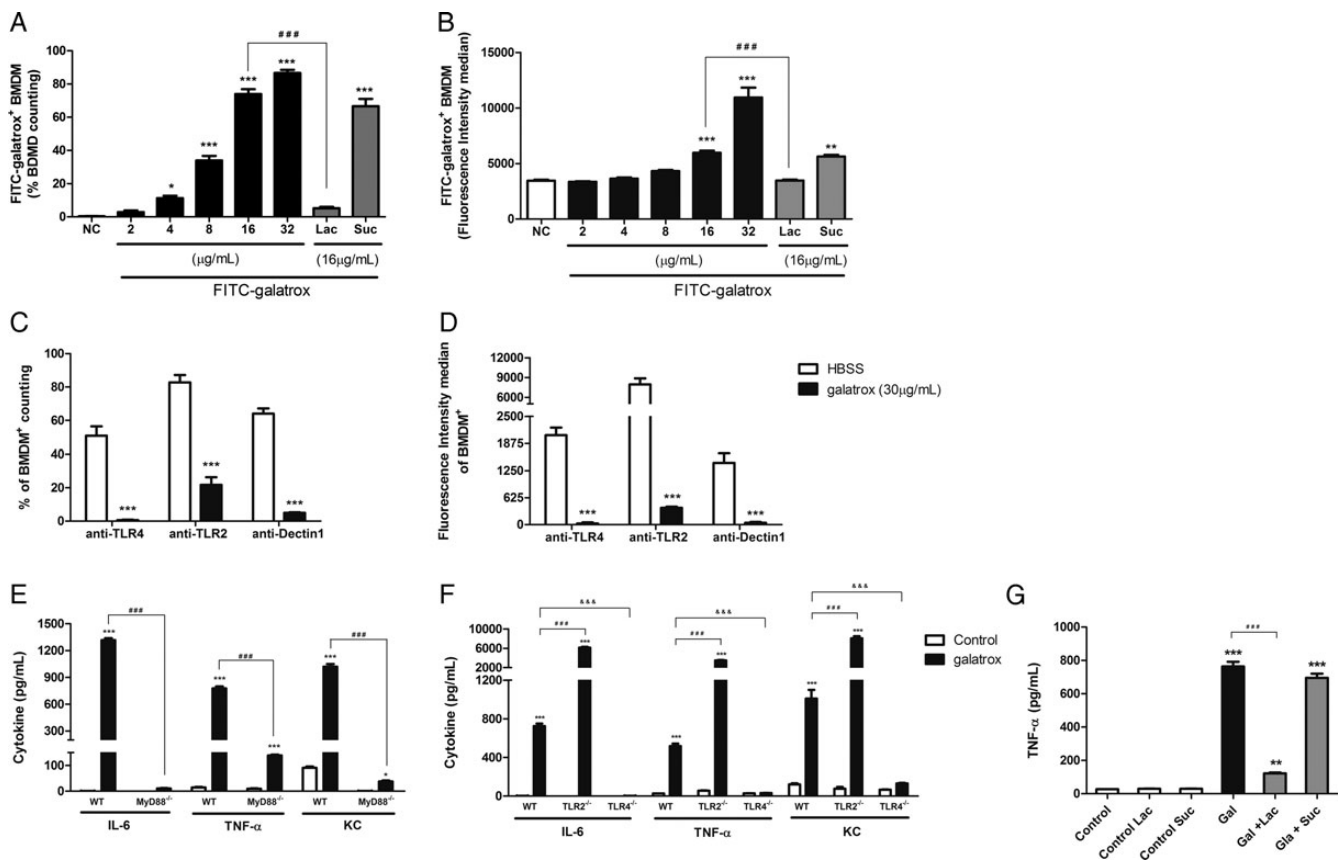
In addition to neutrophils, macrophages can significantly impact inflammatory changes following injury or infection and are likewise mediators of the proinflammatory activity of many galectin



