Serum proteins and fractions, HDL-cholesterol and total IgG and IgE levels in cases of acute and chronic paracoccidioidomycosis

Proteínas séricas e frações, HDL-colesterol e níveis de IgG e IgE totais na paracoccidioidomicose aguda e crônica

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ABSTRACT
This study evaluated serum protein fractions, HDL-cholesterol, total immunoglobulin G and total immunoglobulin E levels in patients with acute and chronic paracoccidioidomycosis, by means of electrophoresis, enzymatic reaction and immunoenzymatic assay. The results demonstrated elevated levels of total immunoglobulin G, total immunoglobulin E, alpha-2 and gamma-globulins, which were more evident in acute than in chronic PCM, but no increase in HDL-cholesterol levels. There was a correlation between the levels of total immunoglobulin E and gamma-globulins and the alpha-2 and beta-globulin fractions in the acute form and between beta and gamma-globulins in both the acute and the chronic form. In conclusion, changes in total immunoglobulin G and immunoglobulin E levels and in the electrophoretic profile may be important markers for the prognosis and therapeutic follow-up of PCM cases, especially because protein electrophoresis is a simple laboratory test that can be applied when specific PCM serological tests are not available. In addition, levels of the gamma-globulin fraction greater than 2.0g/dL may suggest that the patient is developing a more severe form of PCM.


RESUMO
Este trabalho avaliou as frações de proteínas séricas, HDL-colesterol e imunoglobulina G e imunoglobulina E totais em pacientes com paracoccidioidomicose aguda e crônica por eletroforese, reação enzimática e ensaio imunoenzimático. Os resultados demonstraram aumento dos níveis de imunoglobulina G e imunoglobulina E totais, alfa-2 e gama-globulinas, mais evidente na forma aguda que na forma crônica, e não nos níveis de HDL-colesterol. Houve correlação entre níveis de imunoglobulina E total e gama-globulinas e fração alfa-2 e beta-globulinas na forma aguda e entre beta e gama-globulinas nas formas aguda e crônica. Concluindo, alterações nos níveis de imunoglobulina G e imunoglobulina E totais e no perfil eletroforético podem ser importantes marcadores para prognóstico e acompanhamento terapêutico da PCM, especialmente por ser a eletroforese de proteínas um exame laboratorial simples que pode ser empregado em situações onde a sorologia específica para PCM não está disponível. Adicionalmente, níveis da fração gama-globulinas acima de 2,0g/dL podem sugerir que o paciente esteja desenvolvendo uma forma mais grave de PCM.


Paracoccidioidomycosis (PCM) is a systemic granulomatous disease caused by the dimorphic fungus Paracoccidioides brasiliensis. It is considered to be one of the most common systemic mycoses in Latin America, and Brazil has the greatest number of cases. PCM disease presents two clinically distinct forms: the acute or juvenile form (AF) and chronic or adult form (CF). The AF is characterized by rapid clinical evolution with the involvement of multiple organs, while the CF, which accounts for more than 90% of the patients, progresses slowly and silently for years. The multifocal CF is more severe and involves skin, mucosa, lungs and lymph node manifestations.
Serum protein electrophoresis is a simple laboratory test that helps to monitor patients over the course of PCM. It has already been demonstrated that, although total serum proteins may persist at normal levels, the fractionation of the serum shows a decrease in albumin levels and an increase in alpha-1, alpha-2 and gamma-globulin fractions. The variations in these parameters are also more prominent in patients with the acute form than in those with the chronic form of the disease.

Analysis of antibody isotype expression in AF and CF PCM has demonstrated elevated levels of total IgG and IgE and of IgG, IgE or IgA specific to some Paracoccidioides brasiliensis antigens\(^{5,16}\), while in the AF it occurs mainly by the presence of total IgG, IgG\(_4\) subclass and IgE\(^{2,31}\). High levels of total IgE and IgE anti-gp43 have been correlated with polyclonal B lymphocyte activation, which is observed in the AF disease\(^8\). Cellulose-acetate electrophoresis is a simple and routine clinical laboratory procedure that could be used to improve the interpretation and evaluation of PCM patients’ follow-up, taking into account the more recent immunological knowledge of this disease. In addition, since a large proportion of \(\alpha\)-lipoproteins, represented mainly by HDL-cholesterol, migrate into the alpha-1 globulin region and are a major contributor to this protein fraction according to capillary zone electrophoresis\(^{15,18}\), this parameter has also been estimated.

The present study evaluated the levels and possible correlations among serum protein fractions, HDL-cholesterol and total IgG and IgE in acute and chronic PCM patients.

