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## ASSOCIATION BETWEEN HISTOCOMPATIBILITY ANTIGENS (HLA) AND NASAL CARRIAGE OF *STAPHYLOCOCCUS AUREUS*

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**SUMMARY.** We investigated the association between phenotypes of histocompatibility antigen (HLA) and nasal carriage of *Staphylococcus aureus* in two populations—healthy laboratory workers and patients attending an outpatients' clinic. When data from the two sources were pooled, it was evident that the presence of HLA-DR3 was associated with carriage, and the presence of HLA-DR2, HLA-DR1 and HLA-Bw35 with lack of carriage. However, since each person may have two antigenic specificities encoded at the HLA-A, the HLA-B, and the HLA-DR loci, the carriage of the organism was analysed for paired combinations of the more frequent phenotypes. For example, the lack of carriage evident with HLA-DR1 was more marked with the DR1-A11 and DR1-B7 combinations while the predisposition towards carriage shown with HLA-DR3 was more marked with the DR3-DR5 combination. The importance of the analysis of antigen combinations is discussed in relation to association of single antigens with carriage of *S. aureus*.

### INTRODUCTION

There appear to be three main classes of nasal carrier of *Staphylococcus aureus*: the persistent carrier, the persistent non-carrier and the intermittent carrier (Williams, 1963) but no convincing reason for this has yet been put forward. A genetic predisposition to nasal carriage among healthy subjects has been recognized for some time. Noble, Valkenburg and Wolters (1967) analysed carriage within families and found that predisposition to the carrier state was not explained simply by exposure to a common reservoir of bacteria. In addition, from studies on twins, Hoeskma and Winkler (1963) showed that the percentage similarity in carriage was greater in identical twins than non-identical twins or unrelated children. Finally it has been observed that there are some racial differences, e.g., negroids carry *S. aureus* less frequently than do caucasoids (Findlay and Abrahams, 1946; Millian *et al.*, 1960; Noble, 1974).

We investigated the possible association between HLA type and nasal carriage of *S. aureus* in an attempt to explain this genetic predisposition and to study further the colonization of mucosa by this organism.

### MATERIALS AND METHODS

**Subjects.** Nasal carriage of *S. aureus* was investigated in two populations. The first, of 90 adults, was studied in Ireland and consisted of people who had donated blood for HLA typing other studies and was mostly made up of patients attending outpatient clinics in Dublin and Galway. The diagnoses were as follows: Graves's disease 62, coeliac disease 14, psoriasis 8, and juvenile-onset diabetes 6. Antigens of the HLA-A and -B series were detected by the standard NIH Terasaki microlymphocytotoxicity test, and of the HLA-DR series by a cytotoxicity test after B-cell enrichment according to van Rood *et al.* (1975) with a total of 60 B-cell antisera. The second population was of 90 healthy laboratory workers in London on whom HLA typing



had been performed, by standard techniques and for reasons other than the present study, in one of four tissue-typing centres.

*Bacteriology.* Swabs were taken from the anterior nares and placed in Amies's Transport Medium. In the laboratory they were seeded on to nutrient agar and mannitol-salt agar (Oxoid). All coagulase-positive staphylococci isolated were maintained on nutrient-agar slopes and subsequently phage typed. Each person was swabbed only once and those yielding more than three colonies of *S. aureus* were designated carriers.

*Analysis of results.* Antigens Aw23 and Aw24 were grouped with antigen A9, from which they have only recently been separated; similarly, Bw51 was grouped with B5 and Bw44 with B12.

Relations between carriage of *S. aureus* and HLA phenotype were sought by determining the relative risk of carriage associated with the possession of each antigen or antigen combination. This was defined as

$$\frac{a}{b} \times \frac{d}{c}$$

where  $a$  = the number of persons with the antigen or antigens who were carriers,  $b$  = the number with the antigen or antigens who were non-carriers,  $c$  = the number without the antigen or antigens who were carriers, and  $d$  = the number without the antigen or antigens who were non-carriers. Significant departures from a relative risk value of 1.0 were determined by means of Fisher's exact test, but probabilities were not adjusted for the number of antigens for which tests had been made (the Bodmer correction factor) because this would have been meaningless when analysing results in respect of combinations of antigens.

#### RESULTS

The frequency of nasal carriage in the sample of laboratory workers was 30%. This corresponds well with results from previous studies (Williams, 1963). The patient group had a slightly higher frequency (41%) than the laboratory workers but the difference was not significant ( $\chi^2$  test:  $p > 0.05$ ). Length of stay in hospital may increase the frequency of nasal carriage of *S. aureus* (Ayliffe *et al.*, 1977), but none of our patients was in hospital when included in the sample.

Table I shows the frequencies of the more common HLA phenotypes of the two populations studied, together with the associated relative risk of carriage of *S. aureus*. The distribution of individual antigens in the two groups was markedly different, but the patients included a number with diagnoses expected to be associated with the presence of antigens B8 and DR3.