**Fig. 4.** Mice peritoneal leukocyte migration. In vivo leukocyte migration was performed by administration of galatrox (6.25–100  $\mu\text{g/cavity}$ ) into male BALB/c mice peritoneum, and the cavity washed after 4 h administration. NC animals received PBS alone, and PC group carrageenan (500  $\mu\text{g/cavity}$ ). For time-course evaluation, mice were treated with galatrox (100  $\mu\text{g/cavity}$ ) and leukocyte count was evaluated 4–72 h after administration. The peritoneal wash from dose (A) and the time-course (B) assay were assessed for total and differential cell counting. The results were expressed by the mean  $\pm$  SEM of leukocyte, polymorphonuclear neutrophils (PMN) and mononuclear cell (MN) concentration (cells/mL). Peritoneal wash cytokine levels from dose (C) and time-course (D) in vivo leukocyte migration were evaluated by ELISA. Each treatment group consisted of five animals ( $n = 5$ ) in two independent experiments. In cell counting (A and B), \* $P < 0.05$  and \*\* $P < 0.01$  when total leukocytes groups compared with respective control. ### $P < 0.01$  and #### $P < 0.001$  when PMN groups compared with respective control. For cytokine level detection (C and D), \* $P < 0.05$  for IL-1 $\alpha$  compared with respective control, # $P < 0.05$ , and #### $P < 0.001$  for IL-6 compared with respective control.

family members (Garcia-Ramallo et al. 2002; Gordon and Taylor 2005). As a result, we next determine whether galatrox may similarly engage macrophages cell surface carbohydrate receptors and induce alterations in cytokine secretion. To accomplish this, we prepared a differentiated bone marrow-derived macrophage (BMDM) cell culture suspension that was composed of >90% of CD11b<sup>+</sup>F4/80<sup>+</sup> macrophage population (data not shown). As observed for neutrophil surface binding (Figure 2), FITC-galatrox was capable to interact to BMDM surface in a dose-dependent manner (Figure 5A and B). Preincubation of FITC-galatrox with lactose (20 mM) drastically inhibited binding, but not with sucrose (20 mM), suggesting that galatrox recognition of glyco-conjugate(s) on BMDM surface (Figure 5A and B), similar to previous studies examining galectin function.

Cell surface glycosylation can impact the function of many cell surface receptors. For example, the activity of several cell surface receptors, such as Toll-like receptors (TLRs) family members and Dectin-1 receptor, can be modulated following engagement by members of the galectin family and others lectins to induce proinflammatory cytokine secretion (da Silva Correia and Ulevitch 2002; Amith et al. 2010; Esteban et al. 2011; Galván-Moroyocqui et al. 2011). In order to identify possible galatrox glycan targets on the macrophages surface, we pre-incubated BMDM with galatrox to determine whether galatrox possessed the capacity to inhibit interactions with anti-TLR-4, anti-TLR-2 and anti-Dectin1 (Figure 5C and D). Preincubation with galatrox significantly inhibited anti-TLR-4, anti-TLR-2 and anti-Dectin1 binding, strongly suggesting that



