Gut, 1985, 26, 1210–1213

DR and non-DR Ia allotypes are associated with susceptibility to coeliac disease

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SUMMARY We have studied the frequency of HLA-DR, -MT, and -MB antigens in adult patients with coeliac disease and in a group of healthy controls, evaluating the strength of the observed associations by measuring the aetiologic fractions. Among the antigens significantly associated with coeliac disease, MB2 (DQw2) showed an aetiologic fraction higher than those of DR3 and DR7. Our results suggest that MB2, as associated more frequently than other specificities with a hypothetical disease predisposing gene, may play a role in the pathogenesis of coeliac disease. The significant association of DR3 and DR7 with coeliac disease may be ascribed to linkage disequilibrium of these antigens with MB2.

A number of immunologic diseases have been reported in association with HLA-DR antigens which are regarded as human homologues of murine Ia antigens. Recently increasing evidence has suggested that two additional series of Ia genes code for two additional series of Ia antigens termed MB2 and MT. Little is known about the association of these newly defined antigens and disease, but MB1 and MT2 have been reported associated with systemic lupus erythematosus and primary sicca syndrome. Moreover, it is noteworthy that the association between MB1 and systemic lupus erythematosus and between MT2 and primary sicca syndrome has been shown to be stronger than those between DR antigens and the above mentioned conditions.

These clinical studies suggest that MB and MT antigens may be more important than DR in genetic predisposition to some diseases. In view of this we have studied the frequency of MB and MT antigens in coeliac disease, a condition known to be associated with DR alleles DR3 and DR7.

Methods

Patients
Fifty one biopsy proven coeliac patients (aged 15–68 years, mean 38.7 years) were typed for DR and MT antigens. The control group was made up of 56 unrelated healthy subjects matched for sex, age, ethnic background, and geographic origin with coeliacs.

As far as MB2 and MB3 antigens are concerned, typing was carried out in 34 consecutive unselected and unrelated patients of the above mentioned 45 coeliacs and in 49 out of 56 control subjects.

HLA Typing
Purified lymphocyte preparations were obtained from heparinised peripheral blood by Ficoll-Hypaque sedimentation. Immunoglobulin bearing (B) lymphocytes were isolated from neuraminidase rosette forming cells (T lymphocytes) as described by Longo and Ferrara. Cytotoxicity tests were carried out by standard microcytotoxicity assay for B cells.

DR, MT and MB antigens were defined using a panel of 119 antisera; at least three to four antisera were tested to define each antigen specificity. Fifty four antisera were provided by the UCLA Tissue Typing Laboratory (University of California, Los Angeles, Ca), nine antisera were donated by Dr G B Ferrara (Istituto Tumori, Genova, Italy) and the remaining were some of the alloantisera utilised for disease studies in the 8th Histocompatibility Workshop 1980.

Statistical Analysis
The statistical significance of differences in the frequency of HLA antigens between coeliac patients and controls was calculated by $\chi^2$ analysis with
Yates’ correction. p Values were corrected by the number of comparisons made.

The strength of association between HLA antigens and coeliac disease was estimated for each specificity by the relative risk calculated by the method of Woolf, aetiologic and preventive fractions. Aetiologic fraction estimates which of many antigens associated with the same disease has the strongest genetic association with an assumed disease allele and shows how much of a disease is caused by the disease associated factor. The aetiologic fraction ranges from 0, if there is no association between the disease and the antigen, to 1 when the positive association is maximal – that is, all individuals affected by the disease carry the antigen. Preventive fraction is the analogous of aetiologic fraction when an association is negative.

Results

The distribution of HLA-DR, -MT and -MB specificities in coeliac patients and controls is shown in the Table. Positive significant associations were found for DR3, DR7 and MB2. The frequency of DR3 was 68.6% among patients as compared with a control frequency of 25% (p<0.0002), giving a relative risk of 6.56. The frequency of DR7 was 64.7% among coeliac patients as compared with a control frequency of 23.2% (p<0.0005), giving a relative risk of 6.06. The high frequency of the antigen MT2 not only among coeliac patients (92.1%) but also in the healthy population (78.5%) was responsible for the lack of significant association with coeliac disease.

Of the 34 coeliac patients and 49 controls typed with antisera recognising MB2 and MB3 antigens, 6 and 17, respectively, were found negative for both antigens. MB2 antigen was found in 76.4% of 34 patients against 24.4% of 49 controls (p<0.0002), giving a relative risk of 10.02.