For antigens in the HLA-A and HLA-B series, the only statistically significant associations were those between lack of carriage and the presence of antigen A3 or Bw35 in the laboratory-workers' group; in the patients' group there was a non-significant trend in the same direction (relative risk  $< 1.0$ ). For antigens in the HLA-DR series, there was agreement in trend in respect of five of the seven antigens, towards non-carriage in bearers of antigens DR1, DR2 and DR4, and towards carriage (relative risk  $> 1.0$ ) in bearers of antigens DR3 and DR7. These reached statistical significance for DR1 in both groups, and for DR1, DR2 and DR3 only in the patients' group. With two of the antigens—DR5 and DRw6—the trends were contradictory; the presence of DRw6 showed a significant association with non-carriage in the laboratory workers and with carriage in the patients. However, further inspection suggested that these anomalies might be explained by the contribution of other antigens expressed by the same subjects, because each of them may have two antigenic specificities at HLA-A, HLA-B or HLA-DR loci. For example, in the case of HLA-DR5, 12 of the 22 patients expressing DR5 also expressed DR3 but none of the 15 laboratory workers with DR5 expressed DR3.

Thus it may be more relevant to consider antigen combinations than individual antigens. For this purpose we thought it permissible to increase the number of combinations available for comparison by bringing together the results obtained in the two series. It seemed reasonable to assume that, though the two groups differed in HLA constitution, they would show a similar relation of HLA phenotype to nasal carriage. Selected results are shown in table II. The relative



TABLE I

Frequency of occurrence of individual HLA phenotypes, and of carriage of *S. aureus* associated with them, in laboratory workers and patients

HLA antigen	Laboratory workers			Patients		
	Frequency of the stated antigen*	Relative risk of carriage†	p=	Frequency of the stated antigen*	Relative risk of carriage†	p=
HLA-A1	27	1.72	0.106	41	1.48	0.110
HLA-A2	43	1.56	0.115	51	1.08	0.119
HLA-A3	17	0.25	<b>0.045</b>	21	0.84	0.191
HLA-A9/w23/w24	20	0.47	0.109	14	1.08	0.227
HLA-A11	14	0.59	0.199	8	0.44	0.197
HLA-A26	9	1.18	0.281	0	...	...
HLA-A29	6	...	...	8	0.84	0.287
HLA-B5/w51	8	2.56	0.134	9	0.69	0.252
HLA-B7	16	1.07	0.230	20	0.39	0.053
HLA-B8	15	...	...	45	1.71	0.079
HLA-B12/w44	23	0.77	0.190	28	0.89	0.178
HLA-B13	6	...	...	4	...	...
HLA-B14	6	...	...	8	2.60	0.131
HLA-B15	8	0.76	0.307	4	...	...
HLA-B18	9	2.01	0.174	1	...	...
HLA-B27	11	0.85	0.270	11	1.85	0.160
HLA-Bw35	13	0.16	<b>0.043</b>	14	0.33	0.065
HLA-B40 <sup>‡</sup>	6	...	...	6	...	...
HLA-DR1	20	0.51	0.120	21	0.36	<b>0.038</b>
HLA-DR2	28	0.90	0.193	28	0.35	<b>0.021</b>
HLA-DR3	16	1.02	0.240	49	2.51	<b>0.019</b>
HLA-DR4	32	0.86	0.183	24	0.63	0.130
HLA-DR5	15	0.82	0.230	22	2.06	0.067
HLA-DRw6	15	0.13	<b>0.022</b>	9	5.95	<b>0.020</b>
HLA-DR7	27	2.01	<b>0.069</b>	17	1.00	0.210

Figures in bold type indicate a statistically significant association with carriage or non-carriage.

\* Number (of 90 subjects) with the stated antigen.

† See *Materials and Methods*.

risk of carriage is given for each antigen individually, and for antigen combinations when their frequency was greater than five. Although each person may express six antigenic specificities, only paired combinations are shown. In the combined populations, three antigens (Bw35, DR2 and DR1) were significantly associated with lack of carriage while one antigen (DR3) was associated with carriage. However, when the results for combinations of antigens were assessed it was evident that the relative risk was much reduced for certain antigen combinations; for example, the overall relative risk for DR1 was 0.4 whereas that of the combinations DR1-All and DR1-B7 was reduced to zero. The figures in bold type denote a significant association with carriage or lack of carriage. Thus some antigen combinations predispose towards lack of carriage of *S. aureus* while others predispose towards carriage or are indeterminate in effect. It is notable that predisposition towards lack of carriage associated with antigen DR1 or Bw35 seems to be dominant when in combination with other phenotypes, but further data is necessary to clarify this point.

There was no evidence of an association between the phage type of the *S. aureus* strains isolated and the HLA type of the carrier.