**Fig. 5.** BMDM-galatrox binding and cell culture inflammatory cytokine release. BMDM ( $1 \times 10^6$  cells/mL) was incubated with different concentrations of FITC-galatrox (2–32  $\mu$ g/mL) in the presence of  $\alpha$ -lactose or  $\alpha$ -sucrose (galatrox 16  $\mu$ g/mL + 20 mM of respective sugar). Cells treated with HBSS only were used as NC (A and B). BMDM treated previously with non-conjugated galatrox (30  $\mu$ g/mL) or HBSS only was incubated with conjugated anti-TLR-4 (PE), anti-TLR-2 (PE-Cy7) and anti-Dectin 1 (Alexa Fluor 647) (C and D). After incubation, the suspensions were washed with PBS and cells analyzed by flow cytometry. The results were reported as percentage  $\pm$  SEM of BMDM<sup>+</sup> (A and C) and fluorescence intensity median  $\pm$  SEM BMDM<sup>+</sup> (B and D) from two independent experiments. \* $P$  < 0.05, \*\* $P$  < 0.01 and \*\*\* $P$  < 0.001 compared with control and \*\*\*\* $P$  < 0.001 when compared with galatrox 16  $\mu$ g/mL preincubated with  $\alpha$ -lactose 20 mM. Cytokines KC, IL-6 and TNF- $\alpha$  concentrations were determined from BMDM cell suspension isolated from wt, *tlr4*<sup>-/-</sup>, *tlr2*<sup>-/-</sup> and *myd88*<sup>-/-</sup> animals and treated with galatrox 30  $\mu$ g/mL for 24 h. Control represents cells treated with medium only. CBA system kit assay was used to detect IL-6 and TNF- $\alpha$  and ELISA for detection of KC production in the supernatants. Alternatively, BMDM cells isolated from wt mice were treated galatrox (30  $\mu$ g/mL) preincubated or not with  $\alpha$ -lactose (20 mM) or sucrose (20 mM), for 24 h. Culture medium supplemented or not with carbohydrates were used as controls. Cell supernatants were used to determine the TNF- $\alpha$  level by ELISA (G). The results were reported as mean  $\pm$  SEM of cytokine concentration (pg/mL) of BMDM supernatant treated with galatrox ( $n$  = 7). \*\*\* $P$  < 0.001 when compared with Control group and \*\*\*\* $P$  < 0.001 or &&&& $P$  < 0.001 when comparing wt BMDM against knockout groups and \*\*\* $P$  < 0.001 when compared Galatrox groups with Control groups and \*\*\*\* $P$  < 0.001 when compared Galatrox sample with Galatrox sample plus  $\alpha$ -lactose.

galatrox may interact with BMDM cell surface TLR-4, TLR-2 and Dectin-1 ligands.

Although the ability of galatrox to inhibit anti-TLR-4 and anti-TLR-2 binding suggests that TLRs may serve as functional ligands, we next sought to determine whether galatrox-induced proinflammatory activity indeed requires TLR signaling. While galatrox induced significant IL-6-, TNF- $\alpha$ - and keratinocyte-derived chemokine (KC) in wt BMDM (Figure 5E), similar incubation with BMDM from MyD88<sup>-/-</sup> deficient mice, which are insensitive to TLR-2 and TLR-4 signaling (Sorgi et al. 2012; Shin et al. 2008), failed to result in similar increases in these same cytokines, strongly suggesting that galatrox mediates its proinflammatory activity through a TLR MyD88-dependent pathway. Although Dectin-1 may serve as a potential ligand for galatrox, the significant inhibition observed following galatrox incubation with MyD88<sup>-/-</sup> BMDM suggests that galatrox signals through a Dectin-1-independent pathway, as Dectin-1 signaling through a MyD88-independent Syk tyrosine kinase-dependent pathway (Saijo et al. 2007; Leal et al. 2010). As a result, we next sought to determine whether galatrox may preferentially signal through TLR-2 and TLR-4. Importantly, TLR-4<sup>-/-</sup> BMDM displayed significantly reduced sensitivity to galatrox-induced cytokine secretion than wt BMDM (Figure 5F). In contrast, galatrox induced increased cytokine secretion from TLR-2<sup>-/-</sup> macrophages when compared with the same wt BMDM (Figure 5F), similar to other TLR-4 agonists (Sorgi et al. 2012). Interestingly, the capacity of galatrox to promote the release of inflammatory cytokines from macrophages, such as TNF- $\alpha$ , is dependent on their ability to recognize glycans on the surface of this phagocyte, since its proinflammatory activity was dramatically inhibited by lactose (Figure 5G). Interestingly, Park et al. (2010) described a lectin named Korean mistletoe lectin-C that interacts with TLR-4 and promotes macrophage activation as the same as demonstrated to galatrox (Figure 5). These results resolutely suggest that TLR-4 likely plays the primary role in mediating galatrox-induced cytokine secretion and support a major participation of TLR-4 as a mediator of galatrox macrophage stimulation by a carbohydrate-dependent mechanism.

