

Identification and Characterization of the Lipid Transport System in the Tarantula *Grammostola rosea*

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Abstract

Lipids due to its hydrophobic nature are transported in the hemolymph mainly by lipoproteic fractions. In the present work we studied the lipoproteic fractions present in the hemolymph of the spider *Grammostola rosea*. Through ultracentrifugation two lipoproteic fractions are isolated, one of high density named Gr-HDL ($\delta = 1.13 - 1.15$ g/ml) and the other of very high density called Gr-VHDL ($\delta = 1.24 - 1.27$ g/ml) Gr-VHDL in hemolymph is majority in relation to Gr-HDL. In this sense Gr-VHDL fraction has 98.6% of hemolymphatic proteins, and 89.3% of lipids presents in the hemolymph. Both lipoproteic fractions possess phospholipids such as majority lipids (phosphatidylcholine and phosphatidylethanolamine) and 18:1, 16:0, 18:2 and 18:0 as the major fatty acids. In order to confirm the role played by lipoproteic fractions *in vitro* assays with different ^{14}C -lipid were performed. It was observed that Gr-VHDL takes up mainly free fatty acids and triacylglycerols unlike that observed for Gr-HDL in relation to phosphatidylcholine. Through electrophoresis it was observed that Gr-VHDL has three proteins: a predominant band of 68 kDa and two others of 99 and 121 kDa. Gr-HDL displayed a predominant band of 93 kDa, and other minority of 249 kDa. In conclusion, this study reports lipid characterization of the lipoproteic fractions present in the hemolymph of the tarantula, *G. rosea*. The role of each lipoproteic fraction in relation to lipid uptake is sustained by *in vitro* assays. Similarities and differences are found when it is compared to lipoproteins of only the three species of spiders studied.

Keywords

Lipoproteins, Hemocyanin, Arachnids, Lipids

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1. Introduction

Lipids are a heterogeneous group of molecules involved in many vital functions such as energy storage, formation of different structures, source of essential nutrients, vitamins, and chemical messengers. In general, hydrophobic characteristics of lipids determine their circulation mode in the aqueous fluids of animals as particles called lipoproteins.

Arthropod lipid circulation mechanisms are well known only in insects and crustaceans. Insect lipids are mostly carried by a lipoprotein transporting a great quantity of diacylglycerols [DAG] named lipophorin [1]-[5]. In crustaceans, lipids are transported by high-density lipoproteins (HDL) mostly in the form of phospholipids [PL] [6]-[10]. Surprisingly, scarce information is available for arachnids, where only in three out of 43,600 species of the described spiders, lipoproteic fractions have been isolated and characterized [11]-[16].

The first spider which hemolymphatic lipoproteins were characterized was the tarantula *Eurypelma californicum* (a nomen dubium that probably corresponds to *Aphonopelma hentzi*, following Nentwig, 2012) (Araneae, Theraphosidae). Haunerland and Bowers [15], on the basis of density gradient ultra centrifugation of hemolymph, detected three distinct protein fractions in this species. One supernatant corresponds to hemocyanin [Hc], and two others float at lower densities, which were identified as lipoproteins with HDL and VHDL characteristics. HDL with a hydrated density of 1.12 g/ml is composed of approximately 44% lipid (mainly DAG and PL). In contrast, VHDL fraction with density of 1.28 g/ml is composed of approximately 8% lipid [15].

Stratakis *et al.* [16] purified and characterized a VHDL from tarantula hemolymph with a density of 1.23 g/ml. Its lipid content from 16% to 19% (mainly PL and free fatty acids [FFA]) apparently differs from that observed by Haunerland and Bowers [15].

Lipid transport hemolymphatic system of *Polybetes pythagoricus* (Araneae, Sparassidae) is composed of three lipoproteic fractions: two HDLs fractions named HDL-1 ($\delta = 1.13$ g/ml, 28% lipids by wt) and HDL-2 ($\delta = 1.18 - 1.20$ g/ml, 3.6% lipids by wt), and a VHDL fraction ($\delta = 1.21 - 1.24$ g/ml, 3.5% lipids by wt). Their hemolymph concentrations are 2.3, 23.6 and 45.4 mg of protein/ml hemolymph, respectively. Circulating lipids in *P. pythagoricus* (3.41 mg/ml hemolymph) are therefore carried by VHDL (48.7%), HDL-1 (26.2%), and HDL-2 (25.1%) [11] [14]. HDL-1 and VHDL fractions mainly transport phosphatidylcholine [PC], and in a lesser extent FFA, triacylglycerols [TAG] and phosphatidylethanolamine [PE]. The latter has a bluish coloration given by the high amount of Hc which was described as apolipoprotein [11] [14].

