HLA expression in human hepatocellular carcinoma

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Summary
This study examines the expression of MHC class I and II antigens and their related invariant chains in 70 cases of human hepatocellular carcinoma (HCC), using monoclonal (Mabs) and polyclonal antibodies. In comparison with normal hepatocytes, the majority (94.3%) of HCCs show enhancement or acquisition of HLA-A, B, C in either a cytoplasmic or membranous distribution, with staining being uniformly distributed throughout the specimen. HLA-A, B, C was accompanied by β₂-microglobulin expression in all but two cases. Although 44.9% of specimens showed HLA-DR expression, positively staining tumour cells were often sparse and heterogeneously distributed. By contrast, the invariant (I) chain, present in 47.1% of cases, was frequently intensively stained and extensive in distribution. HLA-DR staining was usually cytoplasmic although two cases showed faint membranous enhancement. In addition to HLA-DR and I-chain, two cases also showed HLA-DQ staining. Display of MHC antigens was not related to tumour differentiation or size of the lesion (resected vs. advanced tumours). It is possible that the acquisition of class I antigens by the majority of HCCs may influence tumour behaviour.

Materials and methods

Human tissues
The series comprised 70 samples of hepatocellular carcinoma from 56 black and 14 caucasian patients (66 male, 4 female). Hepatitis B viral (HBV) status and serum alpha-feto protein (AFP) levels were known in 60 cases.

Fresh tissue obtained at autopsy performed within 1h of death (32 cases), by needle biopsy (26), or during resection of small tumours (12), was snap-frozen in OCT compound (Ames Co., Div. of Miles Laboratories, Elkhart) and stored at -70°C.

Histopathological diagnosis was made on routinely processed H&E sections. Tumours were graded as well, moderately or poorly differentiated according to the criteria of the World Health Organization (World Health Organization, 1978).

In view of the current uncertainty regarding the expression of class I antigens on normal hepatocytes, 10 snap-frozen samples of histologically normal liver were included as control material. These were obtained from patients under investigation for conditions other than HCC or HBV infection.

Monoclonal antibodies
An anti-HLA class I Mab was obtained from Cappel Laboratories, Cochranville PA, (working dilution 1:50). This is a murine Mab of IgG₁, class directed against the 45,000 dalton polypeptide associated with β₂m, obtained by immunization of BALB/c mice with concavalin A-activated T-cell blasts followed by fusion of immune spleen cells with NS-1 myeloma cells (Cappel Product Information sheet). Mabs recognising nonpolymorphic regions of HLA-DR (anti-HLA-DR and HLA-DQ (Leu-10), were obtained from Becton-Dickinson, Sunnyvale CA; (working dilutions 1:200 and 1:50 respectively). VIC-Y1 is a monoclonal antibody recognising the invariant (I) chain of HLA-DR (Quaranta et al., 1984). Working dilution for VIC-Y1 was 1:200.

In addition, T-cell subsets were examined with a panel of commercially available Mabs: Leu-7 detecting killer and natural killer (NK) cells (Becton-Dickinson); OKT-4 reacting with helper/inducer T-cells (Ortho Pharmaceutical Co., Raritan, NJ); OKT-8 against suppressor/cytotoxic T-cells (Ortho). All were applied at a dilution of 1:10.
**Polyclonal antibodies**

Rabbit antibodies against the hepatitis B virus (HBV) surface antigen (anti-HBsAg), core antigen (anti-HBcAg), and human β2-m were purchased from Dako Immunoglobulins. They were applied for 30 min at dilutions of 1:50, 1:20 and 1:200 respectively.

**Immunoperoxidase techniques**

Cryostat sections (5 μm) were air-dried overnight at room temperature, fixed for 10 min in absolute acetone and stored wrapped in aluminium foil, at −20°C until use.

Monoclonal antibodies were demonstrated by a three step immunoperoxidase technique. Serial sections were rehydrated in PBS (pH 7.2) and incubated with the Mabs for 30 min at the dilutions specified above. The second and third steps comprised treatment for 30 min with rabbit-antimouse immunoglobulins (Igs) (Dako) at a dilution of 1:50, followed by peroxidase conjugated swine-antirabbit Igs (Dako), dilution 1:100, also for 30 min. Ten percent human AB serum was added to each, to reduce background staining. Each step was followed by vigorous washing in PBS. Sections were then incubated with 0.05% 3-amino-9-ethylcarbazole in 0.05 M acetate buffer (pH 4.9) and 0.01% H2O2. All staining was carried out at room temperature.