**MATERIAL AND METHODS**

**Serum samples.** Serum samples from 30 chronic PCM patients (ranging from 33 to 82 years of age; all females) with unifocal or multifocal disease, who were attended at Londrina State University Clinical Hospital (Londrina, PR, Brazil), and 12 acute PCM patients (ranging from 11 to 23 years of age; seven females and five males) from the Mycosis Immunodiagnostic Laboratory, Immunology Section, Adolfo Lutz Institute, S\(_\alpha\)o Paulo, SP, Brazil, were used. All of the PCM patients had shown positive results in radial immune diffusion tests. The control group (NHS) consisted of 44 serum samples from healthy male and female adult individuals who had already been used as normal controls at the laboratory; all of them were negative for PCM in the IDR and ELISA tests. Informed consent was obtained from all subjects participating in this study, and the study had previously been approved by the Internal Scientific Commission and Research Bioethics Committee of Londrina State University (Londrina, PR, Brazil) and by the Research Ethics Committee of the Adolfo Lutz Institute (CETIAP).

**Enzyme-linked immunoSorbent assay for total IgE.** ELISA immunoplates were sensitized (100\(\mu\)l/well) with the immunoglobulin fraction goat anti-human IgE (1-O632 Sigma Chemical Co, St. Louis, MO, USA) at 1\(\mu\)g/ml in carbonate buffer (pH 9.6), for 1h at 37\(\circ\)C and then overnight at 4\(\circ\)C. After blocking with 5% skimmed milk in PBS and incubated with serum samples (1:4) in 0.5% skimmed milk in PBS for 2h at 37\(\circ\)C, followed by washing and incubation with goat alkaline-phosphatase anti-human IgE (A-3525 Sigma Chemical Co, St. Louis, MO, USA) diluted 1:500 (100\(\mu\)l/well) for 1h 30 min at 37\(\circ\)C. The reaction was developed using p-nitrophenyl phosphate in diethanolamine buffer (mass/volume), at pH 9.8 (100\(\mu\)l/well), with 30 min of incubation. The reaction was stopped with 50\(\mu\)l/well of 3 \(\mu\)M NaOH. The absorbance was read at 405nm in a Multiskan EX reader (Labsystems, Helsinki, Finland). No significant background was observed in the reaction controls. A standard curve was produced using different dilutions of a pool of serum samples of known concentration (ng/ml).

**High-density lipoprotein cholesterol quantification.** Serum samples were analyzed using a commercial kit (Ebram Laboratory Products Ltda), by means of a direct ultra-sensitive colorimetric reaction with cholesterol oxidase in automated equipment (Selectra E, Bayer SA).

**Serum protein electrophoresis.** Serum samples were applied to cellulose acetate strips and electrophoresis was performed in Veronal-Sodic buffer (pH 8.6-8.8) for 25 minutes, followed by 10 min of incubation in Ponceau S stain and destaining in 5% acetic acid. All the strips were dehydrated for 1 min in 5% acetic acid. They were then dried at 80\(\circ\)C, followed by washing and incubation with serum samples (1:4) in 0.5% skimmed milk in PBS for 2h at 37\(\circ\)C, followed by washing and incubation with goat anti-human IgG peroxidase conjugate (A-8775 Sigma Chemical Co, St. Louis, MO, USA) diluted 1:4000 (100\(\mu\)l/well) for 1h 30 min at 37\(\circ\)C, followed by washing and the addition of 10mg of OPD (ortho-phenylenediamine) in 25ml of phosphate citrate buffer (pH 5.0), plus 10\(\mu\)l of 30\% \(H_2O_2\) (100\(\mu\)l/well). After 15 min of incubation, the reaction was stopped with 50\(\mu\)l/well of 4\(N\) \(H_2SO_4\) and the absorbance was read at 492nm in a Multiskan EX reader (Labsystems, Helsinki, Finland). The background interference (evaluated through reaction controls) was discounted from all absorbance results. A standard curve was performed using different dilutions of a pool of serum samples of known concentration (mg/dl).

**Radial Immunodiffusion Test.** The serum samples were analyzed in accordance with the protocol used by Tatibana et al\(^3\).

**Statistical analysis.** For the statistical analysis, all the variables were calculated as log\(_10\) values, except for total IgE, which was calculated as SQRT values. Pearson’s correlation test was applied. Significance was defined as \(p \leq 0.05\) and positive correlations as \(r \geq 0.50\). The correlations were taken to be strong when \(r \geq 0.75\) and weak when \(r > 0.50\) and < 0.75.
globulin fraction analysis among the groups, a box-plot graph was also performed using the Bioestat statistical software.

