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## Anti-inflammatory action of seed extract and polymeric nanoparticles of *Syzygium cumini* in diabetic rats infected with *Candida albicans*

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## ABSTRACT

The role of *Syzygium cumini* against chronic complications of Diabetes mellitus (DM), such as fungal infection and inflammation, has been poorly explored. Here, we evaluated the treatment with *S. cumini* aqueous seed extract (ASc, 100mg/Kg) and polymeric nanoparticles containing ASc (NPASc, 100mg/Kg) in diabetic rats infected or not by *Candida albicans* (CA). Male Wistar rats were divided in: control; DM; CA; CA+ASc; CA+NPASc; DM+CA; DM+CA+ASc; and DM+CA+NPASc. Rats were daily treated for 21 days, when glycemic profile, ectonucleotidase (NTPDase and 5'-NT), adenosine deaminase (ADA), acetylcholinesterase (AChE) and dipeptidyl peptidase IV (DPP-IV) activities and nitric oxide (NO) and cytokine levels were analyzed in serum, platelets, lymphocytes and tissues. The results showed that NTPDase, 5'-NT and ADA activities and NO, IL-1, IL-6, TNF- $\alpha$  and IFN- $\gamma$  levels were increased in *C. albicans*, DM and DM+CA. The treatment with ASc and NPASc decreased ectonucleotidase and AChE activities and NO levels. Both treatments also prevented the increase in ADA activity and pro-inflammatory cytokines in cells and serum. In liver and pancreas, NPASc decreased NO levels more efficiently than ASc. The modulation of ectoenzyme activities can be one of the mechanisms by which *S. cumini* act on cytokines that affect the development of chronic complications in DM.

## **INTRODUCTION**

Diabetes *mellitus* (DM), a disorder characterized by chronic hyperglycemia and impaired insulin signaling, generates metabolic changes and an inflammatory status that will eventually affect all body tissues (Boteanu *et al.*, 2015). Several different immune deficits have been described in diabetic

patients, given that cell-mediated immunity seems to be particularly affected, including dysfunctions of white blood cells such as impaired leukocyte adherence, chemotaxis, oxidative burst and bactericidal activity (Fraga-Silva *et al.*, 2015). The cholinergic anti-inflammatory pathway mediated by acetylcholine (ACh) regulates immune responses to pathogens (Borovikova *et al.*, 2000), but the role of this pathway in infections is not well characterized. Moreover, pro-inflammatory mediators such as nitric oxide (NO) play an important role mediating various cellular signaling pathways and inducing pro-inflammatory and destructive effects, exerting antimicrobial action (Shahani and Sawa, 2011; Samarghandian *et al.*, 2013).

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The association between diabetes and the prevalence of microbial infections such as those caused by fungi has also been reported (Joshi *et al.*, 1999).

Disturbances in cell metabolism caused by DM can lead to changes in intra- and extracellular concentrations of adenine nucleotides and nucleosides. The extracellular metabolism of ATP, which is released in response to cellular stress or damage, to adenosine is usually mediated by a variety of enzymes such as triphosphate diphosphohydrolase (NTPDase), which also hydrolyzes ADP to adenosine.

Additionally, following hydrolysis of ATP to AMP, AMP is dephosphorylated by 5'-nucleotidase (5'-NT) to produce adenosine, which has anti-inflammatory and immunosuppressive functions (Schmatz *et al.*, 2013). The extracellular concentrations of adenosine are regulated through the enzyme adenosine deaminase (ADA), which can also interact with membrane proteins such as dipeptidyl peptidase IV (CD26/DPP-IV) (Gorrell *et al.*, 2001). This process is directly involved in T-cell activation (Kameoka *et al.*, 1993).

Syzygium cumini L. Skeels, a traditional medicinal tropical plant of Myrtaceae family widely known as jambolan or hypoglycemic, anti-inflammatory jamun, possesses and antioxidant properties (for review, see Ayyanar et al., 2013). In Brazil, the most used parts of the plant are fruits, which are consumed fresh, and leaves (Vizzotto, 2008), whose infusion is used by the Southern population to treat DM and by quilombo communities to treat kidney, heart and hyperlipidemia (Da Silva et al., 2012). S. cumini is included in the List of Medicinal Plants of Interest for the Public Health System (Renisus) issued by the Brazilian Ministry of Health, which includes plants considered as potentially valuable for the generation of herbal medicines (Brasil, 2009).