The observed frequency distribution has been also evaluated as aetiologic and preventive fraction.

For MB2 antigen the aetiologic fraction (0.69) was shown to be higher than those for MT2 (0.63), DR3 (0.58) and DR7 (0.54) antigens.

Among negative associations, preventive fraction was particularly high for MT1 (0.26), DR5 (0.23), MB3 (0.20) and DR1 (0.18) antigens.

Discussion

Our results confirm, in adult patients, a significant association of coeliac disease with HLA-DR3 and DR7. As far as the association of the disease with Dw312 and DR313 is concerned, it has been confirmed by all subsequent studies, whereas contrasting results have appeared so far on the association between coeliac disease and DR7. The latter antigen does not seem to be associated with coeliac disease in North America, Ireland and the Netherlands, while in countries such as southern Germany, France, Italy, Spain and Israel, where the association with DR3, though significant, has been shown to be weaker the frequency of DR7 appears significantly increased. All reports but one, on the significant association between DR7 and coeliac disease concern paediatric patients and a recent work from England reports on the association of DR7 with coeliac disease in children but not in adults. These observations may suggest that juvenile coeliac patients are genetically different from adult coeliac patients. Our results are, however, not in keeping with this view. In fact our study confirms, for DR7, previous observations in childhood coeliac disease and also the results reported by

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Table. Frequency of DR and MT antigens in 51 patients with coeliac disease and in 56 controls. MB typing was performed in 34 patients and 49 controls

<table>
<thead>
<tr>
<th>Antigen</th>
<th>Patients (no. (%))</th>
<th>Controls (no. (%))</th>
<th>$\chi^2$</th>
<th>$p_r$</th>
<th>Relative risk</th>
<th>Aetiologic fraction</th>
<th>Preventive fraction</th>
</tr>
</thead>
<tbody>
<tr>
<td>DR1</td>
<td>1 (1-9)</td>
<td>11 (19-6)</td>
<td>6.7</td>
<td>NS</td>
<td>0-08</td>
<td>—</td>
<td>0-18</td>
</tr>
<tr>
<td>DR2</td>
<td>5 (9-8)</td>
<td>9 (16-1)</td>
<td>0-4</td>
<td>NS</td>
<td>0-57</td>
<td>—</td>
<td>0-07</td>
</tr>
<tr>
<td>DR3</td>
<td>35 (68-6)</td>
<td>14 (25-0)</td>
<td>18-7</td>
<td>&lt;0-0002</td>
<td>6-56</td>
<td>0-58</td>
<td>—</td>
</tr>
<tr>
<td>DR4</td>
<td>8 (15-6)</td>
<td>13 (23-2)</td>
<td>0-5</td>
<td>NS</td>
<td>0-61</td>
<td>—</td>
<td>0-09</td>
</tr>
<tr>
<td>DR5</td>
<td>12 (23-5)</td>
<td>23 (41-0)</td>
<td>2-9</td>
<td>NS</td>
<td>0-44</td>
<td>—</td>
<td>0-23</td>
</tr>
<tr>
<td>DR6</td>
<td>1 (1-9)</td>
<td>4 (7-1)</td>
<td>0-7</td>
<td>NS</td>
<td>0-26</td>
<td>—</td>
<td>0-05</td>
</tr>
<tr>
<td>DR7</td>
<td>33 (64-7)</td>
<td>13 (23-2)</td>
<td>17-1</td>
<td>&lt;0-0005</td>
<td>6-06</td>
<td>0-54</td>
<td>—</td>
</tr>
<tr>
<td>MT1</td>
<td>8 (15-6)</td>
<td>21 (37-5)</td>
<td>5-4</td>
<td>NS</td>
<td>0-31</td>
<td>—</td>
<td>0-26</td>
</tr>
<tr>
<td>MT2</td>
<td>47 (92-1)</td>
<td>44 (78-5)</td>
<td>2-9</td>
<td>NS</td>
<td>0-20</td>
<td>0-63</td>
<td>—</td>
</tr>
<tr>
<td>MT3</td>
<td>21 (41-2)</td>
<td>21 (37-5)</td>
<td>0-04</td>
<td>NS</td>
<td>1-17</td>
<td>—</td>
<td>0-06</td>
</tr>
<tr>
<td>MT4</td>
<td>3 (5-8)</td>
<td>3 (5-3)</td>
<td>0-09</td>
<td>NS</td>
<td>1-10</td>
<td>0-05</td>
<td>—</td>
</tr>
<tr>
<td>MB2</td>
<td>26 (76-4)</td>
<td>12 (24-4)</td>
<td>19-8</td>
<td>&lt;0-0002</td>
<td>10-02</td>
<td>0-69</td>
<td>—</td>
</tr>
<tr>
<td>MB3</td>
<td>9 (26-4)</td>
<td>20 (40-8)</td>
<td>1-2</td>
<td>NS</td>
<td>0-52</td>
<td>—</td>
<td>0-20</td>
</tr>
</tbody>
</table>
Betuel et al. in adult patients, suggesting that, at least in southern Europe, no difference seems to exist between HLA status of children and adults with coeliac disease; the genetic heterogeneity of coeliac disease, then, appears to be more a result of different geographic origin than of patients’ age.