Two HDLs fractions were isolated from the hemolymphatic plasma of the spider *Latrodectus mirabilis* (Araneae, Theridiidae). HDL-1 fraction has a density of 1.13 g/ml and HDL-2 fraction of 1.19 - 1.20 g/ml; their hemolymph concentrations are 4.83 and 31.6 mg of protein/ml hemolymph, respectively. Circulating lipids carried by HDL-1 are 80% and those by HDL-2 are 20%. HDL-1 mainly transports PC, FFA, and TAG. HDL-2 characterized by having Hc transports FFA, PC and cholesterol [CHOL] among others [12].

Hemocyanin [HC] plays its principal role as oxygen carrier, but it also has other properties and/or functions such as phenoloxidase activity, antimicrobial activity and apolipoprotein associated with the majority of circulating lipids [12]-[14] [17]-[22]. The hemocyanin-lipids interaction was reported for the first time by Zatta [23] who determined the existence of small amounts of PL bound to Hc of the crab *Carcinus maenas*; he assigned a stabilizing role in protein structure to this lipid fraction.

The amount of lipids was found to be significant in lipoproteins containing Hc as apolipoprotein in the spiders *P. pythagoricus* and *L. mirabilis*. In fact, though VHDL and HDL-2 fractions have a low lipid/protein ratio, they are of great quantitative importance in *P. pythagoricus* hemolymph, since they are bound to two thirds of hemolymphatic lipids [14] and the same occurrence, though in a minor proportion, was observed in the HDL-2 of *L. mirabilis* [12].

Recently, we have demonstrated for the first time that the midgut diverticula of arachnids are a main storage site and a major lipid metabolic center involved in the uptake and mobilization of lipids. It also clearly demonstrates that innovatively the VHDL (fraction with Hc) from *P. pythagoricus* participates in the lipid transport from and toward tissues [24] [25].

The aim of our work was to identify and characterize the components of the lipid transport system in the spider *Grammostola rosea* (Theraphosidae). We show data of the lipid composition of hemolymphatic fractions. Also with the use of radiolabelled lipids we comparatively evaluated the uptake of different lipids (FFA, PC and TAG) in the lipoproteic fractions and hemocytes present in hemolymph.

2. Materials and Methods

2.1. Sampling and Isolation of Lipoproteins

Adults females of *G. rosea* (13 to 18 gr) were maintained in the laboratory at 22°C under a 14/10 h L:D photoperiod. The spiders were fed with cockroaches (*Blattella germanica*) and water. The specimens were anesthetized using CO₂ similar to Baumann *et al.* [26]. Hemolymph samples were collected using a syringe with a needle designed for animal injection (Hamilton Co.) with 0.1 N of sodium citrate as anticoagulant in accordance with Garcia *et al.* [27]; with the addition of a protease inhibitor cocktail 1/1000 v/v (Sigma Chemicals, St. Louis, MO, USA). The extraction was performed by needling one of the back legs. After hemolymph collection, the spiders were fed and watered.

Hemocytes were isolated by diluting hemolymph in buffer Tris-HCl 0.1 M, pH 8, centrifuged at 720 g for 10 min, then the pellet was resuspended and centrifuged again at 720 g for 30 min following the methodology by Laino *et al.* [25].

The isolation of lipoproteic fractions with and without Hc was performed using ultracentrifugation in density gradient, following the methodology previously employed for hemolymph of spiders [12] [14] [18]. Aliquots of clear blue hemolymph (1.4 ml) were overlaid on 3 ml NaBr solution (density 1.35 g/ml) and centrifuged at 178,000 g at 10°C for 22 h in a Beckman L8 70 M centrifuge with a SW 60 Ti rotor. As the plasma density was 1.006 g/ml, a saline solution of the same density was run simultaneously as blank. The total volume of the tubes was fractionated from top to bottom into 0.2 ml aliquots. The protein content of each fraction was monitored spectrophotometrically at 280 nm.

2.2. Protein Analysis

Total protein concentration from different isolated lipoproteic fractions were measured colorimetrically by the method of Lowry *et al.* [28]. The apoproteins were analyzed by dissociating electrophoresis using a 4% - 23% SDS-PAGE [29]. Gels were stained with Coomassie Brilliant Blue R-250 (Sigma Chemical Co.). Molecular weights were calculated as previously described [30].

2.3. Lipid Analysis

Lipids from lipoproteins were extracted following the procedure of Folch *et al.* [31]. Quantitative determination of lipid classes was performed by thin layer chromatography (TLC) coupled to a flame ionization detector (FID) in an Iatroscan apparatus model TH-10 (Iatron Laboratories, Tokyo, Japan), after separation on Chromarods type S-III [24] [32]. Lipid classes were quantified using monoacylglycerol as internal standard. The general procedure for separation, identification, and quantification of lipids was similar to that described in a previous work [14].