Phenols and flavonoids present in high amounts in *S. cumini* are responsible for the several biological activities observed *in vitro*. However, these compounds do not have the same profile of activity *in vivo*. In this sense, polymeric nanoparticles offer a non-toxic and efficient carrier system for enhanced drug bioavailability within the cells, tissues, or both, besides the minimization of the degradation process (Samadder *et al.*, 2012; Bonifácio *et al.*, 2014).

Recently, our group demonstrated that polimeric nanoparticles containing an aqueous extract of *S. cumini* (NPASc) were able to maintain the antioxidant properties of the extract (ASc) and lack toxicity in *Artemia salina* and in rats, highlighting their potential in the treatment of DM and its complications (Bitencourt *et al.*, 2016).

Previous studies from our group have investigated the effects *S. cumini* and adenosine in DM (De Bona *et al.*, 2014; Bitencourt *et al.*, 2015). Nonetheless, as there is paucity of data evaluating the effects of ASc and NPASc on the chronic complications of DM, including increased susceptibility to fungal infections, the present study was aimed: 1) to evaluate changes in cholinergic and purinergic systems in DM rats infected or not by

*Candida albicans*; 2) to evaluate whether ASc and NPASc have potential as anti-inflammatory and antifungal candidates.

## MATERIALS AND METHODS

## Chemicals

Ethyl acetate, methanol, acetonitrile and acetic, gallic, chlorogenic, caffeic and ellagic acids were purchased from Merck (Darmstadt, Germany). Polysorbate 80 (Tween 80<sup>®</sup>), poly-ɛ-caprolactone (PCL, 85 g/mol), sorbitan monooleate (Span 80<sup>®</sup>), adenosine, ATP, ADP, streptozotocin, catechin, epicatechin, quercetin, isoquercitrin, quercitrin, kaempferol and rutin reference standards were acquired from Sigma Chemical Co. (St. Louis, MO, USA). All other chemicals were of analytical grade and were obtained from standard commercial suppliers.

# ASc and NPASc preparation, characterization and phytochemical analysis

*S. cumini* seeds were collected (29°43'22"S and 53°43'47"W, Santa Maria, Rio Grande do Sul, Brazil) fresh, locally and they were cleaned, dried and powdered. They were identified by the Laboratory of Botanic and Pharmacognosy of the Federal University of Santa Maria and a voucher specimen (SMDB 14.001) was deposited in the Herbarium of the institution. ASc was prepared according to Prince *et al.* (1998). NPASc were prepared by the emulsification/evaporation solvent method described by Quintanar-Guerrero *et al.* (1998), with modifications according to Bitencourt *et al.* (2016).

The presence of 11 antioxidant compounds in ASc and NPASc, namely gallic, chlorogenic, caffeic and ellagic acids and catechin, epicatechin, quercetin, quercitrin, isoquercitrin, kaempferol and rutin was investigated by HPLC-DAD. Identification of these compounds was performed by comparing their retention time and UV absorption spectrum with those of the commercial standards. Reverse phase chromatography analyses were carried out under gradient conditions using a C<sub>18</sub> column (4.6 mm x 150 mm, 5 µm). The chromatography peaks were confirmed by comparing their retention time with those of reference standards and by DAD spectra (200 to 500 nm). All chromatographic operations were carried out at ambient temperature and in triplicate. Calibration curves for each substance were: gallic acid, Y = 13973x + 1095.6 (r = 0.9993); catechin, Y = 11840x + 1178.2 (r = 0.9998); epicatechin, Y = 12542x + 1412.7(r = 0.9991); chlorogenic acid, Y = 11864x + 1252.8 (r = 0.9994); caffeic acid, Y = 13178x + 1267.2 (r = 0.9999); ellagic acid, Y = 12681x + 1164.9 (r = 0.9998); rutin, Y = 13077x + 1265.4 (r = (0.9992); isoquercitrin, Y= (11927x + 1306.2) (r = (0.9996); quercitrin, Y = 13470x + 1293.7 (r = 0.9994) and quercetin, Y = 12693x + 1176.0 (r = 0.9997). The limit of detection (LOD) and limit of quantification (LOQ) were calculated based on the standard deviation of the responses and the slope using three independent analytical curves. LOD and LOO were calculated as 3.3 and 10  $\sigma/S$ , respectively, where  $\sigma$  is the standard deviation of

the response and S is the slope of the calibration curve. For the determination of encapsulation efficiency (EE), free phenolic compounds were separated from the nanostructures by ultrafiltration (Bitencourt *et al.*, 2016). The difference between the total and the free concentrations of compounds, determined in the nanostructures and in the ultrafiltrate, respectively, was calculated according to the equation: EE = Total content - Free content / Total content x 100.