## Discussion

Snake venom and other toxins often utilize endogenous cellular receptor systems and signaling pathways to exert their effects. While early studies suggested that several key mediators of snake venom might possess carbohydrate-binding activity, the fine specificity of venom GBPs remained unknown. As elucidation of carbohydrate-binding activity can often provide key insight into potential ligands and function, we utilized recently developed glycan microarrays to evaluate galatrox-binding specificity (Coombs et al. 2006; Stowell, Arthur, Mehta, et al. 2008; Heimburg-Molinaro et al. 2011). The results obtained following glycan microarray analysis provided critical insight into galatrox function and suggest that galatrox may engage similar ligands as endogenous  $\beta$ -galactoside GBPs to exert a proinflammatory response (Stowell, Arthur, Slanina, et al. 2008; Stowell et al. 2009, 2010, 2014; Cerliani et al. 2011).

The results obtained from glycan microarray analysis show that galatrox displays similar glycan interaction characteristics to

others  $\beta$ -galactoside-binding snake lectins, such as BJcuL, from *Bothrops jararacussu* snake venom (Elifio-Esposito et al. 2011) and RSVL from *Crotalus atrox* snake venom (Young et al. 2011) and mammalian lectins such as galectins (Karmakar et al. 2008; Stowell, Arthur, Mehta, et al. 2008; Stowell et al. 2009), SRCL (human scavenger receptor C-type lectin), rat Kupffer cell receptor, rat macrophage galactose lectin and rat hepatocytes asialoglycoprotein receptor lectin (Coombs et al. 2006). The assay also shows that galatrox does not bind to glycans with non-reducing terminal LacNAc modified by sialic acid residues, strongly suggesting that sialylated biological receptors resist galatrox-LacNAc interaction. Similar to galatrox, RSVL interaction to LacNAc glycans is also inhibited by sialylation (Young et al. 2011), as well as the mammalian galectin, galectin-2 (Stowell, Arthur, Mehta, et al. 2008). While other mammalian and snake  $\beta$ -galactoside-binding proteins appear to bind LacNAc following blood group or sialic acid modification (Leppänen et al. 2005; Coombs et al. 2006; Toscano et al. 2007; Stowell, Arthur, Mehta, et al. 2008; Stowell et al. 2009; Young et al. 2011), galatrox binding appeared to be more sensitive to various LacNAc modifications. These results suggest that while galatrox may possess some glycan cell surface receptor overlap with other snake and mammalian  $\beta$ -galactoside containing ligands, galatrox appears to possess a more restricted LacNAc profile and therefore may only engage a subset of the ligands bound by similar GBPs.

The preference for terminal LacNAc glycans by galatrox is most similar to mammal galectin-1 (Stowell, Arthur, Mehta, et al. 2008), the first described galectin family member. Early studies suggested that galectin-1 displays significant affinity laminin, a basement membrane ECM glycoprotein expressing high amounts of LacNAc-containing glycans and also known to participate as a substrate chemoattractant interaction involved in leukocyte migration (Jin et al. 1995; Lauffenburger and Horwitz 1996; Sánchez-Madrid and del Pozo 1999; Camby et al. 2006). Similar to previous studies, galatrox also bound laminin, resolutely suggesting that glycan microarray results accurately predict potential interactions with LacNAc bearing glycoproteins and suggested that galatrox may share similar ligands as members of the galectin family. In contrast, galatrox failed to recognize fibronectin, a heavily sialylated glycoprotein that can also fail to serve as a suitable ligand for galectin family members (Carsons et al. 1987). Given the role of ECM components in galectin-mediated inflammatory regulation (Camby et al. 2006; Elifio-Esposito et al. 2011), galatrox may mediate some of its proinflammatory activities through similar interactions, where GBP gradients cannot only facilitate activation, but play a significant role in leukocyte chemotaxis (Hauzenberger et al. 1995; Lauffenburger and Horwitz 1996; Sánchez-Madrid and del Pozo 1999). Taken together, these results demonstrated that the glycan microarray can accurately predict interactions with intact glycoproteins and that glycan microarray results of toxin may predict potential ligands shared by endogenous mammalian GBPs.