2.4. Fatty Acid Analysis

Fatty acid methyl esters (FAME) from total lipids from hemolymphatic lipoproteins were prepared with BF₃-MeOH, according to the method of Morrison and Smith [33]. The analysis was performed by gas-liquid chromatography (GLC-FID) using an HP-6890 capillary GLC (Hewlett Packard, Palo Alto, CA) fitted with an Omegawax 250 fused silica column, 30 m × 0.25 mm, with 0.25 μm phase (Supelco, Bellefonte, CA). Peaks were identified by comparing the retention times with those from a mixture of standard methyl esters according to Garcia *et al.* [27].

2.5. Lipid Uptake by Hemolymphatic Lipoproteins and Hemocytes

In order to explore in the tarantula *G. rosea* the role played by hemolymphatic lipoprotein fractions and hemocytes in the uptake of different lipids, a series of experiments were performed following the methodology employed in arachnids by Laino *et al.* [25] with modifications. 0.5 ml of hemolymph was incubated *in vitro* for 1 hour in 50 mM potassium phosphate buffer pH 7.4, with the following radiolabelled lipids in two different concentrations (low concentration of 0.16 μCi and high concentration of 0.16 μCi + 500 μg of non-radiolabelled lipid) of: free fatty acid ((1-¹⁴C) palmitic acid), phosphoglyceride ((dipalmitoyl-1-¹⁴C) phosphatidylcholine), and triacylglyceride ((carboxyl-¹⁴C) triolein). The choice of the lipids was based on the difference in polarity, there-

fore different ways of vehiculization in an aqueous means were used. FFA was administered as ammonium salt according to Laino *et al.* [24], PC was administered as extruded liposomes similar to Garcia *et al.* [34] and TAG by means of sodium cholate (Sigma Chemical Co.) following the methodology used by Cunningham *et al.* [18]. FFA was provided by Amersham, TAG and PC by Perking Elmer. All chemicals were of analytical grade. All appropriate controls were performed and discounted.

After incubation, hemocytes and lipoproteins were isolated as mentioned before. With relation to lipoproteins in each ultracentrifugation aliquot; the presence of proteins was determined through absorbance at 280 nm. Radioactivity of hemocytes and lipoproteins was measured by liquid scintillation in a Wallac 1214 Rack Beta equipment.

3. Results

Figure 1 shows a sample of adult female *Grammostola rosea*.

The absorbance at 280 nm measured in each aliquot after ultracentrifugation of the hemolymph showed a protein profile with two peaks (**Figure 2**). One of them was found in the high-density zone (1.13 - 1.15 g/ml) and was named Gr-HDL fraction, the other was found in the very-high-density zone (1.24 - 1.27 g/ml) and named Gr-VHDL fraction.

Fractions corresponding to Gr-HDL and Gr-VHDL were analyzed separately. The amount of proteins in Gr-HDL and Gr-VHDL was 0.59 and 42.35 mg/ml of the hemolymph respectively. The quantity of lipids transported to each fraction was 0.35 mg/ml of hemolymph for Gr-HDL and 2.96 mg/ml for Gr-VHDL. With relation to the distribution of proteins and lipids in the hemolymph, Gr-VHDL fraction has the greatest % of proteins and lipids (98.6% and 89.3%). The lipid % of lipoproteic fraction was 37.2% in Gr-HDL and 6.9% in Gr-VHDL, respectively. The protein % of lipoproteic fraction was 62.8% for Gr-HDL and 93.1% for Gr-VHDL (**Table 1**).

Figure 3 shows the results obtained by electrophoretic analysis of Gr-HDL and Gr-VHDL under dissociating conditions. HDL displayed a predominant band of 93 kDa and 249 kDa, and Gr-VHDL fraction a predominant band of 68 kDa which would correspond to Hc and two others of 99 and 123 kDa.

With relation to the different lipids found in the lipoproteins, it was observed that Gr-HDL and Gr-VHDL fractions have similar % of PC and PE (approximately 49% and 35% for each lipid). Significant differences be-

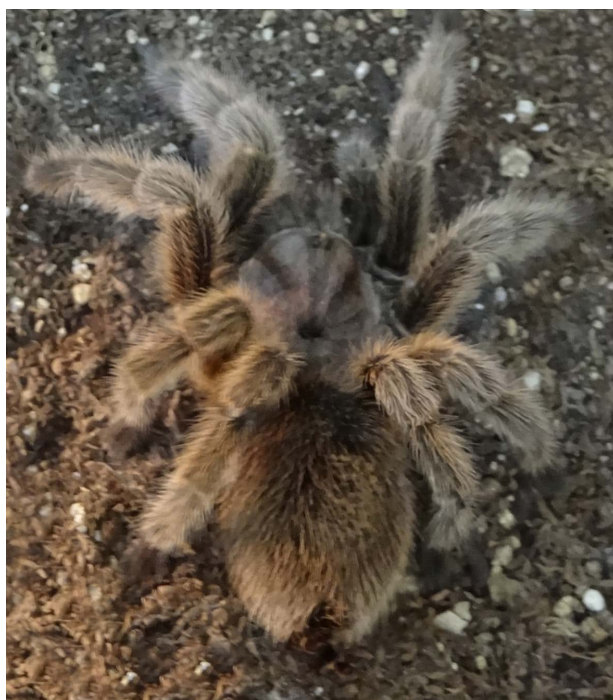


Figure 1. An adult female specimen of *Grammostola rosea* (Araneae, Theraphosidae).

