

A Correlative Study of HLA, Sickle Cell Gene and G6PD Deficiency with Splenomegaly and Malaria Incidence Among Bhils and Pawra Tribes from Dhadgon, Dhule, Maharashtra

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INTRODUCTION

The tribal groups in India form a significant proportion (8.01%) and they are believed to be the earliest settlers or inhabitants of the region. There are 427 tribal communities in India (Singh, 1993). The dominant ethnic group among the Indian tribal group is proto-australoid although groups living in the sub-Himalayan belt have more of Asian characteristics. The three major tribes having a population of more than a million are Gonds of Madhya Pradesh, Maharashtra and Andhra Pradesh, the Bhils of Rajasthan, Gujarat, Maharashtra and Madhya Pradesh and the Santhals of Bihar, Orissa and West Bengal (Vidhyarthi, 1983). The Bhils are one of the largest tribes concentrated mainly in Western Madhya Pradesh, Rajasthan, Eastern Gujarat and Northern Maharashtra. Racially they were classified as Gonds, Malis or Proto-Australoid, but their social history is still a mystery (Bhatia and Rao, 1986). The Pawras resemble a primitive tribe Koli of Western Satpura Hills and are concentrated mainly in Khandesh. Even today these tribal groups hardly perform their marriages outside their tribe, which makes them unique from the cast groups of plains.

There are several reports in literature on population genetics that show prevalence of G6PD deficiency, Sickle cell trait frequency, in different caste groups (Baxi et al., 1963, 1964; Mehta et al., 1971; Bhasin et al., 1992; Joshi et al., 2001, Bhasin and Walter, 2001) and tribal groups (Rao and Bhatia, 1988; Gorackshakar et al., 1987; Bhasin et al., 1992; Kaur et al., 1997; Ramasamy et al., 1994) from India. Further there are reports on the genetic diversity of HLA in different tribal groups from India (Shankarkumar et al., 1999). But still there are very little information on the correlation of Sickle cell trait, G6PD deficiency and HLA in a caste/tribal group from India where Malaria is endemic. This initiated us to undertake

the current study among the Malaria endemic population of Bhils and Pawras from Dhadgon, Dhule district, Maharashtra, Western India.

MATERIALS AND METHODS

Samples: A total of 146 tribal comprising Bhils (91) and Pawras (55) living in Dhadgon, Dhule district, Maharashtra were studied for HLA, Sickle cell Trait and G6PD deficiency. Blood samples were collected in EDTA and analysis was done within 24 hours. They were clinically assessed by (KG) for the presence of splenomegaly and Malarial incidence by standard technique (John Mcleod, 1983).

HLA Typing: The two - stage NIH standard micro-lymphocytotoxicity test was carried out on all the samples as described elsewhere (Shankarkumar et al., 1999).

G6PD Assay: Screening for G6PD deficiency was done by the dichlorophenol-indophenol (DPIP) dye decolorization method (Bernstein 1962). The RBC indices and hematological parameters were estimated using an automated cell counter (ERMA - PC, 608, Japan). Hemoglobin analysis was estimated using the Variant hemoglobin testing system (Bio-Rad, Inc Ohio, USA) using recommended protocols. Solubility test and cellulose acetate electrophoresis using Tris - EDTA borate buffer at pH 8.9 (Dacie and Lewis, 1975) was also done.

RESULTS

HLA Antigen Associations, Sickle Cell Trait and G6PD Deficiency Comparisons: Analysis of the HLA antigens showed that in the splenomegaly and Malaria positive group HLA A1, B40 and B53 ($P < 0.001$) were increased when compared to the negative group where A2, B7 and B35 ($P < 0.01$) were decreased (Fig. 1). HLA A2, B8, B17 and B40 ($P < 0.001$) were significantly

increased in the G6PD deficient group when compared to the normal where A19, A11, and B53 (P<0.01) were decreased (Fig. 2). HLA A2, B40 and B53 (P<0.0001) were increased among the sickle cell heterozygotes when compared to the

homozygotes where A19 and B8 (P< 0.05) were decreased (Fig. 3). Moreover the HLA distribution among the tribes were diverse from other reported tribal populations (Shankarkumar et al., 1999).

