

Calorimetric Markers for Inflammation in *in vivo* Experimental Models

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Received: January 15, 2019

Accepted: March 26, 2019

Published: December 31, 2019

Abstract: In this work differential scanning calorimetry was applied to determine the specific calorimetric features of blood plasma proteome associated with immune response stimulation in experimental model (albino Wistar rats). The thermodynamic behavior of the blood plasma of male and female animals subjected to egg albumin (EA) treatment was investigated. The calorimetric profiles of blood plasma from EA treated rats exhibited reduced heat capacity of the albumin-assigned transition and up-shifted weighted average center of the thermogram as compared to healthy controls, the effect being more pronounced for male animals. Increase in the amplitude of the main transition at 70 °C was observed for female rats after EA treatment, which resulted in higher calorimetric enthalpy. Common feature of the thermograms of EA treated males and females was the broadening of the transitions above 75 °C and the appearance of exothermic transition above 90 °C due to protein aggregation.

Our study clearly revealed gender-specific immune response in rats and contributes to better understanding of the correlation between the calorimetric features of blood plasma and the immunological conditions in the experimental animals.

Keywords: Chronic inflammation, Animal model, Differential scanning calorimetry, Blood plasma, Egg albumin.

Introduction

Inflammation is a protective body mechanism through which the immune system responds to pathogens, injuries and other damages of tissues. Although inflammation is essential for resistance to infections and for healing, long lasting and persistent inflammatory processes can become damaging and thus underlie the pathogenesis of many diseases including cancer. Over the past decade, new proofs have emerged confirming that chronic inflammatory responses facilitate cancer cell proliferation and tumor growth, from tumor promotion to

progression [3, 14]. For example, the risk of occurrence of colorectal carcinoma is increased for patients with chronic ulcerative colitis or Crohn's disease [20]. It is suggested that more than 50% of gastric adenocarcinomas and Lymphoma of gastric-mucosa-associated lymphatic tissue are due to infection with *Helicobacter pylori* [4, 26]. Hepatitis C virus causes acute and chronic infection and is one of the major causes for the development of hepatocellular carcinoma [1]. Chronic inflammation is also linked to the development of lung, breast and skin cancer [16, 21]. In many studies, a relationship between proinflammatory markers and the outcome of the disease in cancer patients has been established [5, 22]. Therefore, in-depth understanding of the inflammatory progression is important for the development of improved therapeutic strategies and to avoid unwanted consequences of the sustained inflammation.

Animal models are often utilized for investigation of the role of chronic inflammation for the promotion and development of cancer and other inflammation related diseases [2, 6, 13, 15, 24], as well as in pre-clinical studies in order to better understand the physiological and biochemical processes, associated with occurrence and progress of diseases. In this respect, it is crucial to develop tools that discriminate those different conditions.

Modern medicine thrives to develop non-invasive screening procedures that identify biomarkers for reliable and specific diagnosis of variety of diseases. In the last decade differential scanning calorimetry (DSC) was recognized as a method with high potential for disease diagnostics and monitoring since the thermal stability of the major blood plasma/serum proteins was shown to be altered in variety of diseases and conditions [9, 11, 17, 18, 25, 29, 30, 33]. However, so far only few studies apply DSC on blood serum derived from animal model [8, 23, 28]. Hence, the correlation between the calorimetric features of plasma/serum proteome and the health status of animal model still needs further in-depth studies.

The aim of this work is to determine the specific thermodynamic features of blood plasma proteins associated with the immune response stimulation in experimental rat models. We studied the thermodynamic behavior of the blood plasma proteome of male and female albino Wistar rats subjected to egg albumin treatment and the levels of the major serum proteins. Data clearly reveal gender-specific immune response in rats and therefore contribute to better understanding of the correlation between the calorimetric features of blood plasma and the immunological condition in *in vivo* experimental rat models.