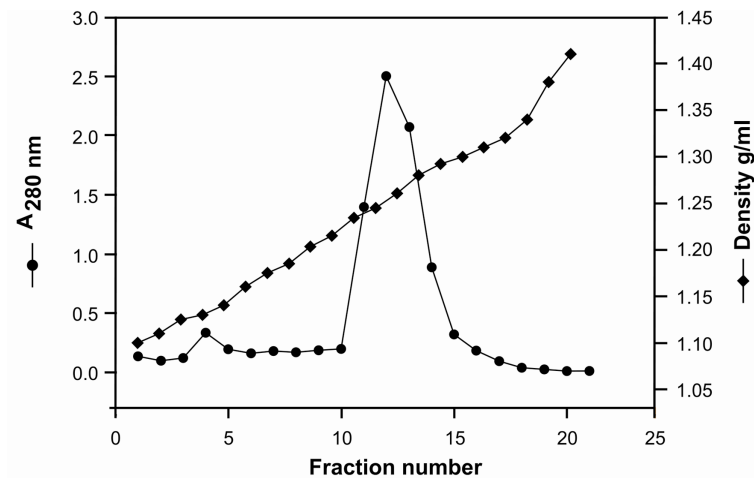


Figure 2. Protein distribution (absorbance at 280 nm) of hemolymph from *G. rosea* obtained by gradient density ultracentrifugation in NaBr 1.35 g/ml.

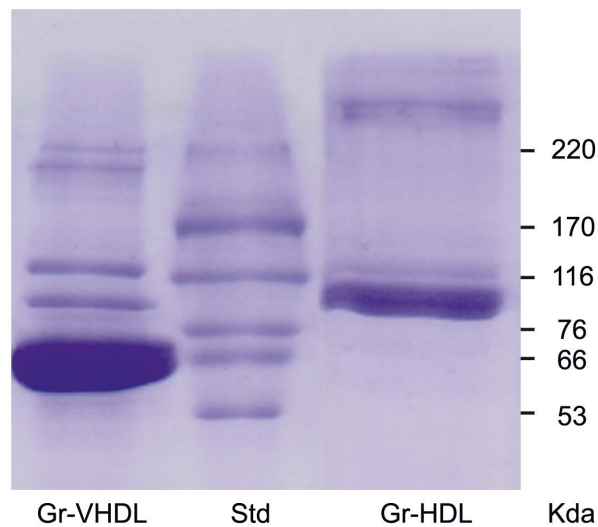


Figure 3. SDS-PAGE analysis of Gr-HDL and Gr-VHDL apoproteins from *G. rosea* hemolymph. The gels were done using polyacrylamide gradient of 4% - 23% wt/vol. The proteins were visualized by Coomassie Brilliant Blue.

Table 1. Quantity and percentage of proteins and lipids in Gr-HDL and Gr-VHDL. Values are the means of three isolations of the hemolymph analyzed separately.

	Gr-HDL	Gr-VHDL
Hydrated density (g/ml)	1.13 - 1.15	1.24 - 1.27
Proteins mg/ml hemolymph	0.59 ± 0.18	42.35 ± 4.6
Circulating proteins %	1.37	98.6
Lipids mg/ml hemolymph	0.35 ± 0.11	2.96 ± 0.32
Circulating lipids %	10.6	89.3
Lipids % of lipoproteic fractions	37.2	6.9
Proteins % of lipoproteic fractions	62.8	93.1
Lipid/Protein ratio	0.59	0.074

tween both fractions were found in minority percentage, thus noticing in Gr-VHDL a lower % of sphingomyelin [SM] and a major % of COL, FFA, TAG and esterified sterol [SE] (Figure 4).

Figure 5 shows a comparative evaluation of the percentage of fatty acids present in Gr-HDL and Gr-VHDL fraction. As the % is similar in both lipoproteins, the predominant fatty acids were in decreasing order: 18:1, 16:0, 18:2, 18:0, 14:0, 16:1 and 18:3, among others.

In vitro assays, FFA, TAG and PC were bound separately to the hemolymph, which was ultracentrifuged in density gradient, and the lipoproteic fractions Gr-HDL and Gr-VHDL were isolated. Radioactivity was measured in each aliquot and the lipoproteins were identified by their absorbance at 280 nm (Figure 6). In this figure it is observed that the radiolabelled TAG and FFA (in these two analyzed concentrations) bind mainly to Gr-VHDL fraction. On the contrary, PC was mainly found in Gr-HDL and in a lesser extent in Gr-VHDL fraction.