## Animals

Male albino Wistar rats (weighing 150–200 g) were housed in colony cages (six rats per cage). All animal experiments were conducted in accordance with principles for laboratory animal use and care, as described in the guidelines of the Ethics Committee for Animal Research of the Federal University of Santa Maria, which approved the experimental protocol ( $n^{\circ}$  074/2014). All efforts were made to minimize the number of animals used and their suffering.

## Induction of DM, animal infection and experimental design

DM type I was induced by the intraperitoneal administration of a single dose of streptozocin (STZ) (60 mg/kg) dissolved in freshly prepared 0.1 M citrate buffer, pH 4.5. Control animals received only citrate buffer. Diabetic rats were allowed to drink 5% glucose solution overnight to overcome the drug-induced hypoglycemia and only diabetic rats with a fasting blood glucose level of at least 250 mg/dl were included in the experiment.

Inoculum was obtained from a clinical *C. albicans* strain grown for 20 h at 30 °C. Yeast cells were harvested by centrifugation at 3000 rpm for 10 min, washed three times with sterile PBS, counted in a hemocytometer and resuspended in sterile PBS to the required inoculum concentration ( $10^5$  colony forming units (CFU)/mL). Fifteen days after DM induction, rats assigned to *C. albicans* infection were inoculated intraperitoneally with 0.2 ml of yeast suspension (Fisher *et al.*, 1989). This inoculum consistently infects >90% of normal rats in this model.

ASc or NPASc were administered by gavage at a dose of 100 mg/kg (Bitencourt *et al.*, 2015). Rats were divided into eight groups of six animals each, according to the following schematic timeline of the experimental design:



At day 21 of treatment, the rats were anesthetized with isoflurane, euthanized and the blood was collected by cardiac puncture. Samples of liver and kidney were rapidly dissected, weighed and placed on ice. Moreover, kidney and liver fragments were aseptically removed and 0.1 g of each organ was homogenized in sterile saline. Samples were cultured onto Sabouraud dextrose agar, incubated at 30 °C for 72 h, and the number of yeasts was expressed as CFU/g of tissue.

## **Platelet preparation**

Platelet-rich plasma (PRP) was prepared by the method of Lunkes *et al.* (2003) and the protein concentration was measured by the method of Lowry *et al.* (1951), using bovine serum albumin as the standard.

## Lymphocyte isolation

Rat lymphocytes were isolated from the spleen under aseptic conditions, as described by Sai Ram *et al.* (1997). The final cell suspension was resuspended in PBS (pH 7.4) and 3 x  $10^6$  cells/mL were used for each analysis. The protein concentration was measured by the method of Lowry *et al.* (1951), using bovine serum albumin as the standard. Acetylcholinesterase (AChE) was determined in lymphocyte suspension by a modification of the spectrophotometric method of Ellman *et al.* (1961). The enzymatic activity was expressed in µmol AcSCh/h/mg of protein.

## **Biochemical and inflammatory analyses**

Serum obtained by centrifugation was analyzed spectrophotometrically for glucose and fructosamine levels using commercial diagnostic kits (Labtest Diagnóstica, Brazil). Plasmatic insulin was measured by radioimmunoassay (DPC Coata-Count kit, CA, USA) in a gamma counter. Serum inflammatory cytokine quantification was assessed by ELISA using commercial kits for interferon (IFN)- $\gamma$ , tumor necrosis factor (TNF)- $\alpha$ , interleukin (IL)-1, IL-6 and IL-10 (R&D Systems, MN, USA), according to manufacturer's instructions.

## **DPP-IV** activity

DPP-IV activity was measured spectrophotometrically in serum and lymphocyte suspension by the method of Jarmolowska *et al.* (2007) and Schön *et al.* (1984), respectively. Samples were incubated with Gly-Pro p-nitroanilide p-toluenesulfonate for 120 min at 37 °C. The reaction was stopped by adding 1 M acetate buffer (pH 4.5). P-nitroanilide was used as standard and the values were expressed in U/l.

## NTPDase and 5'-NT activities

The reaction medium used to assay ATP, ADP and AMP hydrolysis in rat platelets and serum was prepared as described by Lunkes *et al.* (2003) and Oses *et al.* (2004), respectively, and the inorganic phosphate (Pi) released was measured as previously described (Chan *et al.*, 1986). Enzyme activities were expressed as nmol Pi released/min/mg of protein.