Polyclonal antibodies were used in a three-step unlabelled peroxidase-antiperoxidase procedure performed at room temperature. Sections were incubated with primary antibodies for 30 min at the dilutions specified above. This was followed by swine-anti rabbit Igs (dilution 1:100) and, thereafter, rabbit peroxidase-antiperoxidase complex (Dako; dilution 1:300), both for 30 min. Sections were washed in three changes of PBS between steps. The reaction product was developed with 3-amino-9-ethylcarbazole and H2O2. Negative controls consisted of omission of the primary antibody or use of the chromogen alone.

Slides were examined in batches according to primary antibody to allow later comparison between MHC products and their relevant invariant chains. Staining of HLA-A, -B, -C and β2-m was recorded as absent, cytoplasmic and membranous (± cytoplasmic). HLA-DR, -DQ and I chain staining were scored as 0 (absent), 1+ (<10% of hepatocytes positively staining), 2+ (10–50% positive), 3+ (>50% positive) and 4+ (>90% of cells positive).

**Results**

Of 70 cases examined, 23 were well, 33 moderately and 14 poorly differentiated HCC. Twelve were resected specimens, staged as 'early' HCC, and the remainder represented advanced tumours. HBsAg and HBcAg were demonstrated by immunostaining in one case.

**Class I expression**

In addition to sinusoidal staining (endothelial and Kupffer cells), weak, membranous display of HLA-A, -B, -C was observed on variable numbers of hepatocytes in all 10 histologically normal livers (Figure 1).

Positive staining for HLA-A, -B, -C was present in 66 of 70 cases (94.3%). Of the 66 positive tissues, 41 displayed membranous staining accompanied by variable intracytoplasmic positivity (Figure 2), while 23 showed a granular intracytoplasmic stain, sometimes accompanied by sinusoidal enhancement (Figure 3). In 4 specimens the tumour was negative for the antigen although adjacent non-neoplastic hepatocytes showed enhanced membranous HLA-A, -B, -C display (Figure 4). Staining was of uniform consistency in positive specimens with the cytoplasmic or membranous pattern being preserved throughout. In all tumours as well as adjacent non-neoplastic hepatocytes displaying class I

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**Figure 1** Histologically normal liver. Intense HLA-A, B, C staining is observed within sinusoids, while hepatocyte membranes show variable faint positivity (arrowed). (Immunoperoxidase x 640).

**Figure 2** Membranous HLA-A, B, C display in hepatocellular carcinoma. Staining is typically uniform throughout the specimen and is more intense than that observed on apparently normal hepatocytes as illustrated in Figure 1. (Immunoperoxidase x 323).

**Figure 3** Hepatocellular carcinoma showing diffuse cytoplasmic staining for HLA-A, B, C. The membranous enhancement observed in Figure 2 is absent. (Immunoperoxidase x 528).
antigens, staining was considerably more intense than was observed in the control group.

$\beta_m$ staining accompanied HLA-A, -B, -C in all but two cases which were $\beta_m$ negative but showed faint cytoplasmic HLA-A, -B, -C positivity. The pattern of expression was either cytoplasmic or membranous and generally correlated with the heavy chain.

**Class II expression**

Twenty-nine cases (41.4%) were positive for HLA-DR. Staining was invariably heterogeneous (Figure 5). In 10 of these the extent of staining was graded as $1+$, 9 were read as $2+$, 6 as $3+$, and 4 showed $4+$ positivity.

All cases positive for HLA-DR also showed VICY1 staining. In addition a further 4 cases were VICY1"/HLA-DR". In contrast to the relationship between HLA-A, -B, -C and $\beta_m$, however, I chain staining was invariably far in excess of that observed for its associated MHC product (Figure 6). Thus, of 33 cases (47.1%) staining with VICY1, 4 were graded as $1+$, 5 as $2+$, 10 as $3+$, and 14 showed $4+$ positivity.

