Class II HLA antigens in multiple sclerosis

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SUMMARY HLA typing in Wellington revealed a stronger association of multiple sclerosis with DR2 than with DQw1. The association with DQw1 appeared to be due to linkage disequilibrium of this antigen with DR2. These results, when considered in conjunction with other studies, are most easily explained by the hypothesis that susceptibility to multiple sclerosis is influenced by multiple risk factors, with DR2 being an important risk factor in Caucasoid populations.

The cause of multiple sclerosis is unknown. Current evidence suggests that both environmental and genetic factors are of aetiological importance. Multiple sclerosis occurs most frequently in temperate zones, though the prevalence in such climatic regions varies for different ethnic groups. Thus, the disease is common in Caucasians in Northern Europe, North America and Australasia, but is much less common in Orientals and Polynesians, living in the same latitude and environment. Strong evidence implicating genetic factors in susceptibility to multiple sclerosis has come from the Canadian twin study. This population-based study found concordance for multiple sclerosis in 1.9% of non twin siblings, 2.3% of dizygotic twins and 26% of monozygotic twins.

The search for a multiple sclerosis susceptibility gene has been encouraged by the discovery in the last 16 years of a number of consistent associations of HLA antigens and multiple sclerosis. The initial associations in Caucasoid populations were with the Class I antigens A3 and B7. A stronger association with the Class II antigen DR2 (which is in linkage disequilibrium with A3 and B7) was subsequently identified. This association has been observed in Caucasians in the United Kingdom, Europe, North America, Australia and New Zealand. On average, about 50-70% of Caucasoid multiple sclerosis patients are DR2 positive compared with 16-30% of controls.

The association of DR2 with multiple sclerosis is not seen in the Aberdeen region, or the Orkney Islands, areas where the prevalence of multiple sclerosis is exceptionally high. The lack of an association is due to the unusually high frequency of the DR2 antigen in the normal population of these regions.

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Received 25 October 1988.
Accepted 19 December 1988

RR = Antigen+/MS × Antigen-/-control
Antigen-/MS × Antigen+/-control

(41% in Aberdeen and 50% in the Orkneys). However, an association in Aberdeen has been reported with the more recently defined Class II antigen, DQw1.11 The same association has since been reported elsewhere.12,13

To assess the relative significance of DR2 and DQw1 in relation to multiple sclerosis, we tissue typed the Class II DR and DQ antigens in Caucasian multiple sclerosis patients in the Wellington region. The findings were compared with healthy Caucasian controls. HLA data were also collected in Maori controls. A comparison of the findings in Caucasoid and Maori controls was undertaken, in view of the known rarity of multiple sclerosis in Maoris, and because a previous study demonstrated different antigen frequencies between these ethnic groups, in particular a low frequency of DR2 in Maoris.2

Methods

A blood sample for tissue typing was taken from multiple sclerosis patients who were seen as part of a review of all identifiable cases in the Wellington region in 1983. The method of case ascertainment is detailed elsewhere.2 HLA Class I and Class II typing was performed by a standard lympho-cytotoxicity test following T and B cell lymphocyte separation using a nylon wool column. Only the results of Class II typing are discussed in this paper. The typing sera used defined 11 HLA-DR and 3 HLA-DQ specificities. The Caucasoid controls comprised 117 normal healthy blood donors. The Maori control population comprised 82 individuals who self identified as Maoris. The multiple sclerosis group comprised 144 Caucasians with clinically definite or probable multiple sclerosis using the McDonald-Halliday criteria.14 (No Maori patients fulfilling these criteria were identified.)

Statistical evaluation of the data was performed using the chi-squared test with correction, where appropriate, for multiple comparisons. The relative risk (RR) associated with the presence of an antigen for the development of multiple sclerosis was calculated using the formula of Svejgaard et al:2

RR = Antigen+/MS × Antigen-/-control
Antigen-/MS × Antigen+/-control
Results

(1) Comparison of Caucasian controls and multiple sclerosis patients (table 1)

There was a significant excess of both DR2 and DQwl in multiple sclerosis patients. The association with DR2 (RR = 6-9, \( \chi^2 = 50-8 \), \( p < 0-001 \)) was stronger than that with DQwl (RR = 3-3, \( \chi^2 = 17-9 \), \( p < 0-001 \)). All DR2 positive subjects were also DQwl positive. Only 15% of multiple sclerosis patients were DQwl positive/DR2 negative, whereas this antigen combination was seen in 35% of controls. For the subgroup of DR2 negative individuals, the frequency of DQwl was almost the same in multiple sclerosis patients and Caucasian controls (multiple sclerosis: 21/47, 45%; controls: 41/90, 46%; \( \chi^2 = 0-01 \), NS).