#### **ADA** activity

ADA activity in serum, platelets, lymphocytes and tissues (homogenized in 50 mM PBS, pH 7) was estimated spectrophotometrically as previously described (Giusti and Galanti, 1984). The results were expressed as U/L in serum and U/L/mg of protein in lymphocytes and tissues. The protein concentration was measured by the method of Lowry *et al.* (1951).

## Oxide nitric (NO) measurement

NO was determined indirectly by quantifying serum, lymphocytes and tissues (homogenized in 10 mM Tris–HCl buffer solution, pH 7.4). NO was measured by the modified Griess method using the Cobas Mira<sup>®</sup> automated analyzer (Tatsch *et al.*, 2011). The results were expressed as  $\mu$ M/L.

## Statistical analysis

Data were analyzed by one-way analysis of variance (ANOVA) followed by Duncan's post hock test using Statistica 6.0 software (StatSoft. Inc., USA). The limit of statistical significance was set at p<0.05. The results were expressed as mean  $\pm$  SEM.

## RESULTS

## **Characterization of NPASc**

The chromatographic profiles of NPASc showed that nanoencapsulation did not alter the phenolic composition of ASc (Fig. 1A) and the results of encapsulation efficiency showed that almost 100% of phenolic compounds were entrapped within the particle (Fig. 1B). Nanoparticles showed average diameters lower than 200 nm and PDI less than 0.2, which indicates a narrow distribution (data not shown).

#### **Biochemical and immunological parameters**

As expected, there was an increase in the levels of glucose, fructosamine and a decrease in the levels of insulin in DM and DM+CA groups when compared with the control group (Table 1). The administration of ASc (DM+CA+ASc) attenuated these levels, but did not prevent the decrease of insulin levels when compared to DM+CA. NPASc (DM+CA+NPASc) decreased fructosamine and glucose levels and avoided the decrease of insulin levels when compared with DM+CA (p<0.01). Levels of the pro-inflammatory cytokines IL-1, IL-6, TNF- $\alpha$  and INF-y were increased in DM, CA and mainly in DM+CA when compared to control (p<0.01). ASc (CA+ASc) and NPASc (DM+CA+NPASc) were able to reduce these levels when compared to respective controls (CA and DM+CA). Furthermore, we observed a decrease in IL10 levels in DM, CA and DM+CA when compared to control. Interestingly, ASc did not affect this parameter, but NPASc were able to increase IL10 levels in CA+NPASc and DM+CA+NPASc, in comparison to the respective controls. DPPIV activity was not altered.



Fig. 1: HPLC chromatogram obtained from ASc, NPASc (A) and EE (B).

Gallic acid (peak 1a and b), catechin (peak 2a and b), chlorogenic acid (peak 3a and b), caffeic acid (peak 4a and b), ellagic acid (peak 5a and b), epicatechin (peak 6a and b), rutin (peak 7a and b), isoquercitrin (peak 8a and b), quercetin (peak 9a and b) and kaempferol (peak 10a and b). ASc, aqueous seed extract of *S. cumini*; NPASc, nanoparticle suspension containing ASc.

Table 1: Effect of the treatment with ASc and NPASc on serum biochemical and immunological parameters in rats.

Groups	Glucose	Fructosamin	Insulin	DPPIV	IL1	IL6	TNF-α	IFN-y	IL10
Control	83.25±2.58b	0.89±0.06b	0.34±0.02a	33.19±2.34a	36.50±2.02d	43.75±1.49d	59.00±1.87e	74.75±2.28e	96.00±2.34a
CA	83.75±6.90b	0.83±0.11b	0.39±0.02a	35.65±3.69a	130.0±5.46ab	136.0±3.24b	158.3±1.49b	209.5±6.06a	59.50±2.50cd
CA+ASc	90.05±4.53b	0.86±0.10b	0.39±0.02a	34.34±3.82a	99.75±0.85c	113.0±1.95e	137.3±1.79f	187.8±5.61c	66.50±2.25c
CA+NPASc	90.50±1.91b	1.06±0.03b	0.35±0.02a	33.16±3.89a	89.00±0.91c	101.0±2.48ce	127.0±2.55df	177.8±6.14c	79.50±2.25b
DM	358.2±18.89a	2.15±0.13a	0.17±0.01b	37.86±3.13a	86.75±3.70c	93.25±3.27c	121.8±2.32d	126.0±2.04d	66.50±2.72c
DM+CA	346.50±14.99a	2.04±0.27a	0.16±0.01b	39.82±3.95ab	149.8±6.29e	169.3±4.85a	189.3±6.11a	216.8±8.23a	51.00±2.04d
DM+CA+ASc	180.2±8.75c	1.07±0.10b	0.18±0.02b	41.04±3.05b	140.3±5.96abe	149.0±7.03b	175.5±7.63b	203.0±9.11a	54.50±2.10d
DM+CA+NPASc	153.10±8.83d	1.02±0.14b	0.37±0.03a	44.47±2.24b	127.0±6.33b	141.5±6.34b	166.8±7.71b	206.0±1.68a	65.75±1.75c
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Values are expressed as mean  $\pm$  SEM. Values are expressed as mean  $\pm$  SEM. Different letters indicate significant difference among groups (p< 0.05, ANOVA-Duncan's test). Glucose (mg/dL); fructosamine (mmol/L); insulin (ng/mL); DPP-IV (U/L); IL1, IL6, IL10 and TNF- $\alpha$  (pg/mL); IFN- $\gamma$  (µg/mL).