Materials and methods

Experimental models

A total of 20 eight-month old albino Wistar rats (10 females and 10 males) were used as experimental animals in this work. The animals were obtained from the Experimental and breeding base for laboratory animals in Slivnica, and were kept in the vivarium in the Institute of Experimental Morphology, Pathology and Anthropology with Museum. The rats were divided into groups of 5 animals of the same sex and were housed in plastic cages in standard laboratory conditions (at a constant temperature of 23 ± 2 °C, humidity of $55\% \pm 5\%$ and 12 h light/dark cycle), with free access to standard dry pellets and water *ad libitum*. The animals were fed with a standard diet throughout the entire duration of experiment. The treatment of the animals started after 7 days of acclimatization.

The experiments were approved by the ethical committee of the Institute of Experimental Morphology, Pathology and Anthropology with Museum and were in accordance with the ethical committee guidelines.

Immune response induction

Egg albumin (EA) was used as an antigen to induce an immune response in the model animals. 5 female and 5 male rats were injected subcutaneously with the antigen (150 µg/kg) 3 times during one month (i.e., every 10 days). First injection was carried out with 200 µl/rat immunological solution of EA in PBS buffer, Al(OH)₃ and Complete Freund's Adjuvant for sustaining the immune response. Incomplete Freund's adjuvant we used in the solution for the second injection. The solution for the third injection was prepared with only Al(OH)₃ in PBS and EA.

The remaining female and male rats were used as a control groups and were injected with sterile PBS buffer, pH 7.4.

Blood plasma preparation

All rats were sacrificed under deep anesthesia 10 days after the last injection. Blood was drawn by cardiac puncture in vacutainer tubes with anticoagulant (K₂EDTA). The whole blood was centrifuged for 15 minutes at 2800 rpm. The supernatant (blood plasma) was transferred into Eppendorf tubes and stored at -80 °C until thawed for measurement of antibody titers, protein fractions and DSC analysis.

Protein analysis

The total protein content in the blood plasma was determinate by the Biuret method [27]. The levels of the main plasma protein fractions – albumin, α_1 , α_2 , β_1 , β_2 and γ -globulins were determined by Capillary electrophoresis (Capillarys 2, Sebia).

Indirect ELISA

Blood plasma of the experimental animals were assayed by indirect ELISA. 96-well ELISA plates (Greiner Bio-One) were coated with the antigen (egg albumin) in 50 mM carbonate buffer (pH 9.6) to concentration 10 µg/ml. The plate was incubated for 1 h at 37 °C, and than was blocked in 3% skimmed milk in PBS for 40 min. After washing three times for 5 minutes on a shaker with 0.05% Tween 20, the rat's plasma was added in dilution 1:1000 and was incubated for 2 hours at room temperature. The plate was washed with 0.05% Tween-20 three times, after which the secondary antibody Goat Anti-Rat IgG-HRP (Santa Cruz Biotechnology, USA) was added, diluted 1:5000 in PBS and was incubated for 2 hours at room temperature. After the final wash, color was developed by incubating with o-phenylenediamine (0.5 mg/mL in citrate buffer) and 5 µL of 30% H₂O₂ for 20 minutes. The reaction was stopped by adding 50 µL/well of 1M HCl. Optical density at 492 nm was measured with a microplate reader (TECAN, Austria).

Differential scanning calorimetry

DSC profiles were obtained using DASM 4 (Privalov, BioPribor)-built-in high sensitive microcalorimeter. Blood samples were diluted 8-10 times in PBS buffer (pH 7.4) and thermograms were recorded from 30 °C to 105 °C at 1 °C/min scanning rate. A constant pressure of 2 atm was maintained during the experiments. The baseline (buffer–buffer) scans were subtracted from the plasma scans that were further corrected applying a linear baseline fit. Thermograms were normalized to the total protein concentration to obtain the partial heat capacity as a function of temperature. The calorimetric profiles were analyzed using the ORIGIN 8 (MicroCal Software) program package and the following parameters were determined: enthalpy of the thermogram (ΔH), weighted average center of the thermogram (T_{FM}), denaturation temperature (T_m^{max}) and excess heat capacity (c_p^{max}) of the main transition.