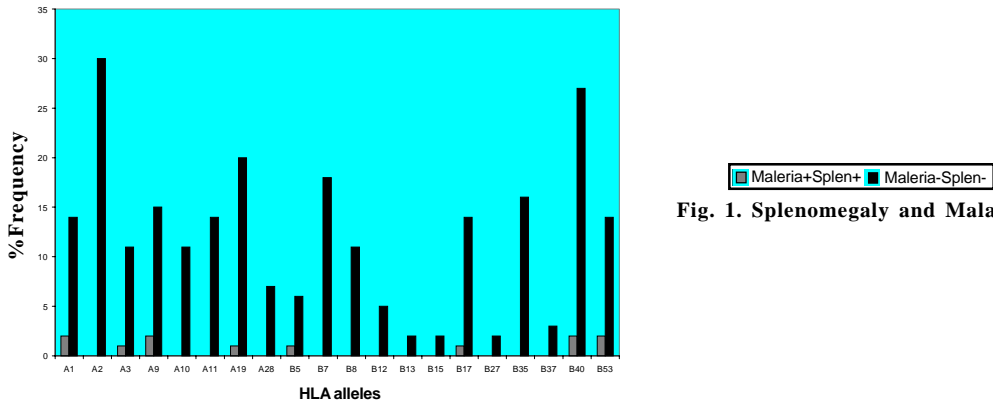


Fig. 1. Splenomegaly and Malaria

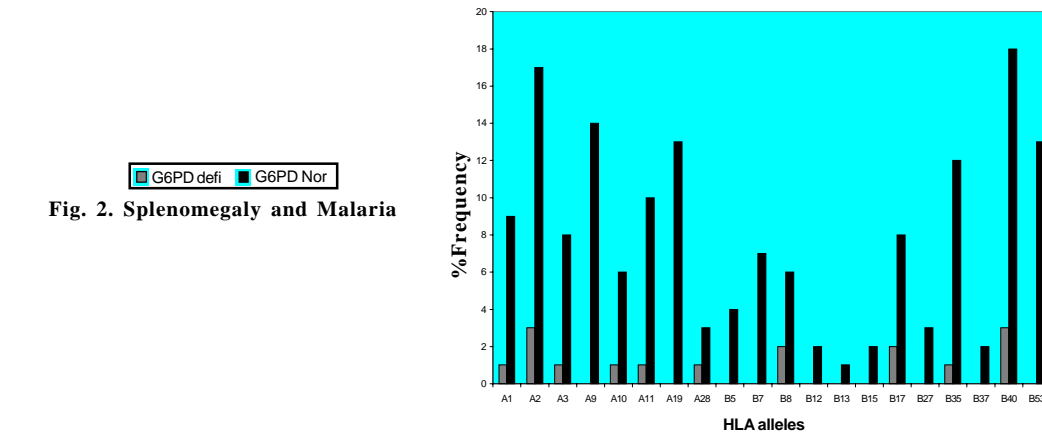


Fig. 2. Splenomegaly and Malaria

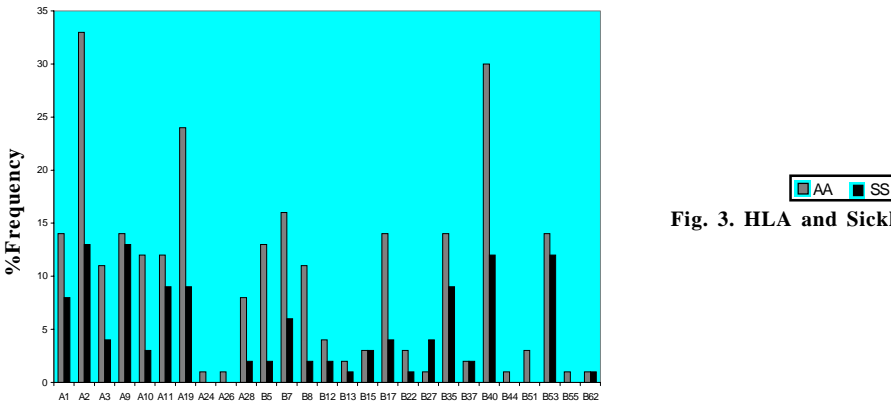


Fig. 3. HLA and Sickle cell status

DISCUSSION

The mortality caused by Malaria in high endemic parts of the world enabled the human survival to evolve a genetic trait, which can defend against malarial infection. These types of protective genetic measures have been observed with relation to polymorphism at the Sickle cell disease, G6PD and the HLA system (Hill et al., 1991, 1992). Such wide genetic variation in different populations has been a result of various factors like a) original gene pool b) migration and genetic drift c) gene selection under different environments and d) inbreeding pattern within the population (Bhatia, 1987). When a gene that has a potential for reducing fitness achieving high frequencies in a population it is necessary to assume the gene confers a survival advantage (Beutler, 1994). There has been significant differences in the HLA gene frequencies between malaria exposed and unexposed populations (Hill et al., 1991, 1992; Brown et al., 1989; Troye-Blomberg et al., 1991; Riley et al., 1991). Comprehensive analysis of HLA association with infectious diseases has allowed precise definition of susceptibility and protective alleles in large population of different ethnic groups. Infectious diseases may have exerted significant pressure on the development and maintenance of HLA polymorphism (Potts et al., 1990). Widespread and frequently fatal parasitic diseases such as Malaria have selectively maintained certain gene frequencies in endemic disease areas (Osaba et al., 1979). Further it has been speculated Sickle cell disease, G6PD deficiency and HLA polymorphism offers protection against malaria in a malaria endemic region.

Haldane proposed that the sickle cell gene provides an advantage in a malaria endemic region. Allison (1954) presented evidence for the balanced polymorphism of sickle cell gene and the sickle cell trait (heterozygotes) and G6PD has been suggested to provide resistance to falciparum malaria (Raper, 1955; Edington, 1965). The HLA polymorphism of Bhils and Pawras along with other Indian tribes (Selvakumar et al., 1987; Pitchappan et al., 1997; Paphia et al., 1983) revealed that all these tribes do not cluster together as a separate entity: suggesting that different tribes have different HLA profile, presumably because of their isolated origin. It is generally believed that the tribes were the early settlers in the Indian Plateau who were driven to