Table 2: Effect of the treatment with ASc and NPASc on biochemical parameters in lymphocyte suspension.

			*	
Groups	ADA	DPPIV	AChE	NOx
Control	2.52±0.15e	104.00±4.91c	0.67±0.04b	57.68±4.75d
CA	6.59±0.22b	153.20±6.54a	0.76±0.03b	239.10±10.72b
CA+ASc	4.65±0.33c	140.50±5.47ab	0.76±0.05b	264.70±8.68b
CA+NPASc	2.73±0.25e	154.40±9.49a	0.78±0.08b	248.00±12.24b
DM	7.30±0.54b	131.30±2.56b	1.13±0.09a	230.10±12.69b
DM+CA	6.24±0.11b	132.50±4.76b	1.11±0.07a	237.00±12.47b
DM+CA+ASc	5.13±0.39c	125.70±7.36b	0.65±0.01b	135.90±7.41c
DM+CA+NPASc	3.30±0.37d	128.90±10.02b	0.73±0.01b	138.3±7.22c

Values are expressed as mean  $\pm$  SEM. Values are expressed as mean  $\pm$  SEM. Different letters indicate significant difference among groups (p< 0.05, ANOVA-Duncan's test). ADA (U/L/mg of protein); DPPIV (U/L); NOx ( $\mu$ mol/L); AChE (umol AcSCh/h/mg of protein).



**Fig. 2:** NTPDase using ATP and ADP and 5'NT using AMP serum activities and NO serum levels in control, diabetic and/or C. albicans-infected rats treated or not with ASc or NPASc. Values are expressed as mean  $\pm$  SEM. Different letters indicate significant difference among groups (p< 0.05, ANOVA-Duncan's test). A= ADA (U/L); B= AMP, C= ADP (nmol Pi/min/protein); D= ATP (nmol Pi/min/protein) and E= NO (µmol/L)

#### **Biochemical parameters in lymphocyte suspension**

Table 2 shows an increase in lymphocytic ADA, DPPIV and AChE activities and in NO levels in DM and DM+CA rats, when compared to control. The administration of ASc and NPASc prevented the increase in these values, except for DPPIV activity in comparison with DM+CA. Moreover, it was observed an increase in ADA and DPPIV activity and in NO levels in *C. albicans*-infected rats. Only ADA activity was reversed by the treatment with ASc and NPASc in these animals.

## ADA, 5'-NT and NTPDase activities and NO levels in serum

An increase in serum ADA, 5'-NT and NTPDase activities was observed in DM and DM+CA when compared to

control. ASc (DM+CA+ASc) and NPASc (DM+CA+NPASc) were able to decrease these activities when compared to DM+CA (Fig. 2). The activities of these enzymes were also increased in CA group in relation to control, and once again, both treatments (CA+ASc and CA+NPASc) effectively decreased ADA, 5'-NT and NTPDase activities when compared to CA.

NO levels were increased in DM (p<0.05) and CA (p<0.05) and an exacerbated increase was observed in DM+CA (p<0.01) in relation to control. The administration of ASc or NPASc reversed these levels when compared to DM+CA. Conversely, ASc did not affect NO levels in CA+ASc, but NPASc treatment (CA+NPASc) was able to reverse this parameter to near normalcy.