Statistical analysis

Wilcoxon rank-sum test non-parametric analysis was used to compare the thermograms recorded for control and EA treated animals, the data were regarded as significantly different at $p < 0.05$.

Results and discussion

To probe the effect of EA induced inflammation on the experimental rat models we employed indirect ELISA analysis. Both female and male animals showed a significant IgG antibody response to the antigen challenge (Fig. 1). After the end of the treatment, the animals had high plasma IgG anti-EA titers, while control rats showed no anti-EA antibodies. Although the difference is not statistically distinct, it should be taken into account that the immune response is higher in male than in female rats.

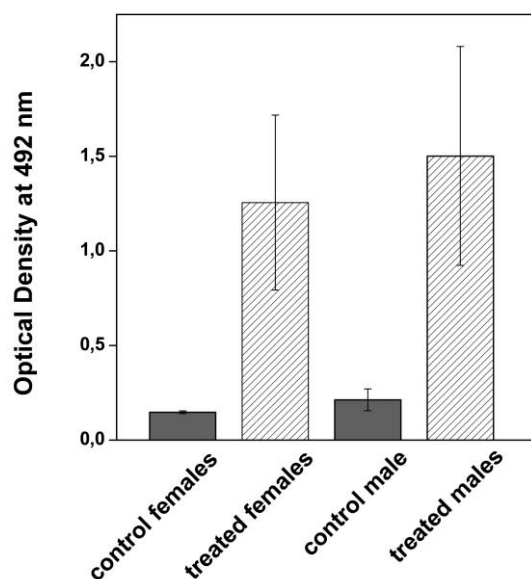


Fig. 1 Indirect ELISA test performed on blood plasma derived from male and female untreated controls and animals treated with egg albumin. Mean \pm SD, $n = 5$.

Next we evaluated the effects of the triggered immune response on the levels and the thermodynamic properties of the major blood plasma proteins by means of capillary electrophoresis and DSC, respectively.

The obtained values for the major plasma proteins (see Table 1) are similar to those previously reported for rats [32] with the exception of the γ -globulins fraction that is higher in this study. No statistically significant difference was found between the untreated and treated animals of the same sex. The plasma protein content of control male rats was characterized with lower albumin content and higher level of β_2 -globulin fraction than that of females (Table 1).

The calorimetric profiles of “healthy” and EA treated rats are presented in Fig. 2. The DSC scans of both female and male rats possess similar characteristics – a dominant transition at 70 °C and several smaller transitions at 51 °C, 62 °C, 81 °C and 89 °C (Fig. 2). On the basis of a number of studies performed on human blood plasma [10, 12, 31] and the data of Kosa et al. [7], demonstrating that albumin isolated from human and rat blood plasma exhibit a single sharp endothermic transition at ca 60 °C, the transitions resolved in the rats thermograms can be assigned to the melting of fibrinogen (51 °C), albumin (62 °C), globulins (70 °C), IgG and

transferrin (above 80 °C). The heat capacity ratio of albumin/globulin assigned transitions is less than 1 for control rats, in contrast to that of human plasma thermograms where it is 2.0 [31]. Most likely this is due to the lower percentage of albumin from the total protein content in rats (ca. 40%) than in humans (ca. 60%) (see Table 1 and [19]). The amplitude of the albumin transition in the thermograms of male rats is much lower than that for female ones and is visible only as a shoulder (Fig. 2) – this effect must be due to the lower albumin content in males than in females (Table 1).

Table 1. The levels of plasma proteins (albumin, α_1 and α_2 , β_1 - and β_2 -, and γ -globulin fractions) determined by capillary electrophoresis for control and EA treated male and female rats. Mean \pm SD, $n = 5$.