Given the similarities between galatrox- and galectin-binding specificity, we determined whether galatrox may similar engage human neutrophils, as previous studies demonstrated that galectins can induce neutrophil activation. Our results demonstrate that galatrox can induce significant neutrophil activation, similar to mammalian lectins, galectin-3 (Nieminen et al. 2005; Farnworth et al. 2008), galectin-1 (Almkvist et al. 2002;

Dias-Baruffi et al. 2003) and galectin-8 (Nishi et al. 2003; Stowell, Arthur, Slanina, et al. 2008). Although the glycosylation patterns of leukocyte glycoproteins have been studied (Wenzel-Seifert and Seifert 2003; Wang et al. 2004; Sun et al. 2006; Babu et al. 2009), the carbohydrate compositions of most individual neutrophil receptor glycans have not fully been characterized. Indeed, the actual receptors for galectin-mediated neutrophil activation remain to be elucidated despite early studies describing this activity nearly 20 years ago. Although a more clear understanding of the glycosylation pattern of leukocyte receptors will be necessary to better define the biological role of GBPs in neutrophil-mediated immune responses, galatrox may induce lectin lattice formation of multiple counter receptors to signal a response, as has been recently reported for several members of the galectin family (Brewer et al. 2002; Boscher et al. 2011; Earl et al. 2011)

During innate immunity, the role of tissue resident macrophages is pivotal for initiation and resolution of inflammatory response (Davies et al. 2013). Therefore, to better understand the involvement of macrophages on galatrox proinflammatory behavior, we assessed the ability of galatrox to stimulate macrophages. BMDM-binding assays suggested that galatrox may interact with proinflammatory cell surface glycoproteins TLR-2, TLR-4 and Dectin-1. Moreover, galatrox stimulation of macrophages, observed by the expression of IL-6, TNF- $\alpha$  and KC, appears to be predominantly mediated by TLR-4 activation through a MyD88 cell signaling pathway. Recent results suggest that other snake GBPs may signal through similar pathways (Moreira et al. 2013). Protazoan  $\beta$ -galactoside-binding GBPs also appear to engage a similar pathway, suggesting that other exogenous proteins may engage TLR-4 through their CRDs (Galván-Moroyoqui et al. 2011). da Silva Correia and Ulevitch (2002) showed that glycosylation of TLR-4 and MD2, a TLR-4 adapter protein, is necessary for cross-linking and the functional integrity of this receptor. Taken together, our results suggest that galatrox likely interacts with LacNAc-containing glycans on TLR-4 to engage MyD88-dependent proinflammatory cytokine secretion by macrophages and subsequent neutrophil recruitment. As a result, galatrox may belong to the venom-associated molecular pattern, a description of venom-associated molecules that possess the capacity to engage and activate pattern recognition receptors (Zoccal et al. 2014).

The results from the present work suggest that galatrox possesses high selectivity for LacNAc-containing glycans. Engagement of LacNAc cell surface glycans appears to induce neutrophil activation and migration, which may be further enhanced by galatrox-induced macrophage activation and subsequent cytokine secretion. Therefore, investigation into the carbohydrate-binding specificity of galatrox provided significant insight into its potential ligands and function. This understanding will likely enable the development of novel approaches to inhibit the deleterious effects of venomation in patients suffering from a snake bite.