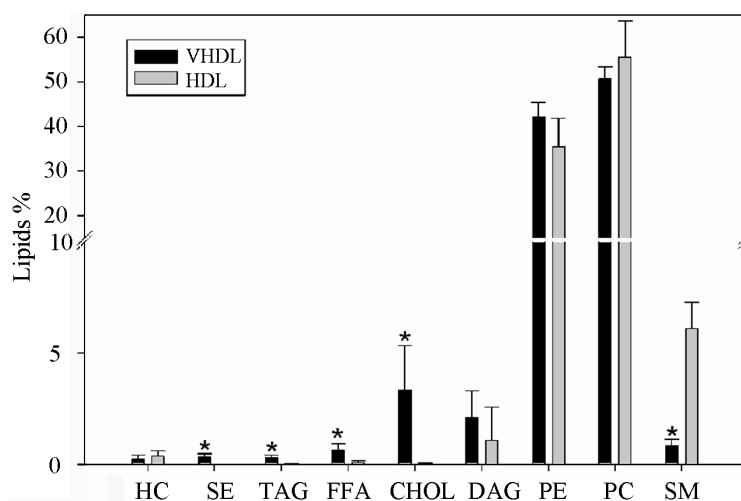


Figure 4. Lipid composition of Gr-HDL and Gr-VHDL from *G. rosea*. Data on lipid classes are expressed as weight percentage as determined and quantified by TLC-FID. Values represent the mean \pm SD of three independent analyses. Student's *t* test was used to compare the significance of the differences between Gr-VHDL and Gr-HDL. **P* < 0.05.

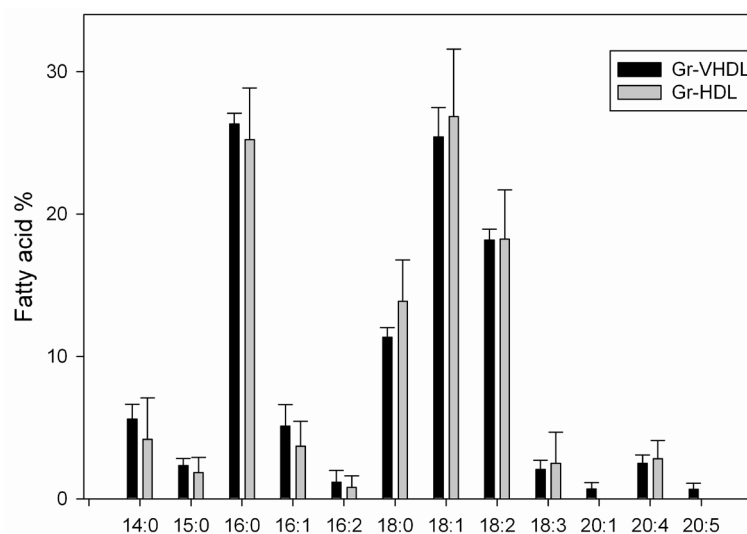


Figure 5. Fatty acid composition of Gr-HDL and Gr-VHDL isolated from hemolymph of *G. rosea*. Data are expressed as weight percentage and quantified by GLC. Values are the mean \pm SD of three independent analyses.

