after a meal. Thus diabetic patients defined as those with random venous concentrations of at least 11 mmol/l (198 mg/100 ml) will be five standard deviations or more above the highest postprandial mean blood glucose concentration. Although we cannot say that we did not miss some diabetic patients in our series, these data suggest that this was unlikely; the incidence too (four in 2403 or one in 600 of a female population of reproductive years) was about the expected figure.

The stages of pregnancy at which samples for random determination of glucose concentration were obtained were those when blood was required for other routine antenatal tests; thus the cost of the needle and syringe was shared with several other tests. If the commonly used Multistix for testing urine, which give much unnecessary data and cost about 8 pence each, were replaced by a simple reagent strip to detect proteinuria only (at about 4 pence) the savings would help to compensate for the automated measurement of the blood glucose concentration. Clinic routine is not disrupted, the patients are not delayed or inconvenienced, and any increased laboratory cost can be balanced by saving on unnecessary urine tests. On the basis of random blood glucose screening two patients were diagnosed as having diabetes mellitus and 32 were correctly referred for an oral glucose tolerance test—that is, excluding the six patients in whom the blood glucose concentrations did not exceed the cut off values. Of these, a further two were found to have diabetes and 11 impaired glucose tolerance. Of 21 women referred for an oral glucose tolerance test for other reasons, three had impaired glucose tolerance. From another viewpoint, none of the diabetic patients was missed by the random checking of blood glucose concentration but only one of the four might have been referred for an oral glucose tolerance test on the basis of her glycosuria.

Is it necessary, however, to look for diabetes mellitus at all? The incidence of diabetes appears to be increasing rather than decreasing,^{8,9} and, though mortality and morbidity rates for females appear to be better than for males, they are still high¹⁰ and show wide regional variations. The sooner pregnant diabetic

women can be detected and returned to a normoglycaemic state the better their prognosis should be. The obstetrician must also be concerned for the fetus. In many hospitals now the perinatal mortality for diabetic mothers is not much greater than for the population at large,¹¹ which almost certainly reflects the increasing care taken in controlling the metabolic state of the mother. If a look for diabetes mellitus is accepted as a necessary part of good antenatal care random measurement of blood glucose concentration offers an easy, efficient, and relatively inexpensive method for doing this.

We thank Professor K G M M Alberti, Dr P A Smith, and our colleagues in the department of clinical biochemistry of the Royal Victoria Infirmary, Newcastle upon Tyne, for their help and co-operation with this study.

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(Accepted 26 September 1984)

HLA-DR typing in coeliac disease: evidence for genetic heterogeneity

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Abstract

Sixty nine probands from a family study of coeliac disease were typed for HLA-DR antigens. Sixty three (91%) were found to carry the antigen DR3, which was a significantly greater proportion ($p=9.6 \times 10^{-24}$) than among the 168 controls (26%). Concurrently 42 children

with the disease were DR typed. Not only was the frequency of DR3 significantly increased in these patients (86% versus 26% in controls; $p=3.1 \times 10^{-12}$) but so also was the frequency of DR7 (patients 60%, controls 29%; $p=5.8 \times 10^{-4}$). When those probands whose coeliac disease presented before the age of 20 were combined with the childhood coeliac group and a comparison made between these patients and the remainder of the probands, all of whom presented when they were older than 20, the childhood onset group had a significant excess of DR7 ($p=2.2 \times 10^{-3}$) and a significant deficiency of DR2 ($p=3.5 \times 10^{-3}$).

These findings indicate that childhood coeliac disease and adult coeliac disease are genetically heterogeneous.

Introduction

An association between coeliac disease and the histocompatibility antigen HLA-DR3 has been reported in all the populations

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in which it has been studied.¹⁻⁸ In some of these studies an association between coeliac disease and another DR locus antigen, DR7, has also been observed.^{2-5, 6} To explain these observations it has been postulated that there is a single coeliac disease gene in linkage disequilibrium with DR3 in all populations and with DR7 in some but not all populations. We have published the results of a family study into coeliac disease carried out in Liverpool.⁹ A preliminary analysis of the HLA-DR frequencies of several of the propiots from that study showed an association with DR3 but not DR7. Because of the difficulties that may sometimes be encountered in differentiating coeliac disease from other causes of a flat mucosa in childhood, and because of the possibility that childhood coeliac disease might be a heterogeneous disorder with permanent or transient gluten intolerance, we decided to look separately at a group of children with coeliac disease who fulfilled all the ESPGAN criteria for the diagnosis.¹⁰ Analysis of that group of patients disclosed an association with both HLA-DR3 and DR7. Those results suggested that the association between coeliac disease and HLA-DR7 may be an age related phenomenon. We therefore decided to study the HLA-DR antigen frequencies in a larger series of both groups of patients and to compare the frequencies in the children with those in the adults. This paper describes our results.