(2) Comparison of Caucasian and Maori controls (table 2)

DQwl was present in 58% of Caucasoid and 55% of Maori controls (\( \chi^2 = 0-2 \), NS). As has been previously reported, only DR2 was more prevalent in Caucasoids than in Maoris, although for the groups in the present study, this difference was not statistically significant (24% versus 13%, \( \chi^2 = 2-9 \), NS). In Maoris, there were significantly higher frequencies of DR5 and DQw3, and lower frequencies of DR3 and DQw2.

(3) Comparison with Cardiff and Aberdeen studies (table 3)

The DR2 and DQwl frequencies for Wellington multiple sclerosis patients and Caucasian controls were compared with those obtained from recent studies centred in Aberdeen \(^{10} \) and Cardiff. \(^{12} \)

(a) DR2. In multiple sclerosis patients, there was a higher frequency of DR2 in Wellington (\( \chi^2 = 11-3 \), degrees of freedom (df) = 2, \( p = 0-004 \)). In controls, there was a higher frequency of DR2 in Aberdeen than that with DR2, \( (\chi^2 = 50-8, p < 0-001) \), was stronger than that with DQwl, \( (\chi^2 = 17-9, p < 0-001) \).

All DR2 positive subjects were also DQwl positive. Only 15% of multiple sclerosis patients were DQwl positive/DR2 negative, whereas this antigen combination was seen in 35% of controls. For the subgroup of DR2 negative individuals, the frequency of DQwl was almost the same in multiple sclerosis patients and Caucasian controls (multiple sclerosis: 21/47, 45%; controls: 41/90, 46%; \( \chi^2 = 0-01 \), NS).

(b) DQwl. In multiple sclerosis patients, the frequency of DQwl was not significantly different between the three regions (\( \chi^2 = 2-6, df = 2, p = 0-28 \)). DQwl was present with a remarkably similar frequency in the three control populations.

Discussion

This study confirms previous ones in showing that both DR2 and DQwl are associated with multiple sclerosis in Caucasian populations. Although the frequency of DQwl positive cases was higher than DR2 positive ones, the association with DR2 was seen to be more significant when compared with the control population.

In the present study, the association of multiple sclerosis with DQwl appears to be secondary to the known linkage disequilibrium of this antigen with DR2, rather than a primary association. Thus, in DR2 negative multiple sclerosis patients, there was no excess of DQwl. It is also notable that DQwl was seen with a high frequency in Maori controls, a population with a very low risk for multiple sclerosis despite living in a high-risk temperate region.

A stronger association of multiple sclerosis with DR2 (than with DQwl) was also found in Cardiff. \(^{13} \) However, in Aberdeen, the stronger association was with DQwl. \(^{10} \) The lack of an association with DR2 in the latter study was largely due to an unusually high frequency of DR2 in the normal population. Indeed, a correlation has been demonstrated between the regional prevalence of multiple sclerosis in the United Kingdom and the frequency of DR2 in the normal population. \(^{17} \)

It has been argued that there is a single multiple sclerosis susceptibility gene on the sixth chromosome in linkage disequilibrium with the HLA system. \(^{18} \) \(^{19} \) If this hypothesis is true, the present results suggest that the putative gene is closer to the DR than the DQ locus and that DR2 is currently the best marker for that gene.
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Alternatively, as Swingler et al have pointed out, the HLA findings in multiple sclerosis may be interpreted by considering that multiple sclerosis susceptibility is related to genetic and environmental factors. Such a multifactorial hypothesis would allow for the lack of association of multiple sclerosis and DR2 in certain populations, different HLA antigens in some non-Caucasoid populations, a high association of DR2 in some populations in which multiple sclerosis is rare, and the occurrence of other genetic associations unrelated to the HLA system.

Swingler et al have discussed the multifactorial hypothesis in terms of multiple risk factors of varying importance. A high frequency of major risk factors in a population will result in a high prevalence of the disease. However, that same high frequency of major risk factors may actually obscure their association with the disease in such a population. Associations may be seen instead with more minor risk factors. Conversely, when major risk factors are less common in a population, the disease prevalence will be less, but the major risk factors will show the strongest associations with the disease.

If we assume that there are multiple risk factors of varying importance predisposing to the development of multiple sclerosis, it might be concluded from the studies of Class II HLA antigens summarised in table 3 that the association with DR2 is of major importance at least in Caucasoid populations. Thus, in Aberdeen where the prevalence of multiple sclerosis is 178/105, DR2 is weakly associated with multiple sclerosis because of its high frequency in the normal population; in Cardiff, where the prevalence of multiple sclerosis is two thirds that of Aberdeen, a stronger and statistically significant association with DR2 is seen; in Wellington, where the prevalence of multiple sclerosis is one third that in Aberdeen, the strongest association of DR2 with multiple sclerosis is seen.

Overall, only a small proportion of the total DR2 positive population actually develops multiple sclerosis. It seems likely therefore that there are other risk factors of major importance for the development of multiple sclerosis which are as yet undetermined.

The study was supported in part by a grant from The New Zealand Multiple Sclerosis Society. Mr G Purdie helped with statistical analysis.

References