the hills by ethnically distinct invading populations (Sanghvi et al., 1981). They have high inbreeding co-efficiency. In India many of the forest and hilly areas that are especially vulnerable are endemic for Malaria. With the current resurgence of malaria in India and its importance as a national health issue it is important to know the prevalence of genetic factors associated with malaria that are present in individuals living in areas endemic for the disease.

KEY WORDS Immunity; genetic markers; tribes; West India; environment

ABSTRACT Haldane proposed that the sickle cell gene provides a survival advantage in malarial endemic regions. A high malarial endemicity has also been attributed to a high prevalence of G6PD deficiency in the Middle East and South East Asia. Moreover, the HLA antigens have been postulated to confer a protective immunity among the malaria endemic population. To evaluate a correlation in between these investigations among 146 tribals comprising of Bhils (91) and Pawras (55) living in Dhadgoan, Dhule district of Maharashtra where Malaria is endemic. It was interesting to note that HLA A1, A9, B5, B40 and B53 alleles were significantly increased among clinically splenomegaly positive patients with a history of Malaria when compared to normal controls (PF > 40% P < 0.0001). HLA A2, B8, B17 and B40 alleles were significantly increased among the G6PD deficient tribals when compared to non-deficient individuals (PF > 30%, P < 0.001). Further, it was observed that B53 was significantly increased among the sickle cell heterozygotes (PF > 45% P < 0.0001) of Bhils and Pawra tribes. A comparison of sickle cell trait and G6PD deficiency status with other Indian tribal populations reported revealed that in our study a 9% G6PD deficiency and 31 % sickle cell heterozygotes with a 17% Sickle cell disease among the tribal population studied. It will be of great interest if we further characterize the HLA, G6PD and Sickle cell gene by molecular methods to study the influence of various environmental factors like hilly terrain, natural calamities and epidemics in the prevalence of Malaria and its association with these parameters at a molecular level among these tribes.

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