Subjects and methods

Patients—Sixty nine propiots from the family study in Liverpool were HLA-DR typed. The age at clinical onset of the disease ranged widely from infancy to adulthood. The diagnosis of coeliac disease was based on a jejunal biopsy showing total or subtotal villous atrophy with a positive clinical response to the withdrawal of gluten from the diet, an improvement in biochemical or haematological values, and,

in most cases, histological improvement. Forty two children who fulfilled all the ESPGAN criteria for coeliac disease were DR typed. There were 24 girls and 18 boys, and all were under 20 at the time of diagnosis (range 1 month to 11 years 10 months, mean 2 years 1 month). There was no overlap between the two groups of patients.

Controls—The control series consisted of 168 medical students and members of staff from the same geographical area.

HLA-DR typing was carried out using a cytotoxicity technique employing an adaptation of the two colour fluorescence method.¹¹ The panel of sera used allowed typing for seven specificities.

Statistical analysis—Fisher's exact test (two tailed) was used for the comparison of antigen frequencies.

Results

Table I lists the frequencies of the seven DR specificities tested in the control group and in the two groups of patients. The only significant positive association seen in the general series of 69 patients with coeliac disease was in respect of DR3, the relative risk being 27.3 ($p=9.6 \times 10^{-24}$). By contrast, the series of 42 patients with disease of early onset showed a positive association for both DR3 (relative risk 15.7; $p=3.1 \times 10^{-12}$) and DR7 (relative risk 3.5; $p=5.8 \times 10^{-4}$). Of additional interest was that the most notable negative association in the former group concerned DR1 (relative risk 0.09; $p=1.5 \times 10^{-4}$), while in the latter group both DR1 and DR2 appeared to be negatively associated (0.05 and 0.09 respectively). (In the first analysis two pairs of subjects were closely related but, as they were ascertained independently, all were included. In the second analysis only the first ascertained of each pair was included.)

Table II gives the relevant antigen frequencies when the patients as a whole were subdivided into those with clinical onset below the age of 20 and those with later onset. Comparison of the two groups showed a significant difference in the frequencies of DR7 ($p=2.2 \times 10^{-3}$). DR2 was significantly less frequent in the early onset group ($p=3.5 \times 10^{-3}$).

TABLE I—Frequencies of HLA-DR antigens in 69 propiots from coeliac family study, 42 children with coeliac disease, and controls

Antigen	Controls (n = 168)		Propositi from family study (n = 69)			p	Children with coeliac disease (n = 42)			p
	No positive	% positive	No positive	% positive	Relative risk		No positive	% positive	Relative risk	
DR1	32	19	1	1	0.09	1.5×10^{-4}	0	0	0.05	8.1×10^{-4}
DR2	48	29	15	22	0.71	0.36	1	2	0.09	1.2×10^{-4}
DR3	44	26	63	91	27.30	9.6×10^{-24}	36	86	15.70	3.1×10^{-12}
DR4	59	35	8	12	0.25	2.4×10^{-4}	6	14	0.33	0.011
DR5	21	13	2	3	0.25	0.03	4	10	0.80	0.82
DR6	26	16	5	7	0.46	0.13	2	5	0.33	0.098
DR7	49	29	23	33	1.20	0.63	25	60	3.52	5.8×10^{-4}

TABLE II—Comparison of frequencies of HLA-DR in adults and children with coeliac disease

Antigen	Onset before 20 years		Onset after 20 years		p
	No positive	% positive	No positive	% positive	
DR1	0	0	1	2	0.84
DR2	3	5	12	26	3.5×10^{-3}
DR3	56	89	41	89	1.0
DR4	9	14	5	11	0.82
DR5	4	6	2	4	1.0
DR6	2	3	4	9	0.41
DR7	36	57	12	26	2.2×10^{-3}

Discussion

These results show that (in Liverpool) there are significant differences in the frequencies of certain HLA antigens between those patients with coeliac disease who present before the age of 20 and those who present later. This suggests that these two subgroups of patients with coeliac disease differ genetically. In particular, coeliac disease in adults appears to be associated with DR3, whereas with an earlier onset the association is with both DR3 and DR7. It is of additional interest that DR2 was significantly reduced in frequency in the early onset group only.