## Materials and methods

### Animals

Male BALB/c and C57BL/6 mice weighting 18–22 g were provided by Isogenic Animal Facility from the School of

Pharmaceutical Sciences of Ribeirão Preto. Male TLR-2, TLR-4 and MyD88 knockout mice (*tlr4*<sup>-/-</sup>, *tlr2*<sup>-/-</sup> and *myd88*<sup>-/-</sup> from C57BL/6 genetic background strain) were provided by the Animal Care Facility at the Ribeirão Preto Medical School, University of Sao Paulo. Animal care procedures were performed according to COBEA (Brazilian College of Animal Experimentation) guidelines and experimental protocols approved by The Committee for Ethics on Animal Use (CEUA) from the Ribeirão Preto Campus Prefecture of the University of São Paulo (Protocol number: 07.1.1495.53.2).

### Peripheral blood sample from human donors

Peripheral blood sample were donated from the Hemotherapy Center of Ribeirão Preto. The samples were collected from both male and female health donors, belonging to the 19- to 40-year-old age group, not undergoing any form of pharmacological therapy. All care procedures were approved by the local Ethical Committee/School of Pharmaceutical Sciences of Ribeirão Preto of the University of São Paulo (protocol number: 118).

### Isolation of galatrox

Galatrox was prepared from *B. atrox* crude venom according to Mendonça-Franqueiro et al. (2011). Briefly, ~150 mg of *B. atrox* whole venom was suspended in phosphate-buffered saline (PBS) buffer and applied into a lactosyl-Sepharose column (10 × 1 cm—Sigma—Aldrich). The column was washed with PBS only to elute the unbound material, and the bound material was eluted with PBS combined with  $\alpha$ -lactose (100 mM). A desalting procedure on a Sephadex G-25 column (8 × 1 cm—Amersham Pharmacia Biotech) was performed to remove lactose from galatrox preparations. The homogeneity of galatrox samples was evaluated by SDS-PAGE and the preservation of lectin activity was confirmed by hemagglutinating assay, as described by Nowak et al. (1976). The protein concentrations were determined using a microassay based on the Bradford dye-binding procedure (Bradford 1976). Before the assays, galatrox samples were treated in polymyxin-B resin to remove remaining endotoxin residues and filtered through 0.22  $\mu$ m pore filters.

### Galatrox glycan array

To determine the glycan recognition pattern of galatrox, the glycan microarray was performed by the Consortium for Functional Glycomics Core H (Emory University School of Medicine, Atlanta, GA, USA). For the analysis, different concentrations of galatrox (0.05, 0.1, 1 and 144  $\mu$ g/mL) were assessed using mammalian printed array version 3.2 which consists of 406 glycans in replicates of 6. The galatrox–glycan interaction was detected by an FITC-galatrox conjugate. The lectin was conjugated to FITC (Pierce, Rockford, IL, USA) by incubating both protein and fluorophore according to manufacture instructions. In an attempt to remove the excess of unbound FITC from conjugated FITC-galatrox, the incubated sample was submitted to a Sephadex G-25 column. The results obtained from the glycan microarray analysis relate to the analysis of FITC-galatrox at a concentration of 0.1  $\mu$ g/mL, expressed by both the mean relative fluorescence unit (RFU) plus the standard error of the mean (SEM). A ranking of the



ligands was also indicated with the following calculation according to Heimburg-Molinaro et al. (2011):  $\text{rank} = 100 \times [\text{RFU bound/highest RFU}]$ . The full table of glycan recognition patterns is available in Supplementary data, Table S1.

#### Extracellular matrix components binding assay

The binding efficiency of galatrox to ECM components, laminin and fibronectin, was evaluated by solid-phase assay. Solid white polystyrene microplate wells (MaxiSorp FluoroNunc) were coated with murine sarcoma laminin, human plasma fibronectin or BSA (5  $\mu\text{g}/\text{well}$ —Sigma—Aldrich) in PBS overnight at 4°C. After incubation, the plate was washed three times with PBS-Tween 20 0.05%, and then blocked with PBS-BSA 1% for 1 h at 37°C. Each well was incubated for 1 h with several concentrations of galatrox-FITC (2–32  $\mu\text{g}/\text{mL}$ ) in a final volume of 50  $\mu\text{L}$ . After washing procedure, the analysis was performed by fluorescence reading using a Power Wave X reader (Bio-Tek Instruments, Inc.) on a emission wavelength of 585/20 nm. The assay was performed in triplicate of each condition. The results were expressed as mean  $\pm$  SEM of the RFU.