RESULTS

The ELISA results for total IgE showed significant differences between all of the groups \((p<0.05)\). For total IgG, the results were significantly higher in AF PCM than in CF PCM or NHS. The statistical analysis did not demonstrate any significant difference in total IgG between the CF PCM and NHS groups (Table 1). The mean values for HDL in AF PCM, CF PCM and NHS were \(28.2 \pm 7.89\, \text{mg/dL}, \, 29.6 \pm 10.45\, \text{mg/dL} \) and \(30.8 \pm 8.52\, \text{mg/dL}\), respectively. In PCM patients, the mean results for HDL-cholesterol were slightly below the normal reference limits (30 to 70 mg/dL). No correlation between HDL and serum protein fractions was observed in this study.

The protein fraction results expressed in g/dL are summarized in Table 2. These show that the mean values for alpha-2 and gamma-globulins are elevated in CF PCM, and that total proteins and alpha-1, alpha-2 and gamma-globulins are elevated in AF PCM. The albumin/globulin ratio was lower in AF PCM than in CF PCM and NHS. There was no correlation between albumin and IDR results \((r = -0.1031)\).

For AF PCM, Pearson’s correlation test demonstrated stronger coefficients between alpha-1 and beta-globulins \((r = 0.8307)\) and between alpha-1 and gamma-globulins \((r = 0.8289)\), but this was not observed in the CF PCM or NHS groups (Figures 1C and 1B). This demonstrated a weak correlation between the beta and gamma-globulin fractions \((r = 0.7142)\) and between gamma-globulins and total IgE \((r = 0.6026)\) (Figures 1A and 1D). In CF PCM, only a weak correlation between beta and gamma-globulin fractions was detected \((r = 0.6506)\).

**TABLE 1**

<table>
<thead>
<tr>
<th>Group (n)</th>
<th>Total IgG</th>
<th>Total IgE</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>OD mg/dL</td>
<td>OD ng/dL</td>
</tr>
<tr>
<td>(1) CF PCM (30)</td>
<td>0.298 ± 0.210(^a)</td>
<td>440 ± 439(^a)</td>
</tr>
<tr>
<td>(2) AF PCM (12)</td>
<td>0.424 ± 0.122(^b)</td>
<td>1704 ± 255(^b)</td>
</tr>
<tr>
<td>(3) NHS (44)</td>
<td>0.243 ± 0.086(^c)</td>
<td>1325 ± 179(^c)</td>
</tr>
</tbody>
</table>

\(^a\)Total IgG 1x2, 2x5 and \(^b\)Total IgG 1x3, 2x5, 1x2, \(p<0.050\).

(n) number of patients or healthy individuals.

Reference values: total IgG \((751 \text{ to } 1,560\, \text{mg/dL in adults})\); total IgE \((> 33.6\, \text{ng/dL for individuals above 10 years old})\).


**TABLE 2**

<table>
<thead>
<tr>
<th></th>
<th>(1) CF PCM (n=30)</th>
<th>(2) AF PCM (n=12)</th>
<th>(3) NHS (n=44)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>g/dL %</td>
<td>g/dL %</td>
<td>g/dL %</td>
</tr>
<tr>
<td>Total proteins</td>
<td>7.60 ± 1.303(^a)</td>
<td>8.64 ± 1.228(^a)</td>
<td>7.43 ± 1.275(^a)</td>
</tr>
<tr>
<td>Albumin</td>
<td>3.64 ± 0.779(^a)</td>
<td>4.79 ± 6.666(^b)</td>
<td>4.28 ± 0.876(^c)</td>
</tr>
<tr>
<td>α1-globulins</td>
<td>0.31 ± 0.115(^a)</td>
<td>4.11 ± 1.466(^a)</td>
<td>0.28 ± 0.113(^b)</td>
</tr>
<tr>
<td>α2-globulins</td>
<td>1.03 ± 0.258(^a)</td>
<td>13.62 ± 2.525(^b)</td>
<td>3.80 ± 1.314(^c)</td>
</tr>
<tr>
<td>β-globulins</td>
<td>0.92 ± 0.296(^a)</td>
<td>12.06 ± 2.766(^a)</td>
<td>10.01 ± 2.035(^b)</td>
</tr>
<tr>
<td>Δ-globulins</td>
<td>1.70 ± 0.492(^a)</td>
<td>22.26 ± 4.166(^a)</td>
<td>18.54 ± 1.366(^c)</td>
</tr>
<tr>
<td>Albumin/globulin</td>
<td>0.919(^a)</td>
<td>0.864(^a)</td>
<td>1.559(^a)</td>
</tr>
</tbody>
</table>

Significant horizontal comparisons between groups (1) CF PCM, (2) AF PCM and (3) NHS are expressed as different vowels \((p<0.050)\), while the same vowels represent results not statistically different.

(n) Number of patients or healthy individuals.