Groups	Albumin, [%]	α_1 , [%]	α_2 , [%]	β_1 , [%]	β_2 , [%]	γ , [%]
Control female	41.3 \pm 0.3	18.8 \pm 0.5	8.7 \pm 0.5	2.9 \pm 0.3	19.9 \pm 0.5	8.4 \pm 0.3
Female + EA	41.1 \pm 0.7	19.3 \pm 1.6	7.8 \pm 0.3	2.4 \pm 0.5	20.4 \pm 0.7	8.9 \pm 1.1
Control male	38.5 \pm 1.1*	18.5 \pm 1.5	9.6 \pm 0.4	1.8 \pm 0.7	22.0 \pm 1.1*	9.7 \pm 1.5
Male + EA	38.0 \pm 2.0**	19.0 \pm 1.5	9.9 \pm 0.8**	2.4 \pm 1.2	21.6 \pm 1.5	9.3 \pm 0.9

* indicates significant difference from control female animals

** stands for significant difference from treated female animals

The amplitude of the dominant transition, c_p^{max} , is slightly lower for female than for male control rats; however, no significant difference between males and females was observed for the enthalpy of the thermogram, the position of the dominant peak and the weighted average center of the thermogram (Table 2).

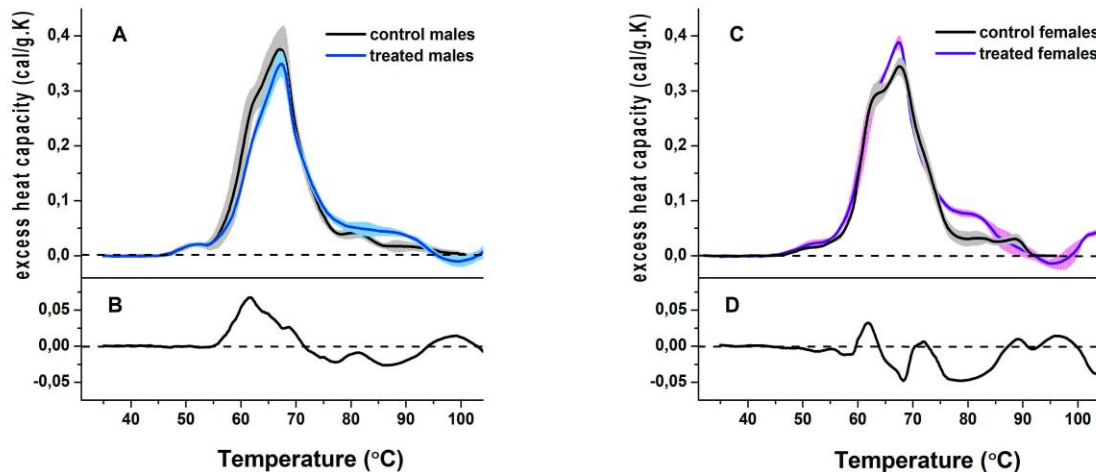


Fig. 2 DSC profiles (average – solid lines and standard deviations – shading) of blood plasma from controls and rats treated with egg albumin. Untreated male and female rats (black solid line and gray shading), A and C; treated male rats (blue solid line and cyan shading), A; treated female rats (violet solid line and magenta shading), C; difference thermograms obtained after subtraction of the averaged plasma denaturation profiles of control and treated rats, B and D.

The effect of EA treatment did not cause significant change in the levels of the major plasma protein fractions (Table 1), like for untreated rats the treated female animals had higher albumin and lower α_2 level as compared to treated male subjects (Table 1). The main effect of the applied treatment is on the amplitude of the albumin assigned transition at 62 °C that is

lower for treated male while higher for treated female rats compared to untreated ones (Table 2). The weighted average center of the thermograms, T_{FM} , is up-shifted by ca. 1 °C for treated males (Table 2) in comparison to untreated ones. For females the peak at 62 °C appears as a shoulder after EA treatment (Fig. 2B). In addition, a slight increase in the amplitude of the main transition at 70 °C was observed, which also resulted in higher enthalpy after EA treatment of female animals (Table 2).