#### Neutrophil surface binding assay

Human neutrophils ( $1 \times 10^6$  cells), obtained according to the *Human neutrophil migration* “in vitro” section, were incubated with different concentrations of galatrox-FITC (2–32  $\mu\text{g}/\text{mL}$ ) for 30 min at 4°C. Aiming to evaluate the role of the lectin CRD on cell surface receptor recognition, galatrox-FITC samples (8  $\mu\text{g}/\text{mL}$ ) were previously incubated with  $\alpha$ -lactose or  $\alpha$ -sucrose (20 mM) for 30 min at 4°C. Cells treated with Hank’s balanced salt solution (HBSS) only were used as a NC. After incubation, the suspension was washed twice with PBS (200  $\times g$  for 10 min), and resuspended to a final volume of 0.5 mL. The samples were analyzed by flow cytometry in a FACSCalibur (BD Biosciences). The results were expressed as the percentage of FITC-galatrox<sup>+</sup> population and fluorescence intensity median of the FITC-galatrox<sup>+</sup> population.

#### Human neutrophil chemotaxis

Chemotactic activity in vitro was performed according to Bignold et al. (1988) using a 48-well acrylic Boyden microchamber (Neuroprobe). The microchamber consisted of two distinct compartments separated by a polycarbonate filter membrane (Millipore) with 5  $\mu\text{m}$  diameter pores.

In order to obtain human neutrophils, heparinized human peripheral blood was layered on a density gradient of a neutrophil isolation medium Monopoly Resolving Medium (density: 1.114—ICN Pharmaceuticals) as described by manufacture instructions. A final neutrophil suspension in HBSS was obtained at a concentration of  $1 \times 10^6$  neutrophils/mL (~94% of polymorphonuclear cells). Galatrox samples in several concentrations (2, 4, 8, 16 and 32  $\mu\text{g}/\text{mL}$ ) were placed in the lower chamber and neutrophil suspension placed in the upper chamber. In order to evaluate the involvement of lectin CRDs in the event, galatrox at its most effective concentration was preincubated with  $\alpha$ -lactose or  $\alpha$ -sucrose (20 mM) at 4°C for 30 min and consequently placed in the lower chamber. Random migration was assessed by using HBSS in the lower chamber. fMLP ( $10^{-7}$  M) was used as the reference chemoattractant. After incubation, the membrane was removed, fixed and stained with Panoptic stain. Cells that migrated through the polycarbonate membrane pore

during the 1 h incubation at 37°C in 5% CO<sub>2</sub> were counted with a light microscope using a 40  $\times$  objective in a total of five fields for each sample.

#### Mice peritoneal cavity leukocyte migration

Male BALB/c mice were injected intraperitoneally with galatrox (6.25–100  $\mu\text{g}/\text{cavity}$ ) in 200  $\mu\text{L}$  of sterile PBS. NC animals received PBS alone, and positive controls (PCs) were given carrageenan (500  $\mu\text{g}/\text{cavity}$ ). After 4 h, animals were anesthetized and the peritoneal cavity was washed with 3 mL PBS supplemented with EDTA (1 mM). For time-course evaluation, mice were treated with galatrox (100  $\mu\text{g}/\text{cavity}$ ) and peritoneal wash was collected 4, 12, 24 and 72 h after administration. The exudate materials from animals were recovered for total and differential cell counting using a Neubauer chamber and Rosenfeld-staining cytopsin preparations, respectively. Each treatment group constituted of five animals. In order to determine the concentrations of inflammatory mediators, IL-1 $\alpha$ , IL-6, IL-10 and TNF- $\alpha$ , peritoneal wash fluid was submitted to centrifugation at 200  $\times g$  for 10 min at 4°C and the supernatant was collected and stored at -80°C for further measurement.