As far as the frequency in coeliac disease of non-DR Ia allotypes is concerned, we have observed for the first time a significant association with the antigen MB2, which shows a relative risk higher than that of the other specificities significantly associated with coeliac disease – that is, DR3 and DR7. It has been suggested, however, that when several antigens with different allele frequencies are associated with the same disease, it is not possible to estimate which of them is more strongly associated with the disease itself solely by means of relative risk; that depends directly on the frequency of the antigen allele. A more accurate estimate of the strength of an association has been made by the calculation of the so-called aetiologic fraction that detects among the different antigens associated with a certain disease, which antigen has the stronger genetic association with a hypothetical disease allele. Our results point to MB2 as such an antigen for coeliac disease, as its aetiologic fraction is higher than that observed for DR3 and DR7. MT2 was more common not only among coeliac patients but also among controls. This is not surprising as MT2 is at present considered a broad supertypic antigen including DR3, DR5, DRw6, and DRw8 specificities, controlled by the DR subregion. That is why the raised aetiologic fraction of MT2 cannot be taken as evidence of a strong association of this antigen and coeliac disease.

In the normal population, MB2 is in strong linkage disequilibrium with DR3 and DR7, and this supports our view that the significant association of coeliac disease with these latter antigens may be secondary to the association with MB2. This is in keeping with the observation that some DR-associated diseases such as lupus erythematosus, ocular hystoplasmosis, and Takayasu’s arteritis, have stronger genetic associations with MB antigens.

Our study is in accordance with a recent paper by Tosi et al., who reported a complete association between coeliac disease and the new specificity DC3, considered the equivalent at the molecular level of MB2. We have found, however, a lower proportion of MB2 positive coeliac patients (76.4%) and this may be because of population differences and/or to minor dissimilarities between DC3 and MB2 and/or to the different techniques used: conventional cytotoxicity assay being used in our study, and RIA typing in that of Tosi et al.

As far as the antigens with a raised preventive}

fraction for coeliac disease are concerned, apart from MT1 and MB3 which have been considered for the first time in this study, we have confirmed the lower frequency, compared with the healthy population, of DR1 and DR5. It seems unlikely, however, that these specificities may play a true protective role for the development of coeliac disease. The concept of ‘protective gene’ would gain strength if a single gene was negatively associated with the disease, but the observation that more genes are less frequent in coeliac patients than in healthy subjects favours the hypothesis of an epiphenomenon compensatory to the increased frequency of DR3, DR7, MT2 and MB2.

In conclusion, our study supports the hypothesis previously formulated by others that MB2, as in linkage disequilibrium with either DR3 and DR7, is the antigen more strongly associated with coeliac disease. Because the association is not absolute, however, MB2 cannot be considered as the direct causative agent of the condition. More likely MB2 is closely associated with a still undefined HLA allele which, together with non-HLA linked genes and environmental factors, plays an important role in promoting the onset of coeliac disease.

This work was presented in part at the National Meeting of the Italian Society of Gastroenterology, Catania, Italy, November 1984. We are indebted to Marinella Cenci for invaluable technical assistance.

Addendum

After the completion of this study the nomenclature and tassonomy of some HLA specificities have been revised by the 9th International Histocompatibility Testing Workshop (Munich, 1984). MB2 is now termed DQw2 and is viewed as a DQ subregion product.

References

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*Gut* 1985 26: 1210-1213
doi: 10.1136/gut.26.11.1210

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