Table 2. Thermodynamic parameters estimated from the calorimetric profiles of control treated male and female rats: enthalpy of the thermogram (ΔH), weighted average center of the thermogram (T_{FM}), denaturation temperature (T_m^{max}) and excess heat capacity (c_p^{max}) of the main transition. Mean \pm SD, $n = 5$.

Groups	ΔH , [cal·g ⁻¹]	T_{FM} , [°C]	T_m^{max} , [°C]	c_p^{max} , [cal·g ⁻¹ ·K ⁻¹]
Control females	4.6 \pm 0.3	66.9 \pm 0.3	67.6 \pm 0.2	0.35 \pm 0.02
Female rats + EA	5.2 \pm 0.3*	67.2 \pm 0.2	67.5 \pm 0.2	0.39 \pm 0.02*
Control males	4.8 \pm 0.3	66.3 \pm 0.4	67.0 \pm 0.6	0.38 \pm 0.05
Male rats + EA	4.6 \pm 0.2	67.4 \pm 0.3*	67.4 \pm 0.3	0.35 \pm 0.02*

* statistically significant difference from control animals, $p < 0.05$

A common feature between the thermograms of EA treated males and females is the broadening of the transitions above 75 °C and the appearance of an exothermic transition above 90 °C that must be due to protein aggregation, an effect that is gender-specific since the exotherm is located at 99 °C for males, while at 95 °C for females (Fig. 2). Similar effect was also observed by Michnik et al. [23] for rats receiving weakly 8 mg·kg_{BM}⁻¹ of testosterone enanthate; according to our data the hormonal treatment might induce immune response. The difference thermograms obtained after subtraction of the DSC profile of plasma derived from EA treated animals from that of control ones for males and females do not overlap in the range of the thermal scanning (Fig. 2B and D), revealing distinct response to the applied treatment for the two sexes. In addition, Wilcoxon non-parametric analysis reveals that the plasma thermograms of control and EA treated male rats differ significantly ($p < 0.01$) in the 57-68 °C and 74-90 °C temperature ranges, while those of females differ in much narrow temperature ranges, i.e., 64-69 °C and 74-85 °C. It should be noted that Garbett et al. [9] also found differences in the thermograms of patients with inflammatory disease (Systemic lupus erythematosus) related to their sex, that involved differences in the albumin assigned transition.

The presented data give insight on the effect of persistent inflammation on the DSC profiles of rat blood plasma which are somewhat similar to that of soft tissue inflammation in humans [31]. The excess heat capacities of albumin assigned transition decreased, whereas that of the globulin increased in humans with inflammation and treated female animals. Increased T_{FM} is also a common feature for the thermograms of patients with inflammation and treated male rats. We should note some differences between the “inflammation” thermograms of rats and of humans as compared to their respective controls: (i) the DSC profiles of plasma from patients with inflammation present the same number of successive transitions as the control one in contrast to rats plasma, where only one broad peak is depicted in the high temperature region (above 75 °C) of EA treated animals instead of two transitions in the respective controls; (ii) protein aggregation is observed for both treated animal groups, that is not occurring in the human profiles.

Conclusion

In summary our data revealed gender-related differences in the albumin content as well as in the calorimetric profiles of male and female rats involving the amplitude of the albumin assigned transition and its behavior after EA treatment. The presence of an exothermic transition appears as a specific feature of the induction of immune response in both males and female rats, its position was also found to be sex-dependent.

The model of chronic inflammation used in this study is valuable tool in exploring the potential of DSC analysis for the characterization of chronic, inflammatory responses in animal model.

The presented data prompt future studies on the effect of different diseases on the calorimetric profiles of rat blood plasma, as well as their gender specific features.

Acknowledgements

This work is supported by grant DFNP-17-133, Program for career development of young scientists and PhD students in Bulgarian Academy of Sciences, 2017 (A.D.).

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