For both Mabs, staining tended to be heterogenous, with staining intensity varying from cell to cell. In tumours showing a pseudoglandular pattern the HLA-DR positivity was often confined to the apical portion of the cell (Figure 5). Both antigens tended to be intracytoplasmic although in two tumours occasional small foci of membranous HLA-DR expression were noted.

Two cases stained for HLA-DQ in addition to HLA-DR and I chain. Staining was less intense and more focal than was observed for HLA-DR on serial sections.

All tumours showed a mild to moderate T-cell infiltrate, with OKT-4+ cells (helper/inducer T-cells) being slightly in excess of OKT-8+ cells (cytotoxic-suppressor T-cells). T-cells were usually confined to intervening stroma and at the margins of resected lesions. There was little infiltration between tumour cells, and NK cells were extremely scarce in all specimens. No relationship between inflammatory infiltrate and MHC status was demonstrated although in two cases there was enhanced HLA-DR and I chain staining in a perisepal distribution and around clusters of helper/inducer T-cells.

No relationship was observed between expression of class I or class II antigens and tumour differentiation, stage or HBV status (data not shown).

**Discussion**

In this study we have examined the patterns of MHC class I and II display in hepatocellular carcinoma. We have also attempted to qualify the relationship between these antigens and their respective invariant chains.

Expression of class I antigens by normal hepatocytes remains controversial. In our series of non-tumorous biopsies there was faint membranous expression of class I antigens by some liver cells. However such expression may represent a facile response of the hepatocyte to a variety of stimuli (Nagafuchi et al., 1985; Fukusato et al., 1986). Biopsies read as histologically normal in this and other studies, and which express HLA-A, -B, -C, may not represent "normal" liver, with unrecognised immunological factors or agents such as drugs or alcohol promoting the expression of HLA products at the hepatocyte membrane. This may explain the variable findings in the different series reported to date (Saunders et al., 1979; Montano et al., 1982; Ponder et al., 1983; Lautenschlager et al., 1984; Nagafuchi et al., 1985; Fukusato et al., 1986; Nagafuchi & Scheuer, 1986). However, if HLA-A, -B, -C is expressed at all on normal liver cells then it would appear to be at very low levels.
Our results suggest, therefore, that malignant transformation of hepatocytes is characterized by expression of class I antigens that is at least enhanced if not acquired de novo, with virtually all tumours in this series showing intense staining for class I antigens. This finding is in keeping with observations reported elsewhere for small series of HCC (Fukusato et al., 1986; Mazzeo et al., 1986).

This contrasts with the situation reported for other human tumours, including colorectal carcinomas in which there is sometimes loss of expression in comparison with the non-neoplastic epithelium (Dhaar et al., 1982; Ciba et al., 1984; Momburg et al., 1986; van den Ingh et al., 1987). Furthermore it is suggested by some investigators that loss of display is related to progressive de-differentiation and prognosis (Momburg et al., 1986; van den Ingh et al., 1987). This observation has also been made for malignant melanomas in which attenuated or absent class I expression is associated with a high degree of malignancy (Ruiter et al., 1986). We have not observed any relationship between MHC expression and tumour differentiation in HCC.

Class II antigens were expressed in nearly half of the cases in our series. Expression was heterogeneous and often rather sparse, however, and was always associated with VCY1 positivity, although the latter typically showed far more extensive and intense staining than HLA-DR

The Mab VICY1 is directed at the core protein of the class II-associated chain (Giacotto et al., 1986). It is an unusual transmembranous glycoprotein with some structural similarity to certain membrane receptor molecules including the transferrin receptor (Creswell et al., 1987). Its function at this time is unknown although it may participate in one or more of a variety of activities: biosynthesis, transport/recycling, immune mediation or other as yet unrecognised functions (Koch & Harris, 1984; Long, 1985; Giacotto et al., 1986; Creswell et al., 1987). Although some cases in our series expressed the I chain in the absence of HLA-DR, it may be that this represents tissue selection rather than true absence of HLA-DR, as the class II antigen is expressed extremely sparsely in some cases showing strong VICY1 staining. It is noted, however, that the murine equivalent of the I chain has been demonstrated in the absence of class II antigen in B-lymphoid and myeloid cell lines (Koch & Harris, 1984), and a subset of tumour cells in one case of mediastinal B-cell lymphoma has been reported to be positive for I chain in the absence of HLA-DR and -DQ (Möller et al., 1986). Furthermore, hyperexpression of the I chain has been demonstrated by electrophoretic techniques in Epstein-Barr virus-transformed leukemic cells (Spirito et al., 1985). To our knowledge the VICY1 antibody has not previously been applied to epithelial neoplasms, and our observations suggest that it may be of value in the study of class II expression in other tumours.