TABLE III—Frequencies of HLA-DR antigens in coeliac disease in present study compared with frequencies in previous studies separated according to age of patients (P) and controls (C)

Antigen	Childhood studies								Present study		Adult studies					
	Italy ³		Spain ⁴		Austria/Southern Germany ²		Finland ⁴		Children	Adults	France ⁵		Holland ⁶		Ireland ⁷	
	P	C	P	C	P	C	P	C			P	C	P	C	P	C
DR1	0	15	4	22	6	20	—	—	0	19	—	—	13	20	12	34
DR2	4	26	4	26	6	28	—	—	5	29	—	—	29	32	14	29
DR3	60	22	77	18	63	20	85	24	89	26	64	13	96	22	88	44
DR4	9	19	0	16	10	27	—	—	14	35	—	—	8	31	14	13
DR5	22	35	23	24	21	6	—	—	6	13	—	—	0	23	19	23
DR6	9	44	4	26	28	20	—	—	3	16	—	—	13	34	9	9
DR7	67	28	69	44	41	28	66	11	57	29	55	10	21	23	27	34

Five of the six previously reported studies in which an association between coeliac disease and DR7 was found were from paediatric centres. Of the two studies recording an association with DR3 alone, one concerned adults only⁹; a second study included a mixed group of children and adults, though 74% were adults⁷ (table III).

The association of coeliac disease with DR3 and DR7 may be interpreted on the basis of more than one model. The condition may, for example, be genetically homogenous and dependent on a susceptibility gene at a locus linked to the DR locus and in linkage disequilibrium with both DR3 and DR7. In this regard it is relevant that the DC3 specificity at the DC locus is in linkage disequilibrium with DR3 and DR7, and Tosi *et al* put forward evidence to suggest that the primary association was with DC3.¹² A recent study of patients with coeliac disease who were DR3 and DR7 negative, however, failed to confirm this.¹³

Our finding of heterogeneity of the DR association argues against this simple model and suggests the possibility of more than one susceptibility gene—namely, DR3 and DR7 or genes in linkage disequilibrium, one with DR3 and one with DR7. If the presence of both genes promotes an early onset of disease, and the presence of only DR3 or a gene in linkage disequilibrium with DR3 a later onset, the present findings would be explained.

Further studies of the association of coeliac disease with DR and DC antigens are being carried out, and it seems important that age of onset of disease should be taken into account. Other than the age of onset, HLA differences do not seem to have any influence on other clinical variables. Kumar *et al* found that HLA factors did not appear to make any difference to the mode of presentation.¹⁴

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(Accepted 26 September 1984)

The sick building syndrome: prevalence studies

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Abstract

Random samples of the entire workforce in nine offices in which similar clerical work was being performed were studied using a doctor administered questionnaire that inquired into symptoms that have been linked with the "sick building syndrome." Five of the offices were fully air conditioned, one had recirculation of air and mechanical ventilation, and three were naturally ventilated. Workers in three air conditioned and three naturally ventilated buildings were interviewed blind. Seven of the buildings were studied at our request in the absence of any known problem. Comparison of prevalences of symptoms between the naturally ventilated and the other buildings showed a repeated pattern of nasal, eye, and mucous membrane symptoms with

lethargy, dry skin, and headaches. There were highly significant excesses of these six symptoms in the air conditioned buildings when compared by χ^2 tests with the naturally ventilated buildings.

It is suggested that these six symptoms represent the sick building syndrome and that the size of the problem is probably greater than is currently recognised. Possible causes are discussed.

Introduction

The "sick building syndrome" is generally taken to describe a building in which complaints of ill health are more common than might reasonably be expected. The affected buildings are usually offices that have full air conditioning. The excess of complaints of ill health is not usually reflected in an increase in sickness absence. No definition of the size of the problem has previously been systematically attempted, although Turiel *et al* noted excesses of eye, upper respiratory tract, and chest symptoms in an open comparison of an air conditioned and a naturally ventilated building.¹ In that study 62% of the workers in the air conditioned building and 66% in the naturally ventilated building were interviewed.

Several causes have been postulated for symptoms in office buildings, and, despite much research, no satisfactory explanation of the problem has been found. The postulated causes include formaldehyde (from cavity wall insulation, office

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