Reference values from literature for each variable are: total proteins \((6.0 - 8.0\, \text{g/dL})\), albumin \((3.5 - 4.9\, \text{g/dL}; \, 50 \pm 68\%)\), α1-globulins \((0.1 - 0.4\, \text{g/dL}; \, 2 \pm 6\%)\), α2-globulins \((0.4 - 0.8\, \text{g/dL}; \, 7 \pm 12\%)\), β-globulins \((0.6 - 1.0\, \text{g/dL}; \, 9 \pm 16\%)\) and Δ-globulins \((0.8 - 1.4\, \text{g/dL}; \, 12 \pm 22\%)\).

Correlations between globulins and total immunoglobulin E, evaluated 2x2, in paracoccidioidomycosis patients with the acute/subacute (AF) or the chronic (CF) form and in healthy controls (NHS). (A) Gamma and beta-globulins in AF PCM (r = 0.7142), CF PCM (r = 0.6506) and NHS (r = 0.3369); (B) Gamma and alpha-1 globulins in AF PCM (r = 0.8289), CF PCM (r = 0.2685) and NHS (r = 0.2790); (C) Beta and alpha-1 globulins in AF PCM (r = 0.8306), CF PCM (r = 0.2303) and NHS (r = 0.3521); (D) Total IgE and gamma-globulins in AF PCM (r = 0.6026), CF PCM (r = 0.0284) and NHS (r = 0.0426).

**DISCUSSION**

Most patients with active PCM demonstrate protein abnormalities that are revealed by a simple serum protein electrophoresis test\(^1\). This test is usually used as an initial evaluation marker for patients and to follow up the treatment evolution\(^13\ 20\). Studies on serum protein profiles for PCM patients have shown decreased albumin and increased alpha-1, alpha-2 and gamma-globulin fractions, along with reduced beta-lipoproteins in some patients, and have correlated these variations with PCM severity\(^13\ 17\). In the present study, increased levels of alpha-2 and gamma-globulins in AF PCM and alpha-1, alpha-2 and gamma-globulins in CF PCM were described. In order to establish a cutoff value for gamma-globulins that could indicate disease severity, data analysis was performed in the Bioestat 5.0 statistical software, which revealed that every CF PCM patient had results below 2.0g/dl, while the mean value for AF PCM patients was 2.08g/dl. Although clinical characteristics, nutritional state and other comorbidities were not appraised in this investigation, it may be suggested through taking into account that AF PCM is considered more severe than CF PCM\(^6\ 23\), that patients with gamma-globulin laboratory results above 2.0 g/dl are probably developing a more severe condition of the disease that requires further investigation. The albumin/globulin ratio was also significantly different between AF or CF PCM and NHS, as expected and described in the literature\(^12\ 21\). The reduced albumin values observed in percentages but not in absolute results, except for CF PCM patients exhibiting high titers in the IDR test (1:16 and 1:32), could be correlated with a more prominent inflammatory response and greater severity of disease state in these patients. Under other conditions, it has been proposed that the main reason for reduced albumin levels relates to synthesis of acute-phase proteins in the liver, which occurs at the expense of albumin production, and that the albumin/globulin ratio is a better parameter for distinguishing between malignant and healthy states\(^1 12\ 21\).

Considering that elevated alpha-1 globulin levels were observed in the AF PCM and not the CF PCM group, we investigated the possible role of HDL in the elevation of this parameter. Our results showed that there was no significant difference in HDL levels between the AF PCM, CF PCM and NHS groups, thus demonstrating that HDL cholesterol is not expected to influence elevation of the alpha-1 globulin fraction in PCM.

The use of serological methods to detect specific antibodies is an important tool in PCM diagnosis, but the problem of antigen standardization still remains. This makes it difficult to compare the results from different laboratories and means that ensuring access to these methodologies for every clinical diagnosis laboratory is problematic\(^8\ 11\ 19\). In this study, higher levels of total IgG and IgE could be seen, but there was only a significant correlation between total IgE and the gamma-globulin fraction in the AF of the disease. This suggests that increased IgE levels could be a better molecular marker for severity than IgG in PCM is.
From these results, it could be confirmed that the serum protein fraction profile changes in PCM, and that there are positive correlations between some of the protein fractions and between gamma-globulin and total IgE levels. It could also be concluded that there are differences in interpreting albumin results when expressed as percentages or absolute levels, and that HDL probably does not contribute towards the changes in the alpha-1 globulin fraction that are observed among PCM patients. In addition, levels of the gamma-globulin fraction above 2.0g/dl may suggest that the patient is developing a more severe form of PCM. Changes in both the serum electrophoretic profile and the total IgE and IgE levels may be important markers for prognostic or even treatment follow-up in PCM. Electrophoresis may be especially important, since it is a simple routine laboratory method that can be applied in regions where specific PCM serological tests are not easily available.

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REFERENCES