		<u> </u>					
Groups		dney	Li	iver	Pancreas		
	ADA	NOx	ADA	NOx	ADA	NOx	
Control	37.72±3.72d	145.30±11.40e	6.98±0.72d	569.0±15.07b	11.85±1.02bc	427.8±41.47d	
CA	45.94±3.12bc	265.00±8.46b	13.63±0.36a	637.2±37.20ab	13.13±0.79bc	633.6±39.53b	
CA+ASc	39.41±2.64cd	198.30±5.82d	12.59±0.44ab	642.7±25.87ab	9.83±0.85c	411.0±6.02d	
CA+NPASc	44.02±2.05bc	192.30±13.12d	12.65±0.38ab	552.9±36.10b	10.91±1.9bc	374.0±33.94d	
DM	55.44±4.92b	283.40±9.48ab	14.75±0.67a	672.8±38.95a	17.35±1.74a	520.9±11.03c	
DM+CA	51.03±1.93b	293.20±5.70a	14.52±1.39a	680.9±28.60a	13.74±0.50b	639.0±33.66b	
DM+CA+ASc	39.98±2.65cd	234.00±8.42c	10.35±0.50bc	703.6±27.27c	10.75±0.91bc	811.5±16.79a	
DM+CA+NPASc	40.79±0.86cd	212.2±7.33cd	9.92±1.23c	559.2±31.93b	10.84±0.86bc	637.9±24.47b	

Table 3: Effect of the treatment with ASc and NPASc on ADA activity and NOx levels in tissues.

Values are expressed as mean  $\pm$  SEM. Values are expressed as mean  $\pm$  SEM. Different letters indicate significant difference among groups (p< 0.05, ANOVA-Duncan's test). ADA (U/L/mg of protein); NOx ( $\mu$ mol/L).



Fig. 3: NTPDase using ATP and ADP and 5'NT using AMP activities and NO levels in platelets of control, diabetic and/or C. albicans-infected rats treated or not with ASc or NPASc. Values are expressed as mean  $\pm$  SEM. Different letters indicate significant difference among groups (p< 0.05, ANOVA-Duncan's test). A= ADA (U/L); B= AMP, C= ADP (nmol Pi/min/protein); D= ATP (nmol Pi/min/protein) and E= NO (µmol/L).

## ADA, 5'-NT and NTPDase activities and NO levels in platelets

Similarly to the effects observed in serum, ADA, 5'-NT and NTPDase activities were also increased in platelets of DM and/or *C. albicans*-infected animals. The enzymatic activities were decreased in the groups treated with ASc and NPASc. Likewise, NO levels were also increased in DM, CA, and DM+CA, but the treatment was only able to normalize NO levels in CA+NPASc (Fig. 3).

## ADA and NO in tissues

ADA activity was increased in animals with DM and/or infected with C. albicans, mainly in the kidney and liver. Kidney and liver ADA activities in DM animals infected with C. albicans and treated with ASc (DM+CA+ASc) or NPASc (DM+CA+NPASc) were decreased when compared to the respective control (DM+CA) (Table 3). This effect was not observed in non-diabetic rats infected with C. albicans and treated with ASc (CA+ASc) or NPASc (DM+CA+NPASc). Pancreas ADA activity did not differ among groups, except for the significant increase observed in DM (p<0.01). Kidney NO levels were also increased in rats with DM and/or C. albicans infection in

comparison to the control group. Groups treated with ASc (CA+ASc and DM+CA+ASc) and NPASc (CA+NPASc and DM+CA+NPASc) showed a significant decrease in kidney NO levels in comparison to their respective controls (CA and DM+CA). In the liver, an increase in NO levels was observed in DM and CA, and the treatment with ASc (DM+CA+ASc) and NPASc (DM+CA+NPASc) had effect when compared to DM+CA (p<0.01). Finally, an increase in NO levels was observed in the pancreas of animals in DM, CA and DM+CA groups when compared to the control group. The treatment was only effective in CA+ASc and CA+NPASc when compared with CA.

#### In vivo antifungal activity

Fungal tissue burden counts in kidney and liver were of  $10.25\pm0.85$  CFU/g and  $2.53\pm0.39$  CFU/g (CA) and of  $13.25\pm1.25$  and  $3.66\pm0.8$  CFU/g (DM+CA), respectively. Kidney and liver of rats treated with ASc showed fungal load of  $9.5\pm0.5$  and  $2.5\pm0.5$  CFU/g (CA+ASc) and  $12.25\pm1.01$  and  $2\pm0.57$  CFU/g (DM+CA+ASc), respectively. Kidney and liver of rats treated with NPASc showed fungal load of  $7.3\pm0.5$  and  $2\pm0.1$  CFU/g (CA+NPASc) and  $11\pm2.05$  and  $1.25\pm0.25$  CFU/g

(DM+CA+NPASc), respectively. No fungi were recovered from animals not infected by *C. albicans* (control and DM).

