Improved Immunodiagnosis of Cystic Hydatid Disease by Using a Synthetic Peptide with Higher Diagnostic Value Than That of Its Parent Protein, *Echinococcus granulosus* Antigen B

GUALBERTO GONZÁLEZ-SAPIENZA,* CARMEN LORENZO, AND ALBERTO NIETO

Cátedra de Inmunología, Facultad de Química, Universidad de la República, Montevideo, Uruguay

Received 5 April 2000/Returned for modification 22 June 2000/Accepted 23 August 2000

The assays are used for the diagnosis of hydatid disease are still imperfect. The reported diagnostic sensitivity and specificity vary greatly depending on the panel of sera used, the laboratory conducting the assay, and, more critically, the antigen used. To contribute to its standardization, we have recently ranked the diagnostic performances of the major parasite antigens and the available synthetic peptides using a large collection of serum samples. That work showed that antigen B (AgB) possesses the highest diagnostic value among these antigens. In the present work we further dissected its antigenicity by analyzing the reactivity of the same panel of sera against a set of synthetic peptides spanning the sequence of both AgB subunits. The N-terminal extension of these subunits appeared to be immunodominant in human infections. A 38-mer peptide (p176) delineated from the N-terminal extension of the AbB1 subunit performed in an enzyme-linked immunosorbent assay with a higher diagnostic sensitivity (80%) and specificity (94%) than native AgB, Ag5, or any other peptide antigen tested against this collection of serum samples. In view of its high diagnostic value and its nature as a well-defined reproducible antigen, p176 could conveniently be used as a reference standard antigen in the diagnosis of hydatid disease.

Cystic hydatidosis, caused by infection with larval *Echinococcus granulosus*, affects both humans and domestic animals and is recognized as one of the world's major zoonoses (15). Clinical diagnosis of the disease is based on symptomatology, epidemiological data, imaging techniques, and immunodiagnosis (1). However, clinical symptoms do not appear until the larval cyst structure of the parasite has reached a certain size, which normally requires years after the primary infection. During this time, immunodiagnosis has also proved effective and can conveniently be used to diagnose the disease in asymptomatic high-risk populations (2, 5). This is particularly important for early diagnosis, which is of great significance, because surgery and chemotherapy are poorly effective if they are applied to patients with advanced infections.

The need for a reliable immunodiagnostic test has prompted abundant research in the field; various tests have been used for detection of specific antibodies, but the key component of the different systems remains the antigen. From the beginning it was evident that the transition from crude preparations to purified fractions improved the sensitivity and specificity of the assays. In this process, two major antigens of hydatid cyst fluid, namely, antigen B (AgB) and Ag5, were identified (17). AgB is a 120-kDa oligomeric lipoprotein composed of multiples of two subunits of 8 kDa (9), namely, AgB8/1 (8) and AgB8/2 (7). Ag5 is a high-molecular-mass lipoprotein complex composed of 57- and 67-kDa components that under reducing conditions dissociate into 38- and 22- to 24-kDa subunits (13). Although both antigens proved to be diagnostically valuable, there are difficulties related to their lack of sensitivity and specificity and problems with the standardization of their use (3). In order to overcome these difficulties, efforts have been made to define discrete epitopes of these antigens that could be mimicked by synthetic peptides. The first application of such peptides was reported by Chamekh et al. (4), who used a 34-mer peptide (p89-122), which was suggested to be a major epitope of Ag5 but which was recently shown to be a fragment of a 29-kDa protoscolex component of *E. granulosus* (10). This was followed by the work of Leggatt and McManus (11), who explored the antigenicity of AgB with three peptides and found that p65, a 27-mer peptide corresponding to residues 12 to 39 of AgB8/1, had potential for use as a diagnostic reagent. In an effort to contribute to the standardization of the immunodiagnosis of hydatid disease, we recently compared the diagnostic value of p89-122, p65, native AgB and Ag5, and Gu4, a 34-mer synthetic peptide corresponding to the C-terminal end of the AgB8/2 subunit (3). This work stressed the relevance of an internal comparison of the different antigens in one laboratory in order to rank their diagnostic performance in a reliable manner and showed that (i) individually considered, native AgB has the highest diagnostic value among these antigens and (ii) the available AgB-derived peptides do not reproduce all the major epitopes of AgB.