#### Macrophage cell culture

BMDM cell suspension were obtained from pathogen-free C57BL/6 mice (wild type—wt) and *tlr4*<sup>-/-</sup>, *tlr2*<sup>-/-</sup> and *myd88*<sup>-/-</sup>, as described by Marim et al. (2010). Briefly, mice bone marrow from both femurs of three different animals from each strain were removed with RPMI medium supplemented with 10 mM of L-glutamine, 30% L929-cell conditioned medium containing macrophage colony stimulating-factor and 10% FBS. Cells were cultured in BD Optilux polystyrene Petri dishes (4  $\times 10^6$  cells/mL) for 7 days (37°C and 5%CO<sub>2</sub>) in the aforementioned medium, with subsequent medium addition after 4 days. The primary BMDM obtained were then detached from the dishes, washed with sterile PBS and treated with galatrox (30  $\mu\text{g}/\text{mL}$ ) for 24 h, with or without  $\alpha$ -lactose (20 mM), in the same incubation conditions as described above. NC represents cells not treated with galatrox or galatrox plus sucrose (20 mM). After incubation, cell suspensions were centrifuged at 200  $\times g$  for 10 min at 4°C and the supernatant collected and stored at -80°C for inflammatory mediators (IL-2, IL-4, IL-6, IL-10, IL-17a, TNF- $\alpha$ , IFN- $\gamma$  and KC).

#### Determination of inflammatory mediator concentrations

*Cytokines detection.* Enzyme-linked immunosorbent assay. The material collected from peritoneal cavity was assessed for inflammatory cytokine release using commercially available enzyme-linked immunosorbent assay (ELISA) kits to measure IL-1 $\alpha$ , IL-6, IL-10 and TNF- $\alpha$  according to the manufacturer’s instructions (BD Bioscience). KC detection was performed for samples of macrophage cell culture material using the same technique as above. The detection sensitivities were >10 pg/mL.

#### Flow cytometry cytokine bead array analysis

Levels of cytokines released into the cell medium supernatant of macrophage culture treated with galatrox were determined using the TH1/TH2/TH17 Cytometric Bead Array (CBA) Kit

(BD Biosciences) according to the manufacturer's instructions. Briefly, cell culture supernatants were incubated with the anti-cytokine beads and then added to the samples and incubated with the included PE-detection reagent. Beads and detection reagent were also mixed with a serially diluted standard of all cytokines (IL-2, IL-4, IL-6, IL-10, IL-17a, TNF- $\alpha$  and IFN- $\gamma$ ) provided by the manufacturer. Samples and standards were incubated in the dark at room temperature for 3 h. The samples were submitted to flow cytometry (FACSCalibur—BD Biosciences) and the events were analyzed using CBA analysis software (BD Biosciences) applying the five-parameter logistic fit for cytokine standard curve.

#### Statistical analysis

Results were expressed as mean  $\pm$  SEM employing one-way ANOVA, followed by Turkey test. Differences with  $P < 0.05$  were considered statistically significant.

#### Supplementary data

Supplementary data for this article are available online at <http://glycob.oxfordjournals.org/>.

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#### Abbreviations

BMDM, bone marrow-derived macrophages; BSA, bovine serum albumin; CRD, carbohydrate-recognition domain; ECM, extracellular matrix; FITC, fluorescein isothiocyanate; Fuc, fucose; Gal, galactose; GalNAc, *N*-acetylgalactosamine; GlcNAc, *N*-acetylglucosamine; HBSS, Hank's balanced salt solution; IL, interleukin; KC, keratinocyte-derived chemokine; LacNAc, *N*-acetylglucosamine; NeuAc, 5-*N*-acetylneuraminic acid; PBS, phosphate-buffered saline; SDS-PAGE, sodium dodecyl sulfate-polyacrylamide gel electrophoresis; TLRs, Toll-like receptors; TNF, tumor necrosis factor.

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