## DISCUSSION

We examined the metabolic changes in a short-term DM model with C. albicans infection, their association with inflammatory changes and disturbances in adenine nucleosides and nucleotides and the treatment with S. cumini seeds in the form of extract and polimeric nanoparticles. DM promoted an increase in 5'-NT and NTPDase activities in serum and platelets as a response to metabolic insult, which is in agreement with other studies (Lunkes et al., 2003; Lunkes et al., 2004; Schmatz et al., 2013). Curiously, the animals inoculated with C. albicans also showed an increase in ATP, ADP and AMP hydrolysis in serum and platelets, contributing to an increase in adenosine production. This was accompanied by an increase in ADA activity, which could be related to an attempt of the organism to compensate organic alterations such as the high levels of ATP released during inflammation, leading to an increase in adenosine deamination. In line with the enhancement of ectonucleotidase activities in serum and platelets. ADA activity was increased in kidney, liver and pancreas of DM animals, corroborating previous studies (Schmatz et al., 2009; Bitencourt et al., 2015). An increase in ADA activity was also observed in the same organs of C. albicans-infected animals. The decrease in adenosine levels is likely to contribute to a high inflammatory state well-known in DM, with increased release of pro-inflammatory cytokines (IL-1, IL-6, TNF- $\alpha$  and IFN-y) and NO, as observed in DM, CA and DM+CA groups. These groups also showed increased ADA and DPPIV activities in lymphocyte suspension. This may be related to the interaction of ADA and DPPIV on T cells, resulting in costimulatory signs responsible for the activation of the T-cell receptor, presenting an important role in immune responses (Gorrell et al., 2001; Elgün et al., 2001).

Taken together, these findings indicate that the upregulation of ectoenzymes in serum and lymphocytes, which reflects in the immune system, and in the platelets, an important source of nucleotides and nucleosides, may play important roles in (1) controlling cellular responses induced by diabetes complications and (2) attenuating systemic changes caused by the fungal infection (Schmatz et al., 2013; Idzko et al., 2014). An important finding of this study was that both ASc and NPASc were able to reduce 5'-NT, NTPDase and ADA activities in cells and serum, and thus facilitating the regulation of the immune system for assisting the host response in an attempt to control the inflammation caused by DM and C. albicans. Moreover, ASc and NPASc decreased the activity of this enzyme in the kidney and liver of the treated groups, thus modulating adenosine concentrations. This effect is highly relevant, since adenosine has been proven to play an important role in the modulation of insulin action on glucose metabolism in different tissues and to reduce inflammation and tissue injury (Rutkiewicz and Górski, 1990; Cuzzocrea *et al.*, 2000). The model of candidiasis used in the present experiment is a sub-acute systemic infection that is usually well tolerated by the animals for several weeks, is not cleared spontaneously and mimics human infection (Fisher *et al.*, 1989). Therefore, as expected, we observed an increase in proinflammatory cytokines in the group infected with *C. albicans*. Moreover, hyperglycemia is known to increase the production of free radicals and to induce inflammation, which can lead to a permanent stimulation of immune cells (Buchta *et al.*, 2013; Kim *et al.*, 2014).

This is in agreement with the extremely strong inflammatory response observed in DM rats infected with *C. albicans*. Nonetheless, ASc and NPASc were able to decrease the levels of pro-inflammatory cytokines and to increase the levels of the anti-inflammatory cytokine IL-10, attenuating the deleterious effects of the exacerbated inflammatory response. These results are likely to be related to the modulation of ectonucleotidase activities by *S. cumini*, reflecting in the increase of extracellular adenosine and, consequently, regulating host immune responses. Moreover, secondary metabolites of *S. cumini* extract have been reported to block the activation of nuclear factor kappa B (NF- $\kappa$ B), a proinflammatory transcription factor that leads to activation of inflammatory mediators in several pathologies (Donepudi *et al.*, 2012; Swami *et al.*, 2012).