In the present work we complete the characterization of the antigenicities of both subunits of AgB by introducing three additional synthetic peptides. The study allowed the identification of a highly antigenic region of AgB residing in the N-terminal extension of the AgB8/1 subunit. An enzyme-linked immunosorbent assay (ELISA) based on the use of a single peptide representing this region exhibited a diagnostic performance that was superior to that obtained by the use of native AgB, and the peptide constitutes a promising candidate for standardization of the serodiagnosis of human cystic echinococcosis.

**MATERIALS AND METHODS**

Human serum sample collection. Sera from 90 patients with surgically confirmed hydatid disease were tested. The samples were not preselected on the basis of previous serologic information and were collected before surgery. Sixty-five of them had records of cyst location, which were as follows: liver (*n* = 40), lungs (*n* = 11), bones (*n* = 8), and multiple sites (*n* = 6). In order to evaluate the specificities of the various antigens, 86 serum samples from patients with the following diseases were included in the study: alveolar hydatid disease (*n* = 27),...
**RESULTS**

The amino acid sequences of the peptides introduced in this study, the amino acid sequences of the previously described peptides, and their alignments with the subunits of AgB are depicted in Fig. 1. The peptides were designed in order to complete the available information on the antigenicity of AgB. p176 covers the N-terminal extension of AgB8/1 (not included in the study of Leggatt and McManus [11]), while p175 and p177 represent the N-terminal and central regions of AgB8/2, respectively, and are therefore complementary to Gu4.

The reactivity of our panel of sera with these peptides and AgB was analyzed by ELISA (total IgG) and is shown in Fig. 2 and Table 1. The peptides exhibited important differences in their diagnostic sensitivities. While peptides p175 and p177 detected only a limited number of confirmed cases of the disease (diagnostic sensitivities of 49 and 38%, respectively), the diagnostic sensitivity accomplished with p176 was 80%, similar to that of AgB (77%). The number of serum samples classified as positive by using p176 was also markedly superior to that obtained with the previously described peptides p65 and Gu4. The correlation between location and seropositivity was similar to what has been reported before for other antigens (3).

All peptides showed a high diagnostic specificity, which, in all cases, was superior to that of AgB. Detailed information about the disease groups that gave place to the limited cases of cross-reactivity is displayed in Table 2. While the cross-reactivity observed with AgB (which has previously been identified as genus specific) was found among the serum samples from patients with alveolar hydatidosis, the few cases of cross-reactivity observed with the peptides covered a wider spectrum of disease groups, and, in general, the cross-reactivity was marginal. The overall diagnostic values of the various antigens, which are best represented by their diagnostic efficiencies, indicate that p176 (diagnostic efficiency, 87%) is superior to AgB (diagnostic efficiency, 81%).

**DISCUSSION**

Native or recombinant antigens are powerful reagents for serology, because they contain a large spectrum of epitopes, which cover variations in the individual responses among patients. However, they represent complex structures, and there is always a balance between diagnostically relevant epitopes.
versus cross-reactive ones, which sometimes compromises the specificity of the system. For this reason, efforts have been made to find alternatives to biological antigens by using synthetic peptides (14, 16). Synthetic peptides mimicking relevant B-cell epitopes are potentially ideal tools for dissecting the antigenicities of the native antigens, making it possible to measure antibodies directed against very specific antigenic determinants. Additionally, synthetic peptides, in contrast to biological products, are easily standardized and can be readily produced in large amounts. Consequently, an increasing number of diagnostic assays based on synthetic peptides are being developed or are already commercially available.

In a previous study we performed an internal comparison of the diagnostic value of the available synthetic peptides and the major antigens of *E. granulosus* (3). That study showed that the diagnostic performance of AgB in ELISA was notoriously superior to that of other native antigens, such as Ag5 or crude hydatid cyst fluid, and also demonstrated that the peptides at hand constituted an unsatisfactory representation of the major epitopes of AgB. In order to find alternative peptide antigens that could mimic these epitopes, we examined new regions of both AgB subunits by using peptides p176, p175, and p177. Since these peptides were analyzed against the same panel of serum samples against which AgB and the other AgB-related peptides were analyzed, it provided an overall view of the antigenicity of AgB, which indicates that the dominant epitopes for humans are localized in the N-terminal extension of the 8-kDa subunits. In effect, those peptides delineated from the middle or C-terminal region of the AgB subunits (p177, Gu4, or p67 and p66 [11]) exhibited comparatively much lower sensitivities than p176, p175, or p65. Among these, p176 emerged, undoubtedly, as the most valuable antigen. Compared to p65, p176 has an additional 12-mer N-terminal fragment which appears to be crucial for recovery of supplementary immunodominant epitopes. As a consequence, the diagnostic sensitivity was augmented from 44% (p65) to 80% (p176) without a significant effect on the diagnostic specificity of p176.