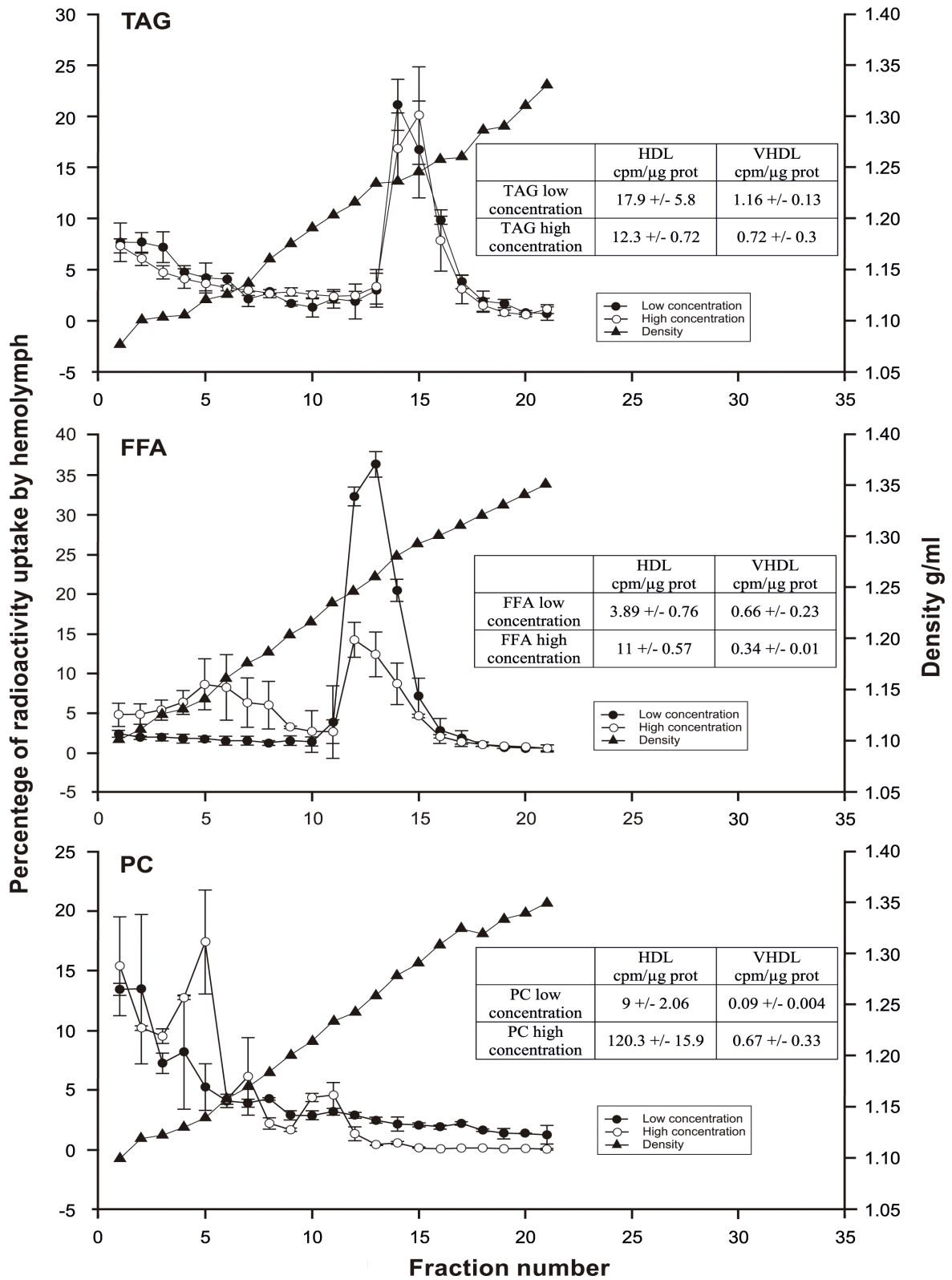


Figure 6. Radioactivity distribution in hemolymphatic fractions of *G. rosea*, after having incubated the hemolymph with different lipids (14 C-FFA, 14 C-TAG and 14 C-PC) and ultracentrifuged in a NaBr gradient. Results correspond to the mean \pm SD of three independent analyses.

Insets show the specific activities of the two concentrations analyzed of the different lipids (TAG, FFA and PC), being majority in Gr-HDL with values ranging from 3.89 to 120 cpm/ μ g of protein and from 0.09 to 1.16 cpm/ μ g of protein for Gr-VHDL.

Table 2 shows the % of radiolabelled lipids, taken up by Gr-HDL, Gr-VHDL and hemocytes in the different assays of *in vitro* incubation. In Gr-HDL it was found about 22% of TAG, 8% and 36% of FFA, and 67% and 78% of PC. In the case of Gr-VHDL it was taken up approximately 75% of TAG, 90% and 62% of FFA, and 30% and 21% of PC. Hemocytes bind from 0.4% to 2.5% of radioactivity associated to the different lipids assayed.

4. Discussion

In the hemolymph of the tarantula *G. rosea* two lipoproteic fractions with densities corresponding to HDL and VHDL were identified and defined as Gr-HDL and Gr-VHDL.

Gr-HDL has a density (1.13 - 1.15 g/ml) similar to that reported for the HDL-1 of the spider *L. mirabilis* (1.13 g/ml) [12]. For the case of spiders *P. pythagoricus* and *Aphonopelma hentzi* in which two HDLs were reported, Gr-HDL coincides with *P. pythagoricus* HDL-1 (1.13 g/ml) [11] and with one of the *Aphonopelma hentzi* HDL (1.14 g/ml) [16].

Molecular weights of apolipoproteins (93 and 249 kDa) related to their electrophoretic mobility of Gr-HDL are similar to the apolipoproteins reported for the tarantula *Aphonopelma hentzi* and for the spider *P. pythagoricus* (80 and 250 kDa) [11] [15]. The term lipophorin proposed by Chino *et al.* [35] to designate insect lipoproteins with densities near 1.14 g/ml, capable of transporting majority DAG and which was previously applied to tarantula HDL [15] seems to be adequate for the present Gr-HDL because the presence of two apolipoproteins of 250 - 280 and 75 - 78 is characteristic of lipophorin [36].