#### DISCUSSION

We have shown that certain HLA phenotypes apparently predispose towards nasal carriage of *S. aureus* and others predispose towards lack of carriage. This is most marked with particular combinations of antigens. The potential importance of combinations of antigens has been



TABLE II

Association between relative risk of carriage of *S. aureus* and some HLA phenotype-combinations in a conflated group of 180 laboratory workers and patients

HLA phenotype	Relative risk of carriage* for single HLA	Relative risk of carriage* for combination of the stated phenotype with phenotype						
		A3	A11	B8	Bw35	DR3	DR2	DR1
A1	1.4(68)	0.5(8)	0.2(7)	1.3(43)	...	1.6(42)	0.7(10)	0.3(12)
A2	0.9(94)	1.4(11)	...	1.2(22)	0(10)	0.9(33)	0.6(25)	0.4(17)
A9/w23/w24	0.6(38)	...	...	0.9(11)	0.5(8)	1.5(13)	0.6(7)	0.4(9)
A3	0.6(38)	...	...	0.6(7)	0.6(7)	...	0.5(20)	0.5(8)
A11	0.4(22)	...	...	...	0(6)	1.7(6)	0.1(11)	0(7)
B27	1.2(22)	...	0.8(7)	...	...	4.5(7)	0.2(7)	...
B8	1.0(60)	0.6(7)	...	...	...	1.8(44)	0.4(10)	0.3(11)
B12/w44	0.8(51)	1.3(9)	...	1.7(12)	0.8(6)	1.3(16)	0.6(17)	0.7(10)
B7	0.5(36)	0.3(18)	0(6)	...	...	0.5(8)	0.4(17)	0(7)
Bw35	0.3(27)	0.6(7)	0(6)	...	...	...	0.6(7)	0.4(10)
DR3	1.9(65)	...	1.7(6)	1.8(44)	...	...	1.1(10)	0.3(11)
DR7	1.4(44)	3.5(6)	...	1.3(9)	...	3.5(6)	1.4(11)	...
DR5	1.3(37)	...	0.8(6)	2.3(7)	...	3.6(9)	0.5(8)	...
DRw6	0.8(24)	...	...	1.2(7)	...	2.3(7)	...	...
DR4	0.6(54)	0(8)	...	0.4(13)	...	1.0(16)	0.4(10)	...
DR2	0.5(56)	0.5(20)	0.1(11)	0.4(10)	0.6(7)	1.1(10)	...	0.4(9)
DR1	0.4(41)	0.5(8)	0(7)	0.3(11)	0.4(10)	0.3(11)	0.4(9)	...

In parentheses: frequency (among 180 subjects) of HLA phenotype or phenotype-combination; ... = frequency < 6; in bold type:  $p < 0.05$ .

\* See Materials and Methods.

emphasized recently (Kaslow and Shaw, 1981). Linkage disequilibrium refers to the situation in which certain combinations occur more frequently than would be expected if the alleles were randomly distributed, e.g., A1-B8-DR3, and suggests some selective advantage. From the present study it is also obvious that in the populations studied certain combinations occur more frequently than others. A gene or genes responsible for carriage or lack of carriage of *S. aureus* may be closely linked with the HLA locus or, rather, it may be in linkage disequilibrium with certain HLA phenotypes. Linkage may be determined by investigating carriage in relation to HLA haplotypes but these can be determined only from family studies.

Recent evidence from antibody responses and T-cell proliferative responses suggests that there is an HLA-linked control of the differential immune responsiveness to microbial antigens (Kaslow and Shaw, 1981). This has been studied by measuring immune responsiveness to naturally acquired immunogens such as streptococcal cell-wall antigens (Greenberg, Gray and Yunis, 1975; Greenberg *et al.*, 1980; Lehner *et al.*, 1981) or to vaccination (Kato *et al.*, 1978; Sasazuki *et al.*, 1980; Nose *et al.*, 1980). Many disease associations have been explained in this way. Ehrenkranz (1966) investigated the nasal rejection of experimentally inoculated *S. aureus* in non-carriers and found that the length of time taken to eliminate the bacteria was less on subsequent occasions than on the first and that this response was specific for the staphylococcal strain. Despite the fact that no difference in serum antibody levels to *S. aureus* has been detected between carriers and non-carriers (Daugharty, Martin and White, 1969), the ability to carry *S. aureus* may possibly be determined by an HLA-linked control of immunity to this organism. Perhaps the Gm genotype of the host may have a controlling influence, as has been recently suggested by Nakao *et al.* (1980) and Whittingham *et al.* (1980).

An alternative hypothesis is that carriers have receptors for this organism. Aly *et al.* (1977) showed that staphylococci adhered more to washed nasal epithelial cells of carriers than to cells of non-carriers. One suggestion for the association between HLA and infectious disease is that the HLA molecule acts as the receptor for organisms, but firm evidence for this is available in only a few instances (Helenius *et al.*, 1978).

The mechanism of interaction between HLA gene products or HLA-linked gene products



and microorganisms remains unclear, but the colonisation of nasal mucosa by *S. aureus* could serve as a model to investigate this further.

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