Although HLA-DR expression has been observed in reactive hepatocytes during HBV infection (Fachetti et al., 1985; van den Oord et al., 1986), data for HLA-DQ expression are not available. We believe that the expression of HLA-DQ in some tumours in this series implies that non-neoplastic liver cells may also express this antigen, but in a sequential or dissociated fashion (Solid et al., 1987).

It has been suggested that the ability to display class II antigens in response to appropriate stimuli may be constitutive to all cells and that it is the failure to express the antigen by a percentage of tumours which represents the ‘aberrant’ state (Moore & Ghosh, 1987). The observation in two cases that both HLA-DR and I chain staining are enhanced in a perisepal distribution may support this, lymphocytic infiltration of the adjacent lymphoid infiltrate being responsible for the enhancement. Conversely, no such accentuation of staining could be demonstrated for the class I antigens and the regular and diffuse nature of antigen expression in all but a minority of cases suggests to us that it represents a constitutive component in malignant transformation of hepatocytes. Whether the lack of display in some tumours represents a pre- or post-translational event is not known.

The MHC antigens function primarily at the cell surface. In a small minority of malignant hepatocytes expressing HLA-A, -B, -C, and in the vast majority displaying HLA-DR, expression is confined to the cytoplasm. This has also been observed with respect to class I antigens in other tumour systems (Lampert et al., 1985; Kadin, 1980). The effect of this distribution on functional integrity is unknown.

Just as carcinogenesis is accepted to be multifactorial in origin, it is evident that subsequent tumour growth and metastasis is the consequence of numerous complex factors including characteristics of the tumour cell and host-tumour interactions. The host’s immune response to the neoplastic cell is one such factor and any modification of the tumour that will enable it to escape immune surveillance could contribute to subsequent growth and metastasis. The possibility that such modification could affect metastatic behaviour has formed the basis of many paradigms involving viral, chemical, or radiation induced cell transformation or neoplasia in culture systems and/or animal models (Goodenow et al., 1985). There is enhancement of metastatic potential in tumours showing loss or attenuation of class I antigens (Schrier et al., 1983), this being reversible by transfection with the appropriate class I gene (Tanaka et al., 1985). Furthermore, modified class I antigens may themselves act as tumour-specific antigens (Philipps et al., 1986), or influence tumour cell behaviour by interaction with non-MHC genes or by an hormonal effect (Demant, 1986).

MHC expression and tumour behaviour may also be related in some human tumours, including colorectal carcinomas (van den Ingh et al., 1987), melanomas (Ruiter et al., 1986), breast carcinomas (Fleming et al., 1981), small cell carcinoma of lung (Dyke et al., 1985), and neuroblastomas (Whelan et al., 1985). The possibility that this relationship may influence the behaviour of HCC merits some consideration.

HCC is a tumour associated with a uniformly fatal outcome unless detected and resected at an early stage (Kew & Geddes, 1984; Okuda, 1986). Although progress from the time of diagnosis to death is typically rapid (Kew & Geddes, 1984), there is a prolonged subclinical period (Sheu et al., 1985) and the tumour shows an early and sustained proclivity for capsular and blood vessel infiltration (Wakasa et al., 1985; Kew & Paterson, 1985).

Advanced HCCs are typically massive and show extensive vascular infiltration and intrahepatic spread. Despite this, metastases are often confined to clinically insignificant hilar lymph node and pulmonary deposits (Anthony, 1978; Kew & Paterson, 1985), with death resulting from hepatic failure or complications of an associated cirrhosis.

While many other factors may contribute to the metastatic potential of HCC, the possibility that class I expression in the majority of HCCs may restrict extensive extrahepatic tumour spread should be further investigated by the examination of defined intra- and extrahepatic metastases and comparison with the primary lesion.

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