The electrophoretic analysis of fraction VHDL under dissociating conditions shows one predominant protein band of 68 kDa and two minority apolipoproteins of 99 and 123 kDa. The band of 68 kDa has a molecular weight similar to the Hc monomers found in other spiders [14] [37]-[39]. Whereas the other two minority bands have molecular weights similar to the apolipoproteins described for VHDL of *P. pythagoricus* and *Aphonopelma hentzi* (105 and 121 kDa) [11] [15]. It is important to highlight that, as in the spider *L. mirabilis* [12], VHDL fraction showed three peaks with the same proteic compounds when subjected to chromatography of molecular exclusion, which was determined in the SDS-PAGE (data not shown).

The quantity of lipids associated with Gr-HDL is 37.2% which coincides with that reported for the tarantula *Aphonopelma hentzi* 40/44% [15], but it is somewhat elevated in relation to that found in other Theraphosidae spiders as in *L. mirabilis* with 20% [12] and in *P. pythagoricus* with 28% [11]. With relation to the quantity of lipids found in Gr-VHDL fraction (6.9%), it is similar to that described for *P. pythagoricus* (3%) and rather higher for *L. mirabilis* (1%) [12] [14].

At present, taking into account the quantity of works performed in this subject, (see review Cunningham *et al.*, [13]) it is not surprising to find a lipoproteic fraction of high (HDL) or very high (VHDL) density with Hc, which was firstly described in the arachnids using *P. pythagoricus* (Sparassidae) by Cunningham and Pollero

Table 2. Percentage of the different lipids (14 C-FFA, 14 C-TAG and 14 C-PC) taken up by lipoproteic fractions after incubating *in vitro* for 1 hour, in 50 mM potassium phosphate buffer pH 7.4.

	Gr-HDL	Gr-VHDL	Hemocytes
% 14 C-TAG (low concentration)	22.40 \pm 2.53	76.55 \pm 2.48	1.04 \pm 0.05
% 14 C-TAG (high concentration)	23.05 \pm 2.63	75.12 \pm 3.18	1.81 \pm 0.57
% 14 C-FFA (low concentration)	8.11 \pm 1.78	90.5 \pm 1.63	1.37 \pm 0.27
% 14 C-FFA (high concentration)	36.0 \pm 2.08	62.15 \pm 2.25	1.74 \pm 0.18
% 14 C-PC (low concentration)	67.17 \pm 10.47	30.28 \pm 9.16	2.54 \pm 1.33
% 14 C-PC (high concentration)	78.58 \pm 7.61	21.01 \pm 7.66	0.40 \pm 0.05

[14], and subsequently it was confirmed the presence of lipids in the lipoproteic fraction with Hc in *L. mirabilis* (Theridiidae) [12]. Using different radiolabelled lipids and *in vivo* and *in vitro* assays, the functionality of lipoproteic fractions with Hc in *P. pythagoricus* was studied throughout the last years by Cunningham *et al.* [18] [40] and Laino *et al.* [24] [25].

In the present work we are not able to find an explanation to the fact that there is a presumably different lipid transport system in a same arachnid family (Theraphosidae). Haunerland and Bowers [15] (using ultracentrifugation in a density gradient with NaBr 1.35 g/ml) describe that there are three proteic fractions of different density in the hemolymph of the tarantula *Aphonopelma hentzi*: HDL of 1.12 g/ml, VHDL of 1.28 g/ml and Hc with other proteins > 1.28 g/ml. In this particular case with the tarantula *G. rosea*, employing a gradient of similar density (NaBr 1.35 g/ml), the Gr-VHDL fraction with Hc showed a density of 1.24 - 1.27 g/ml, and proteins with higher density are not observed. This fact may be possible because Hc interacts with lipids, thus “floating” is achieved in that density. Therefore, Hc functions as lipid transport in arachnids does not seem to be general, since only in three out of the four studied species this function is observed. This discrepancy is not exclusive for arachnids as it was also observed in other invertebrates such as mollusks. In cephalopods Hc transports lipids [41], but in gastropods this phenomenon does not occur [42]. Numerous studies with new models are necessary in order to interpret these observations.

PC is generally in arthropod lipoproteins the majority structural lipid, which can be found in crustacean lipoproteins with 38% of total lipids in *M. borellii* and 46% in *P. leniusculus* [7] [43], and in arachnids lipoproteins with 31% and 25% in HDL-1 and HDL-2 fractions of *L. mirabilis*, 59% in VHDL fraction of *Aphonopelma hentzi* and with 30%, 22% and 40% in HDL-1, HDL-2 and VHDL of *P. pythagoricus* [11] [12] [14] [16]. In the present study the quantity of PC is rather greater, finding values of 55% and 49% in Gr-HDL fraction and Gr-VHDL respectively. PE% in *G. rosea* (35% - 42%) is also greater to that described for arachnid lipoproteins (3.6% - 15.2%) [11] [12] [14] [16].