The antifungal activity of *S. cumini in vitro* is still quite controversial (Chandrasekaran and Venkatesalu, 2004; Shad *et al.*, 2014), possibly due to different concentrations of phytochemical compounds present in extract formulations of plants obtained from different geographical areas. Previously, we have reported that gallic and chlorogenic acids, compounds which are present in greater quantities in the extract, showed low or no antifungal activity against fungi when evaluated alone. Notwithstanding, the formulations containing the whole extract, ASc and NPASc, showed increased antifungal activity *in vitro*, demonstrating a synergistic effect when the compounds are used together (Bitencourt *et al.*, 2016).

Here we observed that DM rats had higher fungal burden than non-diabetic. No antifungal activity was observed for ASc, but liver and kidney of rats treated with NPASc showed lower fungal burden in comparison to controls. The antifungal activity, associate with the immune response developed by the animals, support the hypothesis that the anti-inflammatory activity of the extract *in vivo* is more prominent than the antifungal activity. To the authors' knowledge, this is the first study evaluating the effects of *S. cumini* in DM rats with candidiasis.

The role that host-derived ACh plays in modulating the growth and pathogenicity of microorganisms is unclear. We observed an increase in AChE activity in lymphocyte suspensions in DM, CA and DM+CA groups. ASc and NPASc were equally able to decrease the activity of AChE in DM+CA+ASc and DM+CA+NPASc, increasing ACh levels, which has recently been shown to be an inhibitor of *C. albicans* biofilm formation and pathogenicity (Rajendran *et al.*, 2015).

Indeed, ACh has anti-inflammatory properties that could have modulated the immune responses in the rats, contributing to increase IL-10 levels. In line with Helmstadter (2008), we observed a hypoglycemic effect in the animals treated with ASc, but NPASc treatment was able to reduce the glucose more effectively, probably because of its capacity to normalize insulin levels. PCL polymer has been used in the development of nanoparticulate systems of natural products and has presented several advantages such as maintaining the antioxidant capacity, bioavailability, solubility and stability of polyphenols (Sanna *et al.*, 2015; Ng *et al.*, 2015). Consequently, it can promote an improvement of the known pharmacological activity of these substances such as stimulating insulin secretion from the remaining  $\beta$ -cells and mimicking insulin activity, resulting in peripheral glucose uptake (Ayyanar *et al.*, 2013).

The effects of STZ on  $\beta$ -cells may be mediated by local liberation of NO from STZ within islets, contributing to a part of its diabetogenic potential and exaggerating  $\beta$ -cell necrosis (Turk et al., 1993). In fact, other studies have reported an increased NO concentration in serum and tissues (Yang et al., 2014; Sokolovska et al., 2015; Varsha et al., 2015) and inducible nitric oxide synthase (iNOS) overexpression in STZ-induced diabetic rats (Al-Rejaie et al., 2015). In this context, we observed an increase in NO levels not only in pancreas, but also in the liver and kidneys of DM and/or C. albicans-infected animals, which can be related to the fact that besides of the enhancement in interactions of superoxide with NO in the oxidative diabetic environment, NO is involved in pathogen killing mechanism and contributes to damage in other tissues (Shahani and Sawa, 2011; Samarghandian et al., 2013). The higher effect of NPASc in decreasing the levels of NO in tissues in comparison to ASc can be attributed to the physicochemical characteristics of nanoparticles that can facilitate crossing membranes and interacting more effectively (Jain and Kumar, 2010). A second advantage of this formulation is its high capacity to encapsulate not only hydrophilic but also lipophilic compounds such as rutin and quercitin, which have important pharmacological actions and can be adsorbed in the polymeric wall. In addition, this system provides greater protection against oxidation and other degrading reactions that occurs in the initial segments of intestine, where phenolic substances present in the extract have major absorption (Mora-Huertas, 2010; Ferriz and Vinová, 2010; Bonifácio et al., 2014).

## CONCLUSION

In conclusion, the involvement of ectonucleotidases during infections has been described for several microorganisms. However, there is little information on how these enzymes would contribute to the understanding of pathological condition in fungal infections. Rats with DM and/or *Candida* infection showed increased levels of ectonucleotidases and pro-inflammatory cytokines. ASc and NPASc were able to attenuate hyperglycemia and to prevent the increase in cytokine levels and in ectonucleotidase and AChE activities. As purines are recognized for their important role in modulating processes linked to inflammation, this may take action to regulate cytokine levels which in turn helped to decrease NO levels. Thus, we suggest that the modulation of ectoenzyme activities can be one of the mechanisms by which *S. cumini* act on cytokines that affect the development of chronic complications of DM. Of note, since NPASc was able to maintain the properties of the crude extract, this drug delivery system may be used for other hydrophilic drugs of medicinal plant origin.

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