In general, the use of a combination of two or more peptides

<table>
<thead>
<tr>
<th>Antigen</th>
<th>Sensitivity (%)</th>
<th>Specificity (%)</th>
<th>Diagnostic efficiency (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>p176</td>
<td>80</td>
<td>93</td>
<td>87</td>
</tr>
<tr>
<td>p175</td>
<td>49</td>
<td>94</td>
<td>74</td>
</tr>
<tr>
<td>p177</td>
<td>38</td>
<td>92</td>
<td>68</td>
</tr>
<tr>
<td>AgB</td>
<td>77</td>
<td>85</td>
<td>81</td>
</tr>
<tr>
<td>p65</td>
<td>44</td>
<td>96</td>
<td>73</td>
</tr>
<tr>
<td>Gu4</td>
<td>18</td>
<td>98</td>
<td>63</td>
</tr>
</tbody>
</table>

FIG. 2. Reactivity of our panel of sera against AgB and peptides p175, p176, and p177 as assessed by ELISA. The sera were grouped as follows: NS, sera from healthy donors; Eg, sera from patients with cystic hydatidosis; Em, sera from patients with alveolar hydatidosis; Ts, sera from patients with cysticercosis; Others, other sera used in this study. The cutoff for each assay is displayed as a horizontal line.
as probes in ELISA improves the performance of the peptide-based immunoassay (14). Therefore, we contemplated the possibility of complementarity among our antigens, particularly between p176 and p175. However, examination of individual E. granulosus-specific sera showed that all serum samples negative by ELISA against p176 did not react with any other peptide antigen, including p89-122 (data not shown), which has been shown to be complementary to p65 (3). This suggests that the antigenicity of the AgB molecule is concentrated in the N-terminal extension of the AgB8/1 subunit, which explains why p176 exhibited a better diagnostic performance than native AgB. Indeed, the fact that both the diagnostic sensitivity and the specificity of p176 were superior to those of AgB was a striking finding and provides remarkable experimental support for the peptide approach.

This improvement in the diagnostic efficiency of p176 was more evident in connection with the diagnostic specificity of the peptide (93 and 80% for p176 and AgB, respectively) and was mostly due to a lower level of cross-reactivity with sera from patients with alveolar echinococcosis. E. granulosus and E. multilocularis AgB8/1 subunits have a high degree of identity, and they differ at only five amino acid residues (8). Four of them are located in the stretch represented by p176, which may explain the fine specificity of p176.

Finally, examination of the individual sera did show in this case that a limited number of E. granulosus-specific serum samples reacted with AgB but not with p176, and vice versa. Therefore, there seem to be additional AgB epitopes which could be valuable in increasing the sensitivity of the system. These epitopes cannot be mimicked by linear peptides, as shown in this study. They appear to be discontinuous in nature, as these epitopes cannot be mimicked by linear peptides, as shown in this study. They appear to be discontinuous in nature, as these epitopes cannot be mimicked by linear peptides, as shown in this study. They appear to be discontinuous in nature, as these epitopes cannot be mimicked by linear peptides, as shown in this study. They appear to be discontinuous in nature, as these epitopes cannot be mimicked by linear peptides, as shown in this study.

This work was supported by The Swedish Agency for Research Cooperation with Developing Countries, the Consejo Nacional de Investigaciones Científicas y Técnicas, the Comisión Sectorial de Investigación Científica, Universidad de la República, Montevideo, Uruguay, and the Zafarani Foundation.

ACKNOWLEDGMENTS

This work was supported by The Swedish Agency for Research Cooperation with Developing Countries, the Consejo Nacional de Investigaciones Científicas y Técnicas, the Comisión Sectorial de Investigación Científica, Universidad de la República, Montevideo, Uruguay, and the Zafarani Foundation.

REFERENCES