The present study is the first report that observes the presence of SM on arachnid lipoproteic fractions. It is likely that this lipid is found in other arachnid species where lipid characterization in lipoproteic fractions has been reported, but within the so-called other PL as the case of HDL-1 in *P. pythagoricus* (16%) [11], HDL-2 (17.7%) and VHDL (11.6%) [14], or lipids unidentified in VHDL of *Aphonopelma hentzi* (10%) [16]. Comparing the different lipids found in the two lipoproteic fractions, it is observed that Gr-HDL has greater percentage of SM than Gr-VHDL, and mainly greater % of CHOL and lesser quantities of FFA and SE. With relation to CHOL it is important to highlight that Gr-VHDL possesses 100 more times CHOL in its structure than Gr-HDL, this difference coincides with that described by Cunningham *et al.* [11] [14] where it is observed that VHDL fraction (lipoproteic fraction with Hc) has CHOL in its structure, and being absent in HDL-1 (lipoproteic fraction without Hc). Moreover, VHDL of *P. pythagoricus* has the ability to bind (*in vitro*) CHOL 1070 more times than it normally does [18]. This pattern is also repeated when with-and-without-Hc HDL fractions are compared in *L. mirabilis*, thus finding CHOL 4.7 more times in the HDL with Hc. In 2000 Cunningham and collaborators suggest that lipoproteic fraction with Hc could be specialized in CHOL transport. Although available data are scarce since there are few studies on the subject, it is possible that Hc has different lipid binding or as mentioned by Cunningham *et al.* [12] Hc aggregates give rise to particles of different conformation, which are able to bind certain lipids.

The fatty acid compositions of fraction Gr-HDL and Gr-VHDL determined by GLC are shown comparatively in Figure 6. Here it is observed that both fractions have similar composition of fatty acids being majority fatty acids 18:1, 16:0, 18:2, 18:0 and 20:4. This coincides with that analyzed by Laino and collaborators in VHDL hexamer of *P. pythagoricus* (data not published), by Stratakis *et al.* [16] in VHDL of tarantula *Aphonopelma hentzi* 18:1 + 18:2 (52.3%), 16:0 (24.9%), and also in egg lipoproteins of the spiders *P. pythagoricus* and *Schizocosa malitiosa* with 18:1 (31% and 24%), 18:2 (26% and 25%), 16:0 (18% and 20%) [44] [45]. It is important to point out that also this pattern with 18:1 (30%), 18:2 (23%) and 16:0 (15%) is also found in the biosynthetic organ of spider lipids (midgut diverticula of *P. pythagoricus*) [24]. Although it is described that the different dietary habits of arthropods can generate differences in the fatty acid composition of hemolymph [46], in the present work the comparison of the fatty acids of the lipoproteic fractions with the scarce existing data on hemolymph of *Aphonopelma hentzi* and *P. pythagoricus* leads us to think that either the dietary changes are very few or the present hypothesis does not seem to apply for arachnid lipoproteins.

Binding assays of different lipids to the hemolymphatic components demanded the adaptation of appropriate techniques for the three different lipid bindings. For the binding of TAG, the addition of an emulsifier such as

sodium cholate and optimum incubation conditions were required; for PC, extruded liposomes were necessary; and ammonium salts were performed for FFA. As a consequence of these experiments it was observed that hemocytes participate in the uptake of different lipids (assays of *in vitro* incubation), the % of taken-up lipids is similar to that previously observed in the hemocytes of the spider *P. pythagoricus* [25]. These results suggest that lipids of different polarities are taken up approximately in the same proportion by hemocytes 0.4% - 2.5%, opposed to that observed by Pollero *et al.* [47] where triolein ¹⁴C and palmitic acid ¹⁴C are taken up by mollusk hemocytes, 12% and 30% respectively.

On the other hand, it was observed that Gr-HDL takes up more ¹⁴C-PC than Gr-VHDL coinciding with *in vitro* experiments performed in *P. pythagoricus* with HDL lipoprotein and VHDL analyzed separately [25].

On the contrary, the greatest percentage of radiolabelled FFA and TAG is associated to Gr-VHDL lipoproteic fraction with Hc as apolipoprotein. Cunningham *et al.* [18] with the study of lipid binding capacity of spider Hc explain the possible presence of hydrophobic zones in the protein that would generate zones of lipid binding. This may possibly occur in VHDL fraction of *G. rosea*. The greatest % of TAG and FFA found in Gr-VHDL coincides with this observation. Moreover, despite the higher specific activities of FFA and TAG in Gr-HDL compared to Gr-VHDL fraction, the total amount of proteins, FFA, and TAG remains higher in Gr-VHDL, providing a greater FFA and TAG uptake.

In conclusion, the present study reports the lipid characterization of lipoproteic fractions found in the hemolymph of tarantula *G. rosea*. The role of each lipoproteic fraction in relation to lipid uptake is proven by *in vitro* assays. Similarities and differences according to its lipid and protein compositions are found when they are compared with the lipoproteins of the only three species